Report of the Meeting of the WOAH Biological Standards Commission

Introduction and Member contribution

A meeting of the WOAH Biological Standards Commission (hereafter called 'the Commission') was held from 4 to 8 September 2023 at the WOAH Headquarters in Paris, France. During the meeting, 15 chapters from the WOAH *Manual of Diagnostic Tests and Vaccines for Terrestrial Animals (Terrestrial Manual)* were approved for circulation for first-round Member comment, six Reference Centre applications and eight nominations for replacement experts were also evaluated.

Annexes

Texts in Annexes 4 to 18 are presented for first-round comment.

How to submit comments

Members are reminded that they should submit the rationale for all their proposed changes to the texts, and include references where relevant for the Commission to consult. Your participation in the WOAH Standard-setting process is valued. Thank you for your engagement in the process!

The guidance should be followed when submitting comments:

- 1. Comments may be general or specific, but specific comments are more valuable. General comments should be such that some conclusion and action can be taken in response to them. For example, instead of stating "This test is no longer used in our laboratory", indicate the reasons the test is no longer used and what test is used instead.
- 2. Specific comments should be identified by indicating the line number in the text, to facilitate the editorial process.
- 3. Highlighting typing or technical errors is welcome, but the correct word or figure should be indicated in its place. For example, instead of indicating simply "0.8 M is too high", the preferred value should also be indicated.
- 4. Bear in mind that the introductory chapters (Part 1 of the *Terrestrial Manual*) set general standards for the management of veterinary diagnostic laboratories and vaccine facilities and are not intended to be exhaustive, and indeed none of the chapters can give a completely comprehensive cover of the subject, otherwise the *Terrestrial Manual* would be too long. However, assistance in indicating priorities is always helpful.
- 5. The *Terrestrial Manual* is intended for world-wide use. The chapters need to reflect the development of new technology, while maintaining the established methods, usually requiring less sophisticated apparatus. New technology should not be described in detail until it has gained wide acceptance as a reliable method.
- 6. We recommend that if you have no specific comments, please respond to the WOAH to that effect.
- 7. Any comments, proposed changes or revisions should be supported by clear evidence (the scientific rationale) such that some conclusion and action can be taken in response to them.

Deadline to comment

Comments on relevant texts in this report must reach the Headquarters by <u>7 December 2023</u> to be considered at the February 2024 meeting of the Commission.

Where to send comments

All comments should be sent to the Science Department at: BSC.Secretariat@woah.org

Date of the next meeting

The Commission noted the dates for its next meeting: 5 to 9 February 2024



Science Department BSC.secretariat@woah.org 12, rue de Prony 75017 Paris, France

Table of Contents

1.	Welc	ome from the directors	5				
	1.1.	Director General	5				
	1.2.	Deputy Director General, International Standards and Science	5				
	1.3.	Updates from the WOAH Headquarters	5				
		1.3.1. Transparency of the WOAH process for the elaboration of Standards	5				
2.	Adop	tion of the agenda	6				
3.	Colla	boration with other Specialist Commissions	6				
	3.1.	Horizontal issues among the Specialist Commissions	6				
		3.1.1. Case definitions: infection New World (<i>Cochliomyia hominivorax</i>) and Old World (<i>Chrysomya bezziana</i>) screwworms, and Crimean–Congo haemorrhagic fever (revisited)	6				
	3.2.	Scientific Commission for Animal Diseases	6				
	3.3.	Terrestrial Animal Health Standards Commission	6				
		3.3.1. Updates from the February 2023 Code Commission meeting	6				
		3.3.2. Biological Standards Commission's recommendations to the Terrestrial Animal Health Standards Commission	7				
		3.3.3. Meeting of the Bureaus (7 September 2023)	7				
		3.3.4. Comments on Chapter.5.8. International transfer and laboratory containment of animal pathogenic agents	7				
		3.3.5. Questions on Chapter 6.10. Responsible and prudent use of antimicrobial agents in veterinary medicine	7				
	3.4.	Aquatic Animal Health Standards Commission	7				
4.	Work	Programme	7				
5.	Manu	al of Diagnostic Tests and Vaccines for Terrestrial Animals	7				
	5.1.	. Review of draft chapters received for endorsement for circulation for first round Member comment					
	5.2.	Follow-up from September 2022: conclusions and recommendations from the WOAH <i>Scientific and Technical Review</i> issue on diagnostic test validation science	10				
		5.2.1. Progress on development of a validation report form for tests recommended in the <i>Terrestrial Manual</i>	10				
		5.2.2. Progress on development of a template for a new <i>Terrestrial Manual</i> section on the rationale behind the selection of tests included in Table 1. <i>Test methods available and their purpose</i>	11				
	5.3.	Inclusion of videos on diagnostic techniques on the WOAH website disease portals: review of submitted videos	11				
	5.4.	Further review of Chapter 1.1.6 Validation of diagnostic assays for infectious diseases of terrestrial animals	11				
	5.5.	Follow-up from February: need for a definition of latent period in the chapter on foot and mouth disease	11				
	5.6.	Review and alignment of the validity criteria of the PD ₅₀ and PPG tests for foot and mouth disease between the <i>Terrestrial Manual</i> and the European Pharmacopoeia	11				
	5.7.	Purpose of the list of contributors and addresses in the <i>Terrestrial Manual</i>	12				
	5.8.	Publication of Member comments and need to review the Commission's practice	12				
	5.9.	Terrestrial Manual status: update on chapters selected for the 2024/2025 review cycle	12				
	5.10.	Update on WOAH Standards Online Navigation Tool Project	13				
	5.11.	Terrestrial Manual chapters on non-listed diseases	14				

6.	WOA	H Reference Centres	14
	6.1.	Improving and automating the performance review for annual reports of Reference Laboratories using a risk-based approach	14
	6.2.	Applications for WOAH Reference Centre status	14
	6.3.	Changes of experts at WOAH Reference Centres	15
	6.4.	Review of new and pending applications for laboratory twinning	16
	6.5.	Analysis of the questionnaire sent to all Reference Laboratories	16
	6.6.	Feedback from Laboratories that are not complying with the key ToR	16
	6.7.	Develop a plan of how to evaluate the progress made since the Collaborating Centres submitted their 5-years work plans	16
	6.8.	Update on the three Reference Laboratory network (ASF, PPR and rabies)	17
	6.9.	Review of the current list of Main Focus Areas and Specialties	17
	6.10.	Clarify the role of the Contact Point in providing advice and services to WOAH Members	18
7.	Ad ho	oc Groups: Update on activities of past <i>ad hoc</i> Groups	18
	7.1.	<i>Ad hoc</i> Group on Replacement of the International Standard Bovine Tuberculin (ISBT) and Avian Tuberculin (ISAT)	18
8.	Interr	national Standardisation/Harmonisation	18
	8.1.	WOAH Register of diagnostic kits – update and review of new or renewed applications	18
		8.1.1. Ongoing application for Genelix™ ASFV real-time PCR detection kit	18
		8.1.2. Renewal of Avian Influenza Antibody test kit (registration number 20080203)	19
		8.1.3. Renewal of Newcastle Disease Antibody test kit (registration number 20140109)	19
		8.1.4. Update on WOAH SOP and application form	19
	8.2.	Standardisation programme	19
		8.2.1. Association française de normalisation: follow-up from February 2023	19
		8.2.2. Project to extend the list of WOAH-approved reference reagents: review of guidelines	19
9.	Follo	w-up from the General Session	19
	9.1.	Excerpt from the Final Report: comments from Members	19
	9.2.	The Animal Health Forum and the adopted Resolution on avian influenza	20
10.	Confe	erences, Workshops, Meetings	21
	10.1.	Update on the WAVLD seminar in Lyon, France in 2023 and Commission involvement in future seminars.	21
11.	Matte	rs of interest for consideration or information	21
	11.1.	Update on OFFLU	21
	11.2.	Update on rinderpest	22
	11.3.	Update on VICH activities	22
	11.4.	Update on the Grand Challenge for sustainable laboratories	22
	11.5.	Biosafety Research Roadmap	23
	11.6.	Collaboration with WOAH in the implementation of non-animal testing for veterinary vaccines batch release testing	23
	11.7.	Update on the development of guidelines for alternative strategies for the control of <i>Mycobacterium tuberculosis</i> complex infection in livestock	23
	11.8.	Composition of the WOAH editorial board for the Scientific and Technical Review	23
		Update on activities under the IHSC-WOAH collaboration agreement and consultancy project in Asia	
	11.10	. Update on the virtual biobank project	24

List of Annexes

Annex 1.	Adopted Agenda	
Annex 2.	List of Participants	27
Annex 3.	Work Programme for the WOAH Biological Standards Commission	
Annex 4.	Item 5.1. – Chapter 1.1.5. Quality management in veterinary testing laboratories	30
Annex 5.	Item 5.1. – Chapter 1.1.9. Tests for sterility and freedom from contamination of biological materials intended for veterinary use	40
Annex 6.	Item 5.1. – Chapter 2.2.4 Measurement uncertainty	57
Annex 7.	Item 5.1 Chapter 2.2.6. Selection and use of reference samples and panels	63
Annex 8.	Item 3.1.1. – Chapter 3.1.5. Crimean–Congo haemorrhagic fever	75
Annex 9.	Item 5.1. – Chapter 3.3.6. Avian tuberculosis	77
Annex 10.	Item 5.1. – Chapter 3.4.1. Bovine anaplasmosis	90
Annex 11.	Item 5.1. – Chapter 3.4.7. Bovine viral diarrhoea	110
Annex 12.	Item 5.1 – Chapter 3.4.12 Lumpy Skin Disease	132
Annex 13.	Item 5.1 Chapter 3.6.9. Equine rhinopneumonitis (infection with equid herpesvirus-1)	153
Annex 14.	Item 5.1. – Chapter 3.8.1. Border disease	170
Annex 15.	Item 5.1. – Chapter 3.8.12. Sheep pox and goat pox	183
Annex 16.	Item 5.1 Chapter 3.9.1. African swine fever (infection with African swine fever virus)	196
Annex 17.	Item 5.1 Chapter 3.10.4. Infection with Campylobacter jejuni and C. coli	220
Annex 18.	Item 5.1. – Chapter 3.10.8. Toxoplasmosis	231
Annex 19.	Analysis of the questionnaire for Reference Laboratories	244
Annex 20.	List of Main Focus Areas and Specialties for WOAH Collaborating Centres	245

1. Welcome from the directors

1.1. Director General

Dr Monique Eloit, the WOAH Director General, met the Biological Standards Commission on 5 September and thanked its members for their support and commitment to achieving WOAH objectives.

Dr Eloit remarked on the positive outcomes of the 90th General Session, highlighting the favourable response to the change in the Session's format, which included an Animal Health Forum on Avian Influenza. Dr Eloit emphasised that the forum facilitated interactive discussions and encouraged exchanges from both a policy and technical perspective.

Dr Eloit informed the Commission that WOAH is currently undergoing a consultancy to evaluate the Organisation's *Basic Texts* from both a technical and legal viewpoint. The importance of this consultancy is to introduce a more robust and transparent approach to the organisation's procedures, supported by a solid legal basis. Dr Eloit pointed out the need to determine which fundamental documents or SOPs ¹ necessitate revision and subsequent endorsement by the Assembly. The revision of the *Basic Texts* is essential to maintaining WOAH's credibility among stakeholders, and Members. This assessment will be completed in time for the celebrations of WOAH's 100th anniversary in May 2024.

Finally, Dr Eloit reminded the Commission that the deadline for submission of nominations for election to WOAH Specialist Commissions was 8 September 2023, and that the elections will take place during the General Session in May 2024.

The Commission thanked Dr Eloit for these updates.

1.2. Deputy Director General, International Standards and Science

Dr Montserrat Arroyo, WOAH Deputy Director General, International Standards and Science, welcomed members of the Biological Standards Commission and thanked them for their ongoing contributions to the work of WOAH.

Dr Arroyo informed the Commission that the Organisation is currently dedicating efforts to various IT projects with the aim of creating tools that will facilitate access to WOAH services and practices as detailed in the organisation's *Basic Texts*. Among these tools are the evolution of the system for collecting annual reports from Reference Centres, a digitised system for navigating the Code and Manuals, an improved system for self-declaration of disease status, and a repository of PVS reports, all with the goal of improving and simplifying access to these tools, ensuring transparency, and enhancing the traceability of WOAH's work, while also interconnecting all the tools.

Dr Arroyo also expressed her satisfaction with the past General Session and highlighted that the Organisation will celebrate its 100th anniversary in the coming year. She congratulated the Commission on its interactions with the other Specialist Commissions, emphasising the importance of harmonising and adopting a consistent approach to common work themes.

Finally, Dr Arroyo asked the Commission to discuss the rationale behind the maintenance of chapters in the *Terrestrial Manual* on unlisted diseases, underscoring the challenges and resources required for this effort, both for the Secretariat and the experts, and for the Commission itself (see agenda item 5.11).

The members of the Commission thanked Dr Arroyo for the excellent support provided by the WOAH Secretariat.

1.3. Updates from the WOAH Headquarters

1.3.1. Transparency of the WOAH process for the elaboration of Standards

The Secretariat informed the Commission that the WOAH Director General had agreed to implement a stepwise approach to improve the transparency of the WOAH process for the elaboration of Standards that will include the publication of comments considered, responses, and an evolution of the reports of the Aquatic Animals Commission, the Code Commission and the Biological Standards Commission. This is also to align with the 7th Strategic Plan. The Secretariat also noted that this proposal had been discussed with the Presidents of the three Commissions (Aquatic Animals, Biological Standards and Code Commissions) at a meeting after the 90th General Session in May 2023 and that they supported this approach.

¹ SOPs: Standard operating procedures

The Secretariat explained that this process also aims to ensure that Members can gain a better understanding of the complexity and range of opinions, as well as of Commission decisions, and that this will result in a better understanding of Members concerns and should also improve the quality of the comments received.

The Secretariat explained that this would be a progressive process, that will start in March/April 2024 with the publication of comments considered on new and revised standards during the February 2024 meetings of the respective Commissions on the Delegates only website at the same time as the respective February 2024 report. This process will include an evolution of the Commission reports towards full transparency of comments considered and Commissions responses, which will result in better documentation and traceability of the WOAH process for the elaboration of Standards. The Secretariat noted that Delegates will be kept well informed throughout this process, including a detailed communication that will be sent after the publication of this report.

2. Adoption of the agenda

The proposed agenda was presented and adopted. Dr Emmanuel Couacy-Hymann chaired the meeting and the WOAH Secretariat acted as rapporteur. The agenda and the list of participants can be found at <u>Annexes 1</u> and <u>2</u> respectively.

3. Collaboration with other Specialist Commissions

3.1. Horizontal issues among the Specialist Commissions

3.1.1. Case definitions: infection New World (Cochliomyia hominivorax) and Old World (Chrysomya bezziana) screwworms, and Crimean–Congo haemorrhagic fever (revisited)

The Biological Standards Commission discussed the case definitions for New World (*Cochliomyia hominivorax*) and Old World (*Chrysomya bezziana*) screwworms and gave its recommendations to the Scientific Commission for Animal Diseases (see agenda item 9.3.2.1 of the report of the meeting of the Scientific Commission for Animal Diseases, 11–15 September 2023).

Regarding the case definition for Crimean–Congo haemorrhagic fever that the Biological Standards Commission had reviewed at the February 2023 meeting, the Scientific Commission submitted technical questions on the demonstration of active infection by ELISA² (IgG, competitive and IgM). The Biological Standards Commission agreed with the recommendation of the lead expert on the diagnostic protocols to establish serological evidence of active infection, including using two different serological tests each based on a different antigen for the detection of IgM antibodies given the potential for cross-reactivity, or by seroconversion based on a rise in total or IgG antibody titres on samples taken at 2–4 weeks apart. The lead expert added two footnotes to the purpose *Confirmation of clinical cases in animals* in Table 1 *Diagnostic test formats for Crimean–Congo haemorrhagic fever virus infections in animals*, of *Terrestrial Manual* Chapter 3.1.5 to capture this information. The amended chapter is included in the batch of chapters that will be sent for first-round comment in October 2023 (see agenda item 5.1).

For details of the case definition, refer to the September 2023 report of the Scientific Commission.

3.2. Scientific Commission for Animal Diseases

Nothing for this meeting.

3.3. Terrestrial Animal Health Standards Commission

Matters between the Terrestrial Animal Health Standards Commission and the Biological Standards Commission.

3.3.1. Updates from the February 2023 Code Commission meeting

The Biological Standards Commission was updated by the Secretariat of the Code Commission on the current topics under review by the Code Commission to ensure complementarity and alignment of the two Commission's respective work programmes.

² ELISA: enzyme-linked immunosorbent assay

3.3.2. Biological Standards Commission's recommendations to the Terrestrial Animal Health Standards Commission

At the February 2023 meeting, the Code Commission had referred a Member request to define the latent period for FMD³ virus to the Biological Standards Commission, noting that such detail should be in the *Terrestrial Manual* and not the *Terrestrial Code*. The Biological Standards Commission asked the network of Reference Laboratory experts if a definition of latent period was a useful and necessary addition to the *Terrestrial Manual* chapter, and if it was, to develop a definition. The experts responded that the latent period for FMD is usually defined as the time between infection and the point when an animal is infectious as opposed to the incubation period, which is defined in the *Terrestrial Code* as the time from infection to the first clinical signs. The latent period is usually considered to be shorter than the incubation period (by 0–4 days), but the experts felt that it would be difficult to define in terms of days with a high degree of precision. Although the latent period is an important epidemiological parameter, the experts could not see how it would be useful to include this term in either the *Terrestrial Code* or *Terrestrial Manual* chapter. The Biological Standards Commission therefore would not recommend its addition to the *Terrestrial Manual*, and would inform the Code Commission of its position (see also agenda item 5.5).

3.3.3. Meeting of the Bureaus (7 September 2023)

See agenda item 3.2. of the report of the meeting of the Terrestrial Animal Health Standards Commission, 5– 14 September 2023

3.3.4. Comments on Chapter.5.8. International transfer and laboratory containment of animal pathogenic agents

The advice of the Biological Standards Commission was sought regarding the need to revise *Terrestrial Code* Chapter 5.8. *International transfer and laboratory containment of animal pathogenic agents*. The Biological Standards Commission agreed with a Member comment that *Terrestrial Manual* Chapter 1.1.4. *Biosafety and biosecurity: Standard for managing biological risk in the veterinary laboratory and animal facilities* (last adopted in 2015) no longer refers to classification of pathogens into risk categories, but rather recommends a risk analysis approach to the management of biological risks for biosafety and biosecurity in veterinary laboratories and animal facilities. The *Terrestrial Code* chapter therefore is not harmonised with the *Terrestrial Manual* and should be updated.

3.3.5. Questions on Chapter 6.10. Responsible and prudent use of antimicrobial agents in veterinary medicine

Regarding Chapter 6.10. *Responsible and prudent use of antimicrobial agents in veterinary medicine,* the Code Commission had received comments referring to the establishment of clinical breakpoints. The Code Commission considered that clinical breakpoints should be established in accordance with *Terrestrial Manual* Chapter 2.1.1. *Laboratory methodologies for bacterial antimicrobial susceptibility testing*. The Biological Standards Commission agreed to liaise with the Working Group on Antimircrobial Resistance and other WOAH experts to update chapter 2.1.1, ensure it aligns with Codex Alimentarius Committee guidelines, and that the issue of the establishment of clinical breakpoints is fully addressed. The chapter would be added to the 2024/2025 review cycle, and thus could be proposed for adoption at the same time as *Terrestrial Code* chapter 6.10.

3.4. Aquatic Animal Health Standards Commission

Nothing for this meeting.

4. Work Programme

The updated work programme was agreed and can be found at Annex 3.

5. Manual of Diagnostic Tests and Vaccines for Terrestrial Animals

For this Agenda Item, the Commission was joined by Dr Steven Edwards, Consultant Editor of the WOAH *Terrestrial* Manual.

³ FMD: foot and mouth disease

5.1. Review of draft chapters received for endorsement for circulation for first round Member comment

The Commission reviewed 16 draft chapters and approved 14 for circulation, some subject to clarification of certain points by the experts, for first-round Member comment and eventual proposal for adoption by the Assembly in May 2024. The 14 chapters and a brief summary of the main amendments are provided below:

1.1.5. Quality management in veterinary testing laboratories: updated the references and URLs; clarified that limited availability of suitable material may render validation difficult; included substantial technical updates to the sections on: accreditation; determination of the scope of the quality management system or of the laboratory's accreditation; quality assurance, quality control and proficiency testing; validation of the test method; and estimation of measurement uncertainty; updated the section on strategic planning.

The revised Chapter 1.1.5. *Quality management in veterinary testing laboratories* is presented as <u>Annex 4</u> for first-round comments.

1.1.9. Tests for sterility and freedom from contamination of biological materials intended for veterinary use: given the importance of the laboratory perspective, the chapter has been updated to give the reader an example-based overview of tests and their regulatory background included brief illustrative examples of vaccine contamination in the introduction and in more detail in Section G. Protocol examples – these examples are a strong motivator for extraneous agent testing; updated Section A. An overview of testing approaches with more recent opportunities and challenges; merged sections on living and inactivated viruses and bacteria, e.g. B & C, D & F, to simplify and streamline the chapter; the examples given in Section G. Protocol examples are clearly marked as non-prescriptive and non-exhaustive examples; updated references and weblinks.

The revised Chapter 1.1.9. Tests for sterility and freedom from contamination of biological materials intended for veterinary use is presented as <u>Annex 5</u> for first-round comments.

2.2.4. *Measurement uncertainty:* removed reference to the "WOAH Validation Standard" because the newly adopted validation chapter in the *Terrestrial Manual* will differ from the future one in the *Aquatic Manual* so there is no longer one Standard that applies to both Manuals: the title of chapter 1.1.6 has thus been changed to "*Validation of diagnostic assays for infectious diseases of terrestrial animals* – this change will be made to all chapters in *Terrestrial Manual* Section 2.2. *Validation of diagnostic tests*; clarified that the method described in the chapter is known as the "top-down" approach, and included information of its requirements along with a section on the scope and limitations of the top-down approach; added an example of a measurement uncertainty calculation in molecular tests.

The revised Chapter 2.2.4. *Measurement uncertainty* is presented as <u>Annex 6</u> for first-round comments.

2.2.6. Selection and use of reference samples and panels: updated cross references to Chapter 1.1.6 Validation of diagnostic assays for infectious diseases of terrestrial animals; added a figure on the documentation that is required of reference materials; added a list of references for further reading.

The revised Chapter 2.2.6. *Selection and use of reference samples and panels* is presented as <u>Annex 7</u> for first-round comments.

3.3.6. Avian tuberculosis: updated the nomenclature and classification of species in the genus Mycobacterium; reviewed the ratings of some of the tests in Table 1 Test methods available for the diagnosis of avian tuberculosis and their purpose; updated the section on nucleic acid recognition methods; added a section on the stained antigen test; updated the section on the production and minimum requirements for tuberculin production, and the list of references. NB: the Commission agreed to include the information on avian tuberculin production given in Section C Requirements for diagnostic biologicals in chapter 3.1.13 Mammalian tuberculosis (infection with Mycobacterium tuberculosis complex).

The revised Chapter 3.3.6. Avian tuberculosis is presented as Annex 9 for first-round comments.

3.4.1. *Bovine anaplasmosis:* updated the information in the introduction to the chapter; added an illustration of a stained blood smear showing *Anaplasma marginale* initial bodies; the thoroughly updated the section on the PCR⁴, including adding a table of primer sequences, and the section on the ELISAs including adding a displacement double-antigen sandwich ELISA that has been developed to differentiate between *A. marginale* and *A. centrale* antibodies; emphasised that the complement fixation test has variable sensitivity and removed it from Table 1; reviewed the ratings of some of the tests in Table 1 *Test methods available for the diagnosis*

⁴ PCR: polymerase chain reaction

of bovine anaplasmosis and their purpose – the experts also developed justification tables for the ratings of the tests in Table 1 for each purpose: these tables can be accessed by clicking on the individual purpose. The Commission believes that the information in these justification tables will be extremely useful to users of the *Terrestrial Manual* when deciding which test to use for a given purpose.

The revised Chapter 3.4.1. Bovine anaplasmosis is presented as Annex 10 for first-round comments.

3.4.7. Bovine viral diarrhoea: updated the taxonomy; reviewed the ratings of some of the tests in Table 1 *Test methods available for the diagnosis of bovine viral diarrhoea and their purpose* and developed justification tables for the ratings of the tests in Table 1 for each purpose.

The revised Chapter 3.4.7. Bovine viral diarrhoea is presented as Annex 11 for first-round comments.

3.4.12. *Lumpy skin disease* (vaccine section only): added text on the scarcity of information on the role of wildlife in lumpy skin disease (LSD) epidemiology; thoroughly updated Section C *Requirements for vaccines*.

The revised Chapter 3.4.12. *Lumpy skin disease* (vaccine section only) is presented as <u>Annex 12</u> for first-round comments.

3.6.9. Equine rhinopneumonitis (infection with equid herpesvirus-1): clarified that the chapter covers infection with equid herpesvirus-1: most of the information on equid herspsvirus-4 has been removed as EHV4 is not listed; thoroughly updated Section B *Diagnostic techniques* in particular the section on virus detection by PCR, which now has a table on primer and probe sequences for a number of real-time PCRs and subsections on POC⁵ molecular tests and molecular characterisation, and the sections on virus isolation and on virus neutralisation, and added a section on the complement fixation test; developed justification tables for the ratings of the tests in Table 1 for each purpose.

The revised Chapter 3.6.9. *Equine rhinopneumonitis (infection with equid herpesvirus-1)* is presented as <u>Annex 13</u> for first-round comments.

3.8.1. Border disease: minor update, mainly of the taxonomy.

The revised Chapter 3.8.1. Border disease is presented as Annex 14 for first-round comments.

3.8.12. Sheep pox and goat pox: included fluorescent antibody testing and histopathology in Table 1 Test methods available for diagnosis of sheep pox and goat pox and their purpose; thoroughly updated the section on nucleic acid recognition methods, in particular the conventional and real-time PCR methods; clarified that ELISAs cannot discriminate between antibodies to different capripoxviruses.

The revised Chapter 3.8.12. Sheep pox and goat pox is presented as Annex 15 for first-round comments.

3.9.1. African swine fever (infection with African swine fever virus) (vaccine section only): a consultant working with vaccine developers, subject matter experts, representatives from the scientific community, regulatory authorities and WOAH Reference Laboratories had thoroughly updated Section C Requirements for vaccines on the manufacture of pure, potent, safe and efficacious vaccines for ASF⁶ including key vaccine performance and quality criteria.

The newly drafted section of Chapter 3.9.1. *African swine fever* (infection with African swine fever virus) (vaccine section only) is presented as <u>Annex 16</u> for first-round comments. An appendix to the draft chapter is also included for information comprising the results of the consultation, the key parameters, summaries of discussions, etc.

3.10.4. *Infection with* Campylobacter jejuni *and* C. coli: updated the taxonomy and the references and emphasised that *C. jejuni* and *C. coli* are of interest mainly from the point of view of food safety; reviewed the ratings of some of the tests in Table 1 *Test methods available for the diagnosis of* Campylobacter jejuni *and* C. coli *and their purpose* and developed justification tables for the ratings of the tests in Table 1 for each purpose. Updated the details in the methods for isolation and identification of the agent.

⁵ POC: point of care (tests)

⁶ ASF: African swine fever

The revised Chapter 3.10.4. *Infection with* Campylobacter jejuni *and* C. coli is presented as <u>Annex 17</u> for first-round comments.

3.10.8. *Toxoplasmosis:* thoroughly updated since last adopted in2017. In the interest of clarity, the text has not been marked up.

The revised Chapter 3.10.8. Toxoplasmosis is presented as <u>Annex 18</u> for first-round comments.

The batch of draft chapters also includes revised Chapter 3.1.5 *Crimean–Congo haemorrhagic fever* presented as <u>Annex 8</u> for first-round comments (see agenda item 3.1.1).

	Appendix	Chapter	
1.	4	1.1.5.	Quality management in veterinary testing laboratories
2.	5	1.1.9.	Tests for sterility and freedom from contamination of biological materials intended for veterinary use
3.	6	2.2.4.	Measurement uncertainty
4.	7	2.2.6.	Selection and use of reference samples and panels
5.	8	3.1.5.	Crimean–Congo haemorrhagic fever
6.	9	3.3.6.	Avian tuberculosis
7.	10	3.4.1.	Bovine anaplasmosis
8.	11	3.4.7.	Bovine viral diarrhoea
9.	12	3.4.12.	Lumpy skin disease (vaccine section only)
10.	13	3.6.9.	Equine rhinopneumonitis (infection with equid herpesvirus-1)
11.	14	3.8.1.	Border disease
12.	15	3.8.12.	Sheep pox and goat pox
13.	16	3.9.1.	African swine fever (vaccine section only)
14.	17	3.10.4.	Infection with Campylobacter jejuni and C. coli
15.	18	3.10.8.	Toxoplasmosis

5.2. Follow-up from September 2022: conclusions and recommendations from the WOAH *Scientific and Technical Review* issue on diagnostic test validation science

5.2.1. Progress on development of a validation report form for tests recommended in the *Terrestrial Manual*

At the meeting in February 2023, the Commission simplified and streamlined the validation report template for tests recommended in the *Terrestrial Manual* in the light of comments submitted by experts participating in a pilot scheme to test the template's suitability and usability. The new version of the document was shared again with those WOAH Reference Laboratories in the pilot scheme for further feedback on its utility.

At this meeting, the Commission agreed to make some final adjustments to the template, mainly to improve the introductory paragraph to better clarify the template's scope and purpose. It is important to emphasise that what is important is not how much data can be inserted under each heading in the template but rather are there data to demonstrate that the performance of the assay has been evaluated and that it is proven fit for the chosen purpose in Table 1.

Once the final version has been approved by all members of the Commission, the template will be made available on the WOAH website, and Reference Laboratory experts contributing to *Terrestrial Manual* chapters will be invited to use it to post their validation data online. In this way a repository of validation reports for recommended tests will be created for anyone seeking the validation data available for a given test. The Commission believes that this is an important advance, particularly for new technologies.

As stated in the September 2022 report, the template will also be used for experts requesting to add tests to the *Terrestrial Manual*.

5.2.2. Progress on development of a template for a new *Terrestrial Manual* section on the rationale behind the selection of tests included in Table 1. *Test methods available and their purpose*

At the meeting in February 2023, the Commission agreed to give the template developed to justify the selection of tests considered to be fit for purpose in Table 1, along with their rating, to all experts in the 2023/2024 review cycle when they were invited to update or draft a *Terrestrial Manual* chapter. Experts contributing to four disease chapters had used the template when updating their chapters. The Commission reviewed the submissions and found that this new section is an excellent addition to chapters in the *Terrestrial Manual* providing users with extremely useful information when they are deciding on tests to use for the various purposes while ensuring that the selection process is evidence-based. The new tables provide information on the type of sample, the test's accuracy, the test population, validation report if available, the advantages and disadvantages, and references. The Commission agreed that a link to the justification table would be added to each purpose in Table 1 so that users could click on the purpose for direct access to this information. It is hoped that more contributors to the *Terrestrial Manual* disease chapters will fill in the justification table template in the future.

5.3. Inclusion of videos on diagnostic techniques on the WOAH website disease portals: review of submitted videos

In February 2023, the Commission requested the Secretariat to contact the Reference Laboratories to ask if they have videos of diagnostic techniques they would like added to their chapter. At this meeting, the Commission reviewed the videos submitted. The Commission proposed that a repository of videos could be made available on the WOAH website. For the present, the videos would be from WOAH Reference Laboratories only and linked to diagnostic techniques for tests in the *Terrestrial Manual*. Each video would need to be accompanied by information on its contents, the publication associated with the technique, the software and equipment needed to perform the test and the language of the video. For a video to be added to the repository, the following criteria need to be met:

- i) Clarity: quality of the visual and audio aspects of the video should be high and the technique clearly explained for users;
- ii) Relevant to the Terrestrial Manual chapter;
- iii) Should not contain culturally inappropriate content;
- iv) The technique should be carried out in accordance with the *Terrestrial Manual* standards for quality technical proficiency, quality assurance and biosafety;
- v) Should not contain trade names or advertise certain kits or platform;
- vi) Should have a disclaimer stating that WOAH is not endorsing the contents of the video.

It should also be stressed that the videos are not WOAH standards. Any videos received would be first vetted by the Secretariat to ensure they fulfil the criteria. They will then be sent to the other Reference Laboratories for the disease in question before being submitted to members of the Commission for final approval to be added to the repository.

5.4. Further review of Chapter 1.1.6 Validation of diagnostic assays for infectious diseases of terrestrial animals

The Commission had received some last-minute changes to Chapter 1.1.6 Validation of diagnostic assays for *infectious diseases of terrestrial animals* after it was adopted in May 2023, expanding the information in the chapter on validation of POC tests. For this meeting, the Commission had also reviewed a first draft of a new chapter on the diagnostic validation of POC tests for WOAH-listed viral diseases using field samples and agreed that the chapter needed further development with input from the Reference Laboratory networks. The Commission decided to wait until the new chapter is finalised and adopted before addressing the proposed amendments to chapter 1.1.6.

5.5. Follow-up from February: need for a definition of latent period in the chapter on foot and mouth disease

See agenda item 3.3.2.

5.6. Review and alignment of the validity criteria of the PD₅₀ and PPG tests for foot and mouth disease between the *Terrestrial Manual* and the European Pharmacopoeia

The Commission was informed of the need to harmonise methods used for FMD vaccine potency assessment in the *Terrestrial Manual* and the European Pharmacopoeia (EuPh) to ensure that there is no variability in the interpretation of the methods in both texts. A WOAH Reference Laboratory expert is working with the EuPh to resolve this issue.

5.7. Purpose of the list of contributors and addresses in the Terrestrial Manual

The *Terrestrial Manual* includes a list of contributors to individual chapters with their professional addresses at the time of writing. The Commission believes the list gives credibility to the *Terrestrial Manual* and its inclusion contributes to the transparency of the standard-setting process. However, some of the experts on the current list have retired since contributing to chapters and their former employers do not allow them to use their former professional affiliation in the list. The Commission decided that in such cases to avoid including personal addresses, the name of the expert and their country of residence alone would be included. Experts who had changed employer but not retired, would be asked if the new employer would allow them to use that affiliation in the list.

5.8. Publication of Member comments and need to review the Commission's practice

As a result of the decision to publish Member comments (see agenda item 1.3.1), the Commission will need to change its current working practices. The first implemented change is to annex the chapters for comment to the reports.

5.9. Terrestrial Manual status: update on chapters selected for the 2024/2025 review cycle

The Commission examined the status of chapters that had previously been identified for update in the 2023/2024 review cycle but had not been received. The Commission decided to add to the list the remaining chapters that had last been updated in 2018, and to ask the contributors to chapters last adopted in 2019 if their chapters required an update. The Commission encouraged those Reference Laboratories with outstanding chapters to deliver by the deadline. The following chapters have been identified for update in 2024/2025 (year last adopted in brackets after the title).

- 1.1.2. Collection, submission and storage of diagnostic specimens (2013)
- 1.1.3. Transport of biological materials (2018)
- 1.1.4. Biosafety and biosecurity: Standard for managing biological risk in the veterinary laboratory and animal facilities (2015)
- 1.1.7. Standards for high throughput sequencing, bioinformatics and computational genomics (2016)
- 2.1.3. Managing biorisk: examples of aligning risk management strategies with assessed biorisks (2014)
- 2.1.1. Laboratory methodologies for bacterial antimicrobial susceptibility testing (2019)
- 2.2.1 Development and optimisation of antibody detection assays (2014)
- 2.2.2 Development and optimisation of antigen detection assays (2014)
- 2.2.3 Development and optimisation of nucleic acid detection assays (2014)
- 2.2.5 Statistical approaches to validation (2014)
- 2.2.7 Principles and methods for the validation of diagnostic tests for infectious diseases applicable to wildlife (2014)
- 2.2.8. Comparability of assays after minor changes in a validated test method (2016)
- 2.3.2. The role of official bodies in the international regulation of veterinary biologicals (2018)
- 2.3.3. Minimum requirements for the organisation and management of a vaccine manufacturing facility (2016)
- 2.3.5. Minimum requirements for aseptic production in vaccine manufacture (2016)
- 3.1.2. Aujeszky's disease (infection with Aujeszky's disease virus) (2018)
- 3.1.8. Foot and mouth disease (infection with foot and mouth disease virus) (2021)
- 3.1.9. Heartwater (2018)
- 3.1.14. New World screwworm (Cochliomyia hominivorax) and Old World screwworm (Chrysomya bezziana) (2019)
- 3.1.17. Q fever (2018)
- 3.1.20. Rinderpest (infection with rinderpest virus) (2018)
- 3.1.25. West Nile fever (2018)
 - Introductory note on bee diseases (2013)
- 3.2.4. Nosemosis of honey bees (2013)

- 3.2.5. Infestation of honey bees with Aethina tumida (small hive beetle) (2018)
- 3.2.6. Infestation of honey bees with Tropilaelaps spp. (2018)
- 3.3.1. Avian chlamydiosis (2018)
- 3.3.2. Avian infectious bronchitis (2018)
- 3.3.4. Avian influenza (including infection with high pathogenicity avian influenza viruses) (2021)
- 3.3.7. Duck virus enteritis (2018)
- 3.3.8. Duck virus hepatitis (2017)
- 3.3.11. Fowl typhoid and Pullorum disease (2018)
- 3.3.12. Infectious bursal disease (Gumboro disease) (2016)
- 3.4.9. Enzootic bovine leukosis (2018)
- 3.4.11. Infectious bovine rhinotracheitis/infectious pustular vulvovaginitis (2017)
- 3.4.13. Malignant catarrhal fever (2018)
- 3.4.15. Theileriosis in cattle (infection with Theileria annulata, T. orientalis and T. parva) (2018)
- 3.4.16. Trichomonosis (2018)
- 3.6.1. African horse sickness (infection with African horse sickness virus) (2019)
- 3.6.4. Epizootic lymphangitis (2018)
- 3.6.6. Equine infectious anaemia (2019)
- 3.6.7. Equine influenza (infection with equine influenza virus) (2019)
- 3.6.10. Equine viral arteritis (2013)
- 3.6.11. Glanders and melioidosis (2018)
- 3.8.2. Caprine arthritis/encephalitis and Maedi-visna (2017)
- 3.8.3. Contagious agalactia (2018)
- 3.8.5. Enzootic abortion of ewes (ovine chlamydiosis) (infection with Chlamydia abortus) (2018)
- 3.8.7. Ovine epididymitis (Brucella ovis) (2015)
- 3.8.11. Scrapie (2022)
- 3.8.12. Sheep pox and goat pox (2017) (vaccine section)
- 3.9.2. Atrophic rhinitis of swine (2018)
- 3.9.3. Classical swine fever (infection with classical swine fever virus) (2022: diagnostic tests section)
- 3.9.8. Swine vesicular disease (2018)
- 3.9.9. Teschovirus encephalomyelitis (2017)
- 3.9.10. Transmissible gastroenteritis (2008)
- 3.10.9. Verocytotoxogenic Escherichia coli (2008)

5.10. Update on WOAH Standards Online Navigation Tool Project

The WOAH Standards Department informed the Commission of a project to develop a new WOAH Standards Online Navigation Tool. This project is an initiative to change how WOAH Standards are displayed and made available to Members and other users. The project will enhance the display of the *Aquatic Code*, *Terrestrial Code*, *Aquatic Manual*, and *Terrestrial Manual* on the WOAH website. The project will also comprise a specific tool aimed at providing specific search functions for the visualisation of sanitary measures recommended for the international trade of commodities for terrestrial Animals. Also, the new tool is expected to simplify the annual updating process of the content of the Standards.

The project is aligned with the goals of the 7th Strategic Plan and will provide significant benefits to WOAH and its Members, including enhanced accessibility to WOAH Standards, efficiency in information retrieval, all the while supporting the implementation of WOAH Standards. The project will also bring gains to the Organisation itself, by

improving the efficiency of internal processes and the interoperability across various datasets related to WOAH Standards.

The Commission expressed interest and support for the project and recognised the importance of facilitating Members' access to achieve better understanding and use of WOAH Standards.

5.11. Terrestrial Manual chapters on non-listed diseases

At the meeting of the Bureaus of the Code and Biological Standards Commissions it was noted that some of the *Terrestrial Manual* chapters being updated were on non-listed or delisted diseases. The bureaus discussed the value of keeping and updating such chapters and whether they belong in a WOAH standard. It was noted that the Aquatic Animals Commission decided some time ago to maintain only chapters on listed diseases in the *Aquatic Manual*. The issue was also raised by Dr Arroyo in her welcome presentation. There are a total of 29 such chapters in the *Terrestrial Manual*. The Commission agreed that maintaining these chapters may not be the best use of resources. The Commission agreed to develop evidence-based criteria for keeping or deleting these chapters and to apply them at the meeting in February 2024. As Members could still need assistance with diagnosing and controlling these diseases, the Reference Laboratories and experts could be maintained if desired.

6. WOAH Reference Centres

6.1. Improving and automating the performance review for annual reports of Reference Laboratories using a risk-based approach

Every year, each Reference Laboratory submits an annual report of its activities. The questionnaire comprises 29 questions based on the ToR⁷ of the Reference Laboratories. Currently, each Commission member is tasked with evaluating approximately 40 laboratory reports annually, representing a heavy workload. During the Commission meeting in September 2022, the Members expressed the need to increase the efficiency of the evaluation process while reducing the workload.

At this meeting, the Secretariat proposed for the Commission's review a semi-automated system aimed at creating an effective method of performance assessment able to detect underperforming Reference Laboratories with a high level of sensitivity. Currently, all the annual reports are saved in a database; the system will use this database to generate an Excel file, reflecting an initial quantitative review of the reports. Using a risk-based approach and "weighting" the responses, the initial analysis can be further reviewed. The system will thereby categorise the Reference Laboratories as low risk or high risk of under-performance. This approach ensures that every report is screened uniformly to identify Reference Laboratories that need a more in-depth assessment by a Commission member by directing focus on potentially poor performers. In this way, the number of reports each BSC member evaluates could be reduced by 50 per cent while optimising the effort entailed. The system also ensures that each Reference Laboratory is evaluated at least once every 3 years.

The Commission recognised that some of the activities listed in the ToR for Reference Laboratories were more fundamental to the functioning of the laboratory than others and agreed to weight the responses in three categories: 'essential', 'core' and 'non-core' tasks. In this context, 'essential' signifies an activity that is mandatory to the functioning of the Reference Laboratory, 'core' denotes an activity that is optimal but its undertaking might depend on the Reference Laboratory's situation, and 'non-core' refers to activities that are of benefit to Members but not essential to the overall performance evaluation of the laboratory. The Commission also discussed the existing 'watch list' and the criteria for inclusion on it, such as those Reference Laboratories that did not submit a report or that had previously received a letter requesting clarification of aspects of their performance, as well as Reference Laboratories identified by the Commission based on professional judgement.

The Commission agreed to implement the system for the first time for the review of 2022 reports. The Secretariat will be responsible for distributing the reports identified by the system equally to each Commission member. An extraordinary meeting will be held in October 2023 to finalise the evaluations of the 2022 annual reports, assess the performance of the new system and communicate the findings with the network.

6.2. Applications for WOAH Reference Centre status

The Commission recommended acceptance of the following applications for WOAH Reference Centre status:

WOAH *Reference Laboratory Tularemia* Institute for Bacterial Infections and Zoonoses, Friedrich-Loeffler-Institut (FLI)

⁷ ToR: Terms of Reference

Federal Research Institute for Animal Health Naumburger Str. 96a – 07743 Jena, GERMANY Tel.: (49-3641) 804.2243; E-mail: <u>herbert.tomaso@fli.de</u> Website: <u>Institute of Bacterial Infections and Zoonoses (IBIZ): Friedrich-Loeffler-Institut (fli.de)</u> Designated expert: Dr Herbert Tomaso

WOAH Reference Laboratory for Rabies Laboratory for Emerging Viral Zoonoses, Research and Innovation Department Istituto Zooprofilattico Sperimentale Delle Venezie Viale dell'Università 10, 35020 Legnaro (PD) ITALY Tel.: (+39 049) 808.4385 E-mail: <u>pdebenedictis@izsvenezie.it</u> Website: <u>https://www.izsvenezie.it/</u> Designated expert: Dr Paola De Benedictis

WOAH *Reference Laboratory for Theileriosis* Epidemiology, Parasites and Vectors, Agricultural Research Council – Onderstepoort Veterinary Research, Onderstepoort 0110 SOUTH AFRICA Website: <u>Epidemiology, Parasites and Vectors (arc.agric.za)</u> Designated expert: Dr Barend Johannes Mans

WOAH Collaborating Center for Genome Monitoring of Swine Viral Disease National Bio and Agro-Defense Facility (NBAF) 1980 Denison Ave. Manhattan, KS 66502 UNITED STATES OF AMERICA Tel.: +1-785 477.9006 E-mail: Alfonso.Clavijo@usda.gov; Douglas.Gladue@usda.gov; Manuel.Borca@Usda.gov Website: National Bio and Agro-Defense Facility | USDA Contact Point: Douglas Gladue

Two applications had been received from the same institute in a Member in the Africa Region for Collaborating Centres, one for field epidemiology and one for risk assessment for animal health. Though the applicant is active and collaborates with institutes both within and outside the region, the Commission found that neither application provided sufficient details of the activities they propose to undertake and the services that they will provide to Members in the region. Neither did either application include a 5-year work plan. The Commission also noted that both applications chose Animal health management as the main focus area, with overlapping specialties in epidemiology, surveillance and risk assessment. The applicant will be requested to merge the two applications into one application for Collaborating Centre status and at the same time to strengthen the application by including details of proposed activities and outputs, and a 5-year work plan.

6.3. Changes of experts at WOAH Reference Centres

The Delegates of the Members concerned had submitted to WOAH the following nominations for changes of expert at WOAH Reference Laboratories. The Commission recommended their acceptance:

Rabbit haemorrhagic disease: Dr Patrizia Cavadini to replace Dr Lorenzo Capucci at the Istituto Zooprofilattico Sperimentale della Lombardia e dell'Emilia Romagna, Brescia, ITALY

Lumpy skin disease and *sheep pox and goat pox*: Dr Georgina Limon-Vega to replace Dr Pip Beard at the Pirbright Institute, UNITED KINGDOM

Salmonellosis (Salmonella spp.): Dr Francesca Martelli to replace Dr Rob Davies at the Animal and Plant Health Agency, UNITED KINGDOM

Foot and mouth disease:

Dr Vivian O'Donnell to replace Dr Consuelo Carrillo at the APHIS National Veterinary Services Laboratories, UNITED STATES

6.4. Review of new and pending applications for laboratory twinning

As of September 2023, 85 projects have been completed and 19 projects are underway. Of the completed projects, 15 Reference Laboratories and four Collaborating Centres have achieved WOAH designation status.

Two Laboratory Twinning project proposals were presented for the Commission's review:

- 1. **South Africa China (People's Rep. of)** for lumpy skin disease: the Commission supported the technical contents of this project proposal.
- 2. **South Africa Ghana** for African swine fever: the Commission supported the technical contents of this project proposal.

6.5. Analysis of the questionnaire sent to all Reference Laboratories

The questionnaire developed by the Commission was sent to all experts in charge of Reference Laboratories. Of 180 experts, 126 responses were received (70% response rate), representing all five WOAH regions. The survey was focused on the Reference Laboratory system and processes, covering topics not included in the annual report template. It placed particular emphasis on the point of contact between Reference Laboratories and WOAH, as well as the Commission. The survey provided an opportunity for experts to suggest changes and improvements to the system. It was divided into seven sections addressing various aspects of the Reference Laboratory system.

The survey results indicated overall satisfaction among Reference Laboratories. Multiple responses highlighted the need to simplify and make the application process, annual reports, and *Terrestrial Manual* chapter reviews more transparent. The Reference Laboratory experts also expressed a desire for closer contact with the Commission and WOAH staff, preferably in-person, to strengthen networking, align and harmonise efforts, and set joint goals.

The Commission discussed the benefits that accrue from face-to-face meetings of the designated experts responsible for the Reference Laboratories and noted that almost a decade had elapsed since the most recent global conference of WOAH Reference Centres in Incheon, Korea (Rep. of).

The survey will serve as a fundamental information source for enhancing processes related to Reference Laboratories. The questionnaire results will be shared with all Reference Laboratories as informational feedback as well as attached as <u>Annex 19</u> in this report.

6.6. Feedback from Laboratories that are not complying with the key ToR

The Commission was made aware of two Reference Laboratories that are not accredited to ISO 17025 or equivalent quality management system despite this being an essential requirement of all WOAH Reference Laboratories. The laboratories would be given 2 years to achieve accreditation and provide certificates or proof of equivalency.

6.7. Develop a plan of how to evaluate the progress made since the Collaborating Centres submitted their 5-years work plans

Collaborating Centres are designated for a period of 5 years, during which they adhere to a 5-year work plan, which was submitted at the beginning of the designation period. At the end of this period, the Director General sends a letter requesting a report of the achievements of the 5 years as outlined in the work plan. The Commission evaluates this report and decides if the Collaborating Centre's designation should be renewed or not based on their performance and the need to maintain a Centre for the specific topic.

This system of designating Collaborating Centres for a 5-year period was introduced in 2020 with the adoption of the SOPs (<u>Collaborating Centre - Procedures for Designation</u>). The first Centres to reach the end of their 5-year designation will do so at the end of 2024.

The Commission discussed the best approach to obtaining information on the activities of the Centres during their 5year designation period. The Commission agreed that a letter requesting a final report of their activities over the past 5-years in relation to the 5-year work plan originally submitted should be sent in the last quarter of the fifth year of the designation (September). At the February 2024 meeting, the Commission will develop a template for this final report along with performance criteria: it should include evidence of the Centre's impacts and achievements, benefits to the region, etc. The Centres will also be requested to submit the regular annual report, and both will be assessed by the Commission. The Commission will conduct an initial evaluation of the final reports, with the first results announced in the relevant February meeting. Those Collaborating Centres with an accepted final report will be notified after the February meeting that their designation can be renewed and will be asked to submit a new 5-year plan. Those Centres whose level of performance was not accepted will have a 6-month appeal period until the next Commission meeting in September at which their designation will be further evaluated and could lead to their withdrawal from the list.

6.8. Update on the three Reference Laboratory network (ASF, PPR⁸ and rabies)

African swine fever

The WOAH ASF Reference Laboratory network held regular virtual meetings to exchange scientific and technical expertise, including recent developments on ASF vaccines, and discussed activities in developing training programmes to assist at-risk countries, including the organisation of proficiency tests. The network is finalising a laboratory manual, including diagnostic algorithms to detect low virulent and novel emergent ASFV variants, and to explore user requirements on an open access information sharing platform for ASF virus genome sequence data. In the upcoming year, the Network plans to review its <u>overview</u> of ASF diagnostic tests for field application (POC tests).

Peste des petits ruminants

The WOAH PPR Reference Laboratory network continues to regularly update its <u>website</u> and organise activities in support of its members. Its second annual newsletter was circulated in July 2023 and published on its website. The newsletter provides an update on recent and forthcoming activities of the network as well as updates on activities of its members including other relevant networks (FAO/IAEA⁹ VETLAB Network). The main activities planned for 2023 include the organisation of a webinar in September 2023 on the harmonisation of PPR diagnostic methods through proficiency tests and the third workshop of the network, which will be held virtually in December 2023.

Rabies

The WOAH Reference Laboratory Network for Rabies (RABLAB) increased meetings to bimonthly in order to improve information sharing and alignment of activities between reference laboratories. Following recommendations from the RABLAB meeting in December 2022, *Istituto Zooprofilattico Sperimentale delle Venezie* (IZSVe), the FAO¹⁰ Reference Centre for Rabies, has joined RABLAB as an affiliate member, helping improve coordination of international efforts for rabies control.

Following recommendations from the Commission February 2023 meeting, the <u>WOAH website</u> has been updated to better highlight the role of RABLAB in supporting WOAH Members. RABLAB recommendations, including the 'Procedure for production of in-house (internal) positive control serum for rabies antibody testing' and the 'RABLAB statement on the use of commercial rapid immunochromatographic tests' can also be found online. To improve support for WOAH Members in their rabies control efforts, the RABLAB network has also developed a guidance document to assist WOAH Members in compiling information to support applications for WOAH endorsement of official control programmes for dog-mediated rabies and the network continues to participate in several Twinning projects to build laboratory capacity for rabies diagnosis. RABLAB members contributed to guidance on the role of oral rabies vaccination for dogs, and the development of a United Against Rabies Country Partnership programme.

There are ongoing discussions about the use of lateral flow devices (LFDs) for rabies virus detection. The RABLAB network has <u>developed guidance</u> for WOAH Members regarding this. The RABLAB network is continuing discussions with relevant manufacturers to explore how protocols and tests can be improved to support rabies surveillance.

6.9. Review of the current list of Main Focus Areas and Specialties

The Commission reconsidered the recommendation made at its February 2023 meeting to change the title of one of the Main Focus Areas for WOAH Collaborating Centres from "Wildlife Health and Biodiversity" to "Climate Change and Environment". The Commission noted the importance of maintaining the original title, Wildlife Health and Biodiversity, to reflect WOAH's commitment to the implementation of WOAH's Wildlife Health Framework, which included an objective to develop the network of Collaborating Centres specialising in wildlife health.

⁸ PPR: Peste des petits ruminants

⁹ IAEA: International Atomic Energy Agency

¹⁰ FAO: Food and Agriculture Organization of the United Nations

However, the Commission also identified that the text describing the Main Focus Area did not currently reflect a wildlife theme. The Commission therefore suggested that the descriptive text be amended.

The Commission also suggested integrating climate change and its impacts as specialties in three of the Main Focus Areas, namely Animal Health Management, Animal Production, and Wildlife Health and Biodiversity.

Following the meeting, the list was also reviewed and further amended by the Aquatic Animal Health Standards Commission. The updated document with the changes shown is available at <u>Annex 20</u>.

6.10. Clarify the role of the Contact Point in providing advice and services to WOAH Members

Each WOAH Collaborating Centre has one designated Contact Point to supervise the Centre's activities and act as the liaison between WOAH, the Commission, and WOAH Members. The Contact Point is often the Director of the institute that hosts the Centre, though in reality, other members of the Centre's personnel often take responsibility for responding to administrative issues or requests for assistance from Members on behalf of the official Contact Point. The Commission highlighted the need for a reliable and available point of contact. This was deemed essential for situations that required seamless communication and immediate assistance from a Reference Centre.

The official Contact Point will be asked if they are willing to continue to receive and re-direct requests or if they would prefer to nominate another staff member to be this first point of contact for their Centre for administrative purposes. This member will not replace the official Contact Point of the Centre but will facilitate all contact between the Centre and WOAH. This approach would enhance collaboration and also ensure timely response to the demands of those who depend on the services of the Reference Centre. The continual availability of a point of contact is considered essential for the success of these centres by the Commission.

7. Ad hoc Groups: Update on activities of past ad hoc Groups

7.1. Ad hoc Group on Replacement of the International Standard Bovine Tuberculin (ISBT) and Avian Tuberculin (ISAT)

In April 2023, the *ad hoc* Group on Replacement of the International Standard Bovine Tuberculin (ISBT) met virtually to continue discussions on the ISBT replacement process. Led by the UK Health Security Agency (UK HSA), a series of experiments were conducted to assess the potency of a newly manufactured PPD diagnostic reagent for *Mycobacterium bovis* in sensitised guinea-pigs. The goal was to establish the best method for evaluating the reagent's potency compared with standard PPD reagents in guinea-pigs sensitised with *M. bovis*. The experiments are currently in a new phase of testing with the aim of obtaining standardised results that will enable the assessment of the validity of the potential tuberculin replacement. The results are expected by the last quarter of the year.

The *ad hoc* Group also addressed the depletion of avian PPD reserves, providing comments on the technical criteria specifications and identifying potential manufacturers to launch the process of identifying a suitable replacement.

Following the update, the Commission noted the substantial commitment of time and resources being dedicated to the project to develop replacement tuberculin.

8. International Standardisation/Harmonisation

8.1. WOAH Register of diagnostic kits – update and review of new or renewed applications

The Secretariat for Registration of Diagnostic Kits (SRDK) informed the Commission of the status of ongoing applications. At present, there are 16 diagnostic test kits in the WOAH Register of Diagnostic Kits.

8.1.1. Ongoing application for Genelix[™] ASFV real-time PCR detection kit

The assessment of the application for *GenelixTM ASFV real-time PCR detection* kit (Sanigen) is under evaluation. No major issues have been identified in the 3rd Assessment Report. The applicant has submitted the revised dossier with the answers to the Panel of Experts (PoE). The review is under evaluation by PoE. The endorsement of the Review Panel Final Report and Validation Studies Abstract is planned to take place during the next Commission meeting (February 2024) and preparation of a new Resolution in May 2024 is underway.

8.1.2. Renewal of Avian Influenza Antibody test kit (registration number 20080203)

The applicant (BioChek [UK]) has provided the notification letter and declared that the test is still viable and that no changes have been introduced since the last renewal. The renewal procedure was initiated via the Reference Laboratories on 9 August 2023 according to the current Renewal Procedure. In expectation of a 5-year renewal of the Avian Influenza Antibody test kit, a new Resolution is being prepared for presentation in 2024.

8.1.3. Renewal of Newcastle Disease Antibody test kit (registration number 20140109)

The applicant (BioChek [UK]) has provided the notification letter and declared that the test is still viable and no that changes have been introduced since the last renewal. The renewal procedure was initiated via the Reference Laboratories on 9 August 2023. In expectation of a 5-year renewal of the Newcastle Disease Antibody test kit, a new Resolution is being prepared for presentation in 2024.

8.1.4. Update on WOAH SOP and application form

The SRDK updated the WOAH webpage on the registration of diagnostic kits to reflect the new WOAH logo and brand in the SOPs and application form.

8.2. Standardisation programme

8.2.1. Association française de normalisation: follow-up from February 2023

Following the February 2023 meeting, the Commission had informed AFNOR¹¹, on behalf of CEN/TC¹² 469, that proposals for amendments to the *Terrestrial Manual* should be submitted through the European Union representative and that CEN could not be involved directly in the work and discussions of the Commission. In response, the Chair of CEN/TC 469 clarified that CEN is not a European Union institution but a private international non-profit organisation recognised as the European standard-setting organisation. The Commission was informed that CEN/TC 469 does not hold the status of an international organisation having signed an agreement with WOAH, and therefore cannot submit proposals directly on WOAH standards. WOAH has signed an agreement to be a liaison organisation to participate in CEN/TC work. CEN/TC would need to investigate other means for submitting its proposals on introductory chapters of the *Terrestrial Manual*, through national Delegates or the European Commission.

8.2.2. Project to extend the list of WOAH-approved reference reagents: review of guidelines

The project to extend the list of WOAH-approved reference reagents is not progressing, with no WOAH Reference Laboratory submitting candidate reagents in the past few years. The main barrier is that the guidelines for antibody standards¹³, antigen standards¹⁴ and PCR assays¹⁵, are too demanding and thus costly to be an incentive for Reference Laboratories. As the Commission would like to maintain and extend the list, it was agreed to ask the disease-specific networks, namely ASF, FMD, rabies and PPR, to establish minimum criteria for the development of reference reagents so that the guidelines could be made more achievable while maintaining the quality of the reagents produced. At the next meeting in February 2024, the Commission will review the advice received, amend the guidelines and develop a list of priority reagents.

9. Follow-up from the General Session

9.1. Excerpt from the Final Report: comments from Members

At the General Session, a Member had informed the Assembly that a new vaccine for *Paenibacillus larvae* had been authorised for use in the country and requested that the vaccine be included in Chapter 3.2.2. *American foulbrood of honey bees (infection of honey bees with* Paenibacillus larvae). As this is a new diagnostic advance for this disease, the Commission would request more information from the Member in question and from the WOAH Reference Laboratories before proceeding with the request.

¹¹ AFNOR: Association française de normalisation

¹² CEN/TC: European Committee for Standardization/Technical Committee

^{13 &}lt;u>https://www.woah.org/app/uploads/2021/03/a-guideline-antibody-standards.pdf</u>

¹⁴ https://www.woah.org/app/uploads/2021/03/a-guideline-antiegen-standards.pdf

¹⁵ https://www.woah.org/app/uploads/2021/03/a-guideline-pcr-standards.pdf

Another Member drew the Commission's attention to the WOAH website, where previous amendments to the *Terrestrial Code* chapters are available in chronological order and can be accessed. The Delegate proposed that the same service be provided for the *Terrestrial Manual*.

The Commission discussed the value of this project and concluded that the Secretariat make available the chapters amended each year since 2013, based on the documents circulated with the February reports, i.e. the versions of the chapters proposed for adoption with all the amendments marked up. A table similar to the one for the *Terrestrial Code* would be created and updated on an annual basis, indicating the year (edition), revised chapter, Resolution, and the published chapter. Until 2012, the *Terrestrial Manual* was published every 4 years and no marked up chapters were circulated. These editions are available on the WOAH document portal.

9.2. The Animal Health Forum and the adopted Resolution on avian influenza

In light of the ongoing global avian influenza crisis, WOAH hosted its first <u>Animal Health Forum</u> (AHF), fully dedicated to the disease. The AHF served as a platform for international experts, the private sector and government representatives to engage in meaningful discussions about the current challenges and perspectives for combatting avian influenza. The AHF, held during 90th General Session, convened key stakeholders and WOAH Members to discuss how to minimise the impacts of avian influenza across sectors. Based on the Technical Item entitled <u>Strategic Challenges in the Global Control of High Pathogenicity Avian Influenza</u> presented at the event, participants discussed the impact of the disease, the fitness for purpose of existing prevention and control tools, international trade impact, and the necessity to enhance global coordination. WOAH Members adopted a <u>Resolution</u> that will serve as a basis for shaping future avian influenza control activities, while protecting wildlife, supporting the poultry industry and the continuity of trade. The Resolution notably underscores the importance of Members respecting and implementing WOAH international standards to effectively combat avian influenza.

The Commission was updated on the WOAH avian influenza resolution implementation framework that defines the activities, outputs and expected outcomes for the next 2 years to address the strategic challenges in the global control of HPAI that were discussed during the 90th WOAH General Session. This framework has been developed in consultation with the WOAH scientific network, the technical departments at headquarters and regional and subregional offices.

The Commission was briefed on the meeting of the WOAH Working Group on Wildlife which was held in June 2023 at the WOAH Headquarters. Given the global level of concern regarding avian influenza and its potential impact on wildlife, the Working Group prepared a brief statement on considerations associated with emergency vaccination of high conservation value species against avian influenza. The Working Group noted that the current *Terrestrial Code* chapter on avian influenza does not include information relevant to surveillance and reporting of highly pathogenic avian influenza (HPAI) in wild mammals, and to address this, the group submitted comments on the *Terrestrial Code* chapter for consideration. The Working Group discussed the available guidance on response to HPAI outbreaks in marine mammals and proposed next steps to draft a practical guide for field response to HPAI outbreaks in marine mammals, with a focus on biosecurity and sample collection and carcass disposal with the assistance of the WOAH Collaborating Centre Health of Marine Mammals.

The Commission was briefed on the update of the GF-TADs¹⁶ avian influenza strategy. which was last updated in 2008. The strategy is expected to be a short high-level document presenting the background, objectives, theory of change and the governance that rely on strong involvement at regional level. The strategy's purpose is to guide and create a global coordination framework to support regional and country action plans dedicated to the prevention and control of HPAI. The final version of the strategy is expected to be available by the end of the year.

The Commission commended the various activities presented to address the current global avian influenza crisis. In accordance with the <u>adopted Resolution</u>, the Commission agreed on the need to review the *Terrestrial Manual* chapter on avian influenza by Reference Centre experts to ensure the information is up to date with the latest science and fit for purpose. The Commission decided to pay special attention to new developments in diseases causing significant global impacts (e.g. avian influenza, African swine fever) and to prioritise those chapters in the work plan. To this end, the WOAH Reference Laboratories will be asked to review the current *Terrestrial Manual* chapter on avian influenza for important amendments as needed on an immediate basis. The amended chapter will go for one review round and be annexed to the February 2024 report for Member comments with the aim of proposing it for adoption in May 2024. The Commission emphasised the importance for Members to respect and implement of WOAH international standards to effectively combat the disease.

¹⁶ GF-TADs: Global Framework for the Progressive Control of Transboundary Animal Diseases

10. Conferences, Workshops, Meetings

10.1. Update on the WAVLD seminar in Lyon, France in 2023 and Commission involvement in future seminars

The 1-day WOAH Seminar had been held on 30 June 2023 during the ISWAVLD (International Symposium of the World Association of Veterinary Laboratory diagnosticians) in Lyon, France. ISWAVLD is an essential biennial meeting for all veterinarians, biologists, scientists and students who are involved to different degrees in research and laboratory diagnosis. The theme of the WOAH seminar was "Towards the Veterinary Diagnostics of the Future"; the One Health concept was a major theme throughout the agenda of the WAVLD symposium. The WOAH Seminar, a parallel daylong session, promoted the work of laboratory diagnostics, One Health in the laboratory setting, and capacity building and attracted a global audience of veterinary laboratory diagnosticians to advocate the needs of WOAH Members; the private sector was also present.

The WOAH Seminar had presentations on sustainable laboratory biosafety and biosecurity from recognised experts. Other topics included the challenges of sustainability of veterinary laboratory networks, supporting laboratory leadership in investment case development, countries perspectives of the benefits of the sustainable laboratories approach for national veterinary laboratory networks, update on the global laboratory leadership programme using a One Health approach, outcomes of the WOAH biosafety research roadmap, grand challenge for sustainable laboratories, safe and cost-effective protocols for shipment of suspected FMD samples for laboratory diagnostics, experience of the laboratory twinning project in ensuring high quality FMD vaccines, the Nagoya protocol and animal health, diagnostic kits and challenges to detection in wildlife. The WOAH Seminar ended with a panel discussion on laboratory equipment management solutions.

The response to the Seminar, as well as WOAH's engagement and involvement throughout the Symposium was positive from both the WAVLD Executive Board and the Symposium participants. The next ISWAVLD will occur in Calgary, Canada in June 2025. The Commission agreed to be fully involved by providing technical inputs for the theme and agenda of the WOAH Seminar. The Commission discussed topics that could be of interest for the next Seminar, including emerging transboundary animal diseases that are of importance to WOAH, novel technologies for animal disease diagnosis, updates on established successful networks such as ASF, FMD, rabies, PPR and avian influenza and the road to eradication of these diseases, how to integrate pen-side tests for disease diagnosis and information on validation techniques.

11. Matters of interest for consideration or information

11.1. Update on OFFLU

In response to the global surge in avian influenza outbreaks, the OFFLU ¹⁷ experts participated in various teleconferences and tripartite risk assessments and shared important data with the scientific community and policy makers. The network released scientific statements to address emerging animal influenza threats which include statement on high pathogenicity avian influenza caused by viruses of the H5N1 subtype, avian influenza events in mammals and cats.

The Commission was briefed on OFFLU's contribution to the <u>February 2023 WHO¹⁸</u> Consultation on the Genetic and antigenic characteristics of zoonotic influenza A viruses and development of candidate vaccine viruses for pandemic preparedness. The network provided sequence data gathered from laboratories in Europe, Asia, Africa, Oceania, and the Americas. For the <u>avian influenza report</u>, they collected 795 avian influenza virus sequences of H5, 34 of H7, and 305 of H9 subtypes. Additionally, for the <u>swine influenza report</u> they gathered 69 swine influenza virus sequences of H1 and 7 of H3 from WOAH Reference Centres, national veterinary laboratories, and research networks via the OFFLU network.

An <u>OFFLU avian influenza matching (AIM) initiative</u> to provide information on the real-time antigenic characteristics of contemporary avian influenza viruses is underway. This information will facilitate selection of appropriate vaccines for poultry and updating of poultry vaccine antigens in places where vaccines are being used. A report presenting the results of the pilot project will be made available to stakeholders in October 2023 and networking and expanding the geographical reach of this project with select partners is ongoing.

The OFFLU wildlife technical activity has been sharing data and offering support to countries and working closely with their local public health counterparts to track and monitor risk in response to the H5 mammalian spillovers experienced throughout 2022 and 2023. OFFLU experts released statements to update the H5N1 events in wild birds

¹⁷ OFFLU: Joint WOAH-FAO Network of Expertise on Animal Influenza

¹⁸ WHO: World Health Organization

in the Americas and the Europe and also contributed to the Scientific task force on avian influenza and wild birds statement.

The swine influenza virus expert group contributed invaluable information to the WHO vaccine composition meeting for influenza prepandemic preparedness.

The Expert Surveillance Panel of Equine Influenza comprising OFFLU and WHO influenza experts reviewed the recent equine Influenza virus activity in various countries, characteristics of the viruses isolated and provided vaccine recommendations.

11.2. Update on rinderpest

The Commission was updated on the rinderpest post-eradication activities. WOAH continues to work in partnership with FAO to reduce the RVCM¹⁹ holdings around the world, with the exception of diagnostic materials and vaccines. All seven FAO-WOAH designated RHFs²⁰ had their designations extended as a result of the site inspections of five of these facilities, conducted in 2022. The two remaining facilities will be inspected in 2024. WOAH will host a meeting on the 25th October 2023 to review inspection SOPs together with representatives of the Smallpox and Polio secretariats and the EuFMD²¹.

The third meeting of the RHF Network will take place at the WOAH HQ on the 6 and 7 December 2023. The Pirbright Institute will develop a competitive ELISA that only uses non-infectious material in 2024–2025, and this test will be validated by the members of the RHF Network. Currently, two members of the RHF Network are undertaking 'sequence and destroy' projects.

After the site inspection done by independent inspectors in October 2022, and a review of corrective actions by the FAO-WOAH Rinderpest Joint Advisory Committee (JAC) in September 2023, the Ethiopian National Veterinary Institute has been authorised to start rinderpest (RBOK) vaccine production to replenish the reserve at AU-PANVAC. There has not been any progress in sequestration or destruction of RVCM in the five Members that hold these materials outside of FAO-WOAH designated RHFs, in spite of several in-person and virtual discussions having been held.

Finally, the GF-TADs Management Committee is overseeing the review of the size and membership of the JAC done by the FAO-WOAH Rinderpest Secretariat. The JAC will have five members instead of seven and the new membership will be announced during the last trimester of 2023.

11.3. Update on VICH²² activities

The next VICH Steering Committee and Outreach Forum meeting will take place in Tokyo from 13 November to 16 November 2023. There are no vaccine- or biologicals-related VICH Guidelines or Concept Paper for comment at this time. VICH is undergoing a structural reorganisation to better address the needs of the Members who may have diverse expectations of the VICH Forum. This pre-meeting will be the second meeting to discuss the expectations for the future of the VICH Forum and the impact of past 5 years of being a member of the Forum. The agenda for this pre-meeting is being prepared by WOAH.

11.4. Update on the Grand Challenge for sustainable laboratories

The Commission was updated on plans to launch a Grand Challenge to identify solutions to improve the sustainability of laboratories. A short motion picture was shown that had been premiered at the Biological Weapons Convention in August 2023. The Commission was also updated on a study to assess the feasibility of running such a Grand Challenge, and informed that a high level meeting would take place in the UK in November 2023 with the aim of convening investment partners to take forward the initiative.

A member of the Commission had been part of the advisory board that reviewed the outcomes of the feasibility study. The member highlighted the importance of engaging local communities and experts in the development of solutions, and pointed out that technological leaps in diagnostic technology could reshape the current laboratory model.

¹⁹ RVCM: Rinderpest virus-containing materials

²⁰ Rinderpest holding facilities

²¹ EuFMD: European Commission for the Control of Foot-and-Mouth Disease

²² VICH is a trilateral (EU-Japan-USA) programme aimed at harmonising technical requirements for veterinary product registration. Its full title is the International Cooperation on Harmonisation of Technical Requirements for Registration of Veterinary Medicinal Products

11.5. Biosafety Research Roadmap

The Commission was updated on the Biosafety Research Roadmap announcing that it had delivered results in two areas: 1. A series of papers had been published in *Applied Biosafety* identifying gaps in the current evidence base for biosafety and biosecurity for 8 selected pathogens, and 2: a review of laboratory acquired infections and laboratory escapes over the past 20 years had been submitted for publication in *Lancet Microbe*. The outcomes of this work would be synthesised into a Chatham House policy paper calling for greater transparency in reporting laboratory accidents and for greater investment in the evidence base for biological risk management.

11.6. Collaboration with WOAH in the implementation of non-animal testing for veterinary vaccines batch release testing

The WOAH Antimicrobial Resistance and Veterinary Products Department requested the Commission to provide analysis and comments on WOAH's assessment on the future collaboration with the Animal Free Safety Assessment Collaboration (AFSA), HealthforAnimals and the International Alliance for Biological Standardization (IABS). The aim is to continue to collaborate on the implementation of non-animal testing for veterinary vaccines batch release testing in the *Terrestrial Manual*: "Implement 3Rs (*Replace, Reduce and Refine*) in Veterinary Vaccines Batch Release Testing". More specifically, the collaboration searches to:

- 1) Review relevant chapters of the *Terrestrial Manual*, add information on all available non-animal testing methodologies, and consider including additional guidance on these methodologies in the disease-specific chapters. The objective of this work is to support industry and regulatory stakeholders to use such new approaches.
- 2) Explore further collaboration on training and scientific education on consistency and non-animal safety and potency tests. The objective of this work is to improve access to and use of alternatives globally.

The collaboration started on 9 May 2022: two webinars, for the Americas and Asia-Pacific regions, and a face-to-face workshop were organised to determine what was needed to facilitate alternative methods validation, implementation and regulatory acceptance, including global harmonisation, of non-animal-based batch release testing for veterinary vaccines. The <u>report</u> from the webinars and workshop was published in the journal *Biologicals* in July 2023.

The Commission agreed on the value of continuing this collaboration, and requested further information on the 3Rs process to define and select alternative methods to ensure quality, validation and equivalency with existing methods.

11.7. Update on the development of guidelines for alternative strategies for the control of *Mycobacterium tuberculosis* complex infection in livestock

The Commission was informed of the development of Guidelines for Alternative Strategies for the Control of *Mycobacterium tuberculosis* complex infection in livestock. The guidelines are being developed by two consultants who began their work in May 2023. The objective is to illustrate selected tuberculosis control strategies that are flexible and adaptable to the changing conditions of real-world scenarios, taking into account socioeconomic and cultural settings.

The consultancy will identify existing science-based strategies for the control of *M. tuberculosis* complex infection in livestock other than slaughter and testing. The methodology will involve a literature review and the elicitation of expert opinion through surveys, interviews, and focus group discussions. This document will be reviewed upon an *ad hoc* Group expected to convene in January 2024.

The Commission emphasised the importance of presenting this document during the General Session in 2024.

11.8. Composition of the WOAH editorial board for the Scientific and Technical Review

The Head of the Publications Unit explained why a new Editorial Board was being established for WOAH's peer reviewed journal, the *Scientific and Technical Review*. Although the content is of high quality and robust editorial and reviewing processes are in place, the publication lacks governance to ensure its scientific credibility.

The Editorial Board will monitor and foster the quality and impact of the *Scientific and Technical Review* and will also advise on WOAH's overall publications strategy on request. The role of the Board will be mainly advisory but they will also participate occasionally in reviewing content and will attend two meetings per year.

WOAH would ask the Commission members if anybody would be willing to join the Editorial Board. Given that the mandate of the current Commission will end in May 2024, the term of the first nominated candidate will run until

September 2024. If no member of the Commission could commit to the role, then WOAH could ask other experts outside the Commission if they would like join the board.

The Commission agreed that the creation of a new Editorial Board would be a positive step forward for WOAH's publications.

11.9. Update on activities under the IHSC²³-WOAH collaboration agreement and consultancy project in Asia

The Commission was updated on the activities carried out between 2022 and 2023 under the collaboration agreement between WOAH and the IHSC. These activities include:

- The collaboration of the horse industry to improve WOAH Standards with regard to equine-related diseases;
- The development of diagnostic capabilities (i.e. glanders) and vaccines for horses (i.e. African horse sickness inactive vaccine and vaccine banks);
- The implementation of tools to facilitate the safe movements of competition horses.

The Commission was also briefed on the state of play and activities of two consultancy projects in South America and in Asia and the Pacific Region, particularly in terms of the assessment of laboratory capacity to diagnose equine diseases and the organisation several webinars on equine-related diseases to enhance preparedness.

The Commission commended the work conducted under this collaboration agreement and emphasised the importance of enhancing the diagnostic capabilities for equine encephalitides in the Americas.

11.10. Update on the virtual biobank project

The Commission was briefed on the reactivation of the Virtual Biobank project in April 2023, with the WOAH Collaborating Centre for Veterinary Biologicals Biobank, host by Istituto Zooprofilattico Sperimentale della Lombardia e dell'Emilia Romagna (IZSLER), Italy, taking the lead. The project was on hold due to the SARS-CoV-2 pandemic; the new timeline aims at completing the development of the web platform in 2024.

The project is currently in the development phase, with ongoing efforts to create the design a prototype. IZSLER's IT technical team is working in collaboration with the WOAH IT and Communication teams to build the project's website. Discussions are currently taking place between both IT teams regarding the web page server, where the architecture and design of the website will be stored.

To ensure that the project progresses, monthly meetings are being convened to deliberate on the project's advancement.

.../Annexes

²³ IHSC: International Horse Sports Confederation

Annex 1. Adopted Agenda

MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023

1. Welcome

- 1.1. Director General
- 1.2. Deputy Director General, International Standards and Science
- 1.3. Updates from the WOAH Headquarters

2. Adoption of Agenda

3. Collaboration with other Commissions

- 3.1. Horizontal issues among the Specialist Commissions
 - 3.1.1. Infection New World (*Cochliomyia hominivorax*) and Old World (*Chrysomya bezziana*) screwworms, and Crimean–Congo haemorrhagic fever (revisited)
- 3.2. Scientific Commission for Animal Diseases
 - 3.1.1. Nothing for this meeting.
- 3.3. Terrestrial Animal Health Standards Commission
 - 3.3.1. Updates from the February 2023 Code Commission meeting
 - 3.3.2. Biological Standards Commission's recommendations to the Terrestrial Animal Health Standards Commission
 - 3.3.3. Meeting of the Bureaus (7 September)
 - 3.3.4. Questions on Chapter 12.6 Infection with equine influenza virus
 - 3.3.5. Comments on Chapter.5.8. International transfer and laboratory containment of animal pathogenic agents
 - 3.3.6. Questions on Chapter 6.10. Responsible and prudent use of antimicrobial agents in veterinary medicine
- 3.4. Aquatic Animal Health Standards Commission
 - 3.4.1. Nothing for this meeting.

4. Work Programme

5. Manual of Diagnostic Tests and Vaccines for Terrestrial Animals

- 5.1. Review of draft chapters received for endorsement for circulation for first round Member comment
- 5.2. Follow-up from September 2022: conclusions and recommendations from the WOAH *Scientific and Technical Review* issue on diagnostic test validation science
 - 5.2.1. Progress on development of a validation report form for tests recommended in the Terrestrial Manual
 - 5.2.2. Progress on development of a template for a new *Terrestrial Manual* section on the rationale behind the selection of tests included in Table 1. *Test methods available and their purpose*
- 5.3. Inclusion of videos on diagnostic techniques on the WOAH website disease portals: review of submitted videos
- 5.4. Further review of Chapter 1.1.6 Validation of diagnostic assays for infectious diseases of terrestrial animals
- 5.5. Follow-up from February 2023: need for a definition of latent period in the chapter on foot and mouth disease
- 5.6. Review and alignment of the validity criteria of the PD₅₀ and PPG tests for foot and mouth disease between the *Terrestrial Manual* and the European Pharmacopoeia
- 5.7. Purpose of the list of contributors and addresses in the *Terrestrial Manual*
- 5.8. Publication of Member comments and need to review the Commission's practice
- 5.9. Terrestrial Manual status: update on chapters selected for the 2024/2025 review cycle
- 5.10. Update on WOAH Standards Online Navigation Tool Project
- 5.11. Terrestrial *Manual* chapters on non-listed diseases

6. WOAH Reference Centres

- 6.1. Improving and automating the performance review for annual reports of Reference Laboratories using a riskbased approach
- 6.2. Applications for WOAH Reference Centre status
- 6.3. Changes of experts at WOAH Reference Centres
- 6.4. Review of new and pending applications for laboratory twinning
- 6.5. Analysis of the questionnaire sent to all Reference Laboratories Reference Laboratories – Implementation of the SOPs
- 6.6. Feedback from Laboratories that are not complying with the key ToRs: Collaborating Centres – Implementation of the SOPs
- 6.7. Develop a plan of how to evaluate the progress made since Centres submitted their 5-year work plans Reference Centre networks
- 6.8. Update on the three Reference Laboratory networks (African swine fever, peste des petits ruminants and rabies)
- 6.9. Review of the current list of Main Focus Areas and Specialties
- 6.10. Clarify the role of the Contact Point in providing advice and services to WOAH Members

7. Ad hoc Groups: Update on activities of past ad hoc Groups

7.1. *Ad hoc* Group on Replacement of the International Standard Bovine Tuberculin (ISBT) and Avian Tuberculin (ISAT)

8. International Standardisation/Harmonisation

- 8.1. WOAH Register of diagnostic kits: update and review of new or renewed applications
 - 8.1.1. Ongoing application for Genelix™ ASFV real-time PCR detection kit
 - 8.1.2. Renewal of Avian Influenza Antibody test kit (registration number 20080203)
 - 8.1.3. Renewal of Newcastle Disease Antibody test kit (registration number 20140109)
 - 8.1.4. Update on WOAH SOP and application form
- 8.2. Standardisation programme
 - 8.2.1. Association française de normalisation: follow-up from February 2023
 - 8.2.2. Project to extend the list of WOAH approved reference reagents: review of guidelines

9. Follow-up from the General Session

- 9.1. Excerpt from the Final Report: comments from Members
- 9.2. The Animal Health Forum and the adopted Resolution on avian influenza

10. Conferences, Workshops, Meetings

10.1. Update on the WAVLD seminar in Lyon, France in 2023 and Commission involvement in future seminars

11. Matters of interest for consideration or information

- 11.1. Update on OFFLU
- 11.2. Update on rinderpest
- 11.3. Update on VICH activities
- 11.4. Update on the Grand Challenge for sustainable laboratories
- 11.5. Biosafety Research Roadmap
- 11.6. Collaboration with WOAH in the implementation of non-animal testing for veterinary vaccines batch release testing
- 11.7. Update on the development of guidelines for alternative strategies for the control of *Mycobacterium tuberculosis* complex infection in livestock
- 11.8 Composition of the WOAH editorial board for the Scientific and Technical Review
- 11.9. Update on activities under the IHSC²⁴-WOAH collaboration agreement and consultancy project in Asia
- 11.10. Update on the virtual biobank project

²⁴ IHSC: International Horse Sports Confederation

Annex 2. List of Participants

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023

MEMBERS OF THE COMMISSION

Prof. Emmanuel Couacy-Hymann

(President) Professor of Virology CNRA/LIRED Abidjan CÔTE D'IVOIRE

Dr Joseph S. O'Keefe (Member) Head of Animal Health Laboratory Ministry for Primary Industries Upper Hutt NEW ZEALAND

Prof. Ann Cullinane (Vice-President) Head of Virology Unit Irish Equine Centre Naas IRELAND

Dr Satoko Kawaji (Member) Principal Scientist National Institute of Animal Health Naro JAPAN

Dr John Pasick (Vice-President) Formerly National Centre for Foreign Animal Disease Winnipeg CANADA

Prof. Chris Oura (Member) Professor of Veterinary Virology The University of the West Indies St-Augustine TRINIDAD AND TOBAGO

CONSULTANT EDITOR OF THE TERRESTRIAL MANUAL

Dr Steven Edwards c/o WOAH, Paris, FRANCE

WOAH HEADQUARTERS

Dr Gregorio Torres Head Science Department

Dr Charmaine Chng Deputy Head Science Department **Ms Sara Linnane** Scientific Officer Science Department

Dr Mariana Delgado Scientific Secretariat Officer Science Department **Dr Gounalan Pavade** Scientific Coordinator Science Department

Annex 3. Work Programme for the WOAH Biological Standards Commission

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023

Subject	Issue	Status and Action
	 Circulate the chapters approved by the BSC to Members for first-round comment 	October 2023
	 Remind authors of the chapters identified previously for update but not yet received and invite authors of chapters newly identified for update 	On-going
	3) Create a database of validation reports to be published on the WOAH Website for tests recommended in the <i>Terrestrial Manual</i>	On-going
	a) Finalise the template for the validation report for current and future tests recommended in the <i>Terrestrial Manual</i> and make available to Reference Laboratory experts for submission for new or existing tests	December 2023
Updating the Terrestrial Manual	4) Add a new section to the disease-specific chapters to describe the rationale behind the selection of tests for different purposes given in Table 1 <i>Test methods</i> <i>available and their purpose</i> and an explanation for their score. Subsequently, add links to the validation reports (point 3 above)	Accomplished
	 a) Send the template for this new section to the experts updating Manual chapters asking they use it or provide a justification in an alternative format of their choice 	Accomplished
	5) Ask Reference Centres to provide links to suitable instructional videos to be added to the end of the disease-specific chapters. Videos to be reviewed by the BSC when the chapter is up for review	On-going
	 Develop criteria for removing chapters for nonlisted diseases and assess those chapters against the criteria 	For February
	 Review new developments in diseases causing significant global impacts (e.g. avian influenza, African swine fever) and prioritise those chapters 	On-going
	1) Implementation of the adopted SOPs:	
Collaborating Centres	 a) Develop a template for the Collaborating Centres for the report of their assessment of their performance in the past 5-years to be compared with the 5-year work plan 	For September 2024
control control	 Review the designations of those Centres that completed 5 years 	February 2025
	 Ask the Contact Point to nominate a first point of contact to address requests for assistance, enquires, etc. on behalf of the Centre 	October/November 2023

Subject	Issue	Status and Action
	1) Put under-performing labs on watch list	On going
	 Implement the new system for evaluating annual reports and provide list of assigned reports to BSC members 	For October 2023
Reference Laboratories	3) Send feedback to the Reference Laboratory network on the questionnaire	October 2023
	4) Develop a concept note for a global conference of Reference Centres	February 2024
	5) Explore enhancements to the annual report process: the possibility of filling in the annual report template throughout the year	On-going
Reference Centre Networks	 Follow up with the three newly launched Reference Laboratory networks (ASF, PPR and rabies) 	On-going
	1) Project to extend the list of WOAH-approved reference reagents	On-going
Standardisation/ Harmonisation	 Ask the networks to review the template and three guidelines for approved reagents with the view to making the procedure less stringent so more Reference Laboratories may apply for recognition 	For February 2024
	 Project to develop Replacement International Standard Bovine and Avian Tuberculin: finalise report and propose for adoption 	On-going
Ad hoc Groups	1) Ad hoc Group on Sustainable Laboratories	On-going
Projects	1) Veterinary Biobanking (project)	On-going
Conferences,	1) Biosafety research roadmap	On-going
Workshops and Meetings with participation by BSC Members	2) ISWAVLD OIE Seminar: develop a theme and programme and speakers	June 2025 in Canada
Performance	 Engage with the ongoing processes around performance issues with Reference Laboratories 	On-going
Develop laboratory standards for emerging diseases	1) Discuss the <i>Terrestrial Code</i> chapter once adopted with the aim of introducing a corresponding chapter for the <i>Terrestrial Manual</i>	After May 2024
Case definitions	1) Follow up the implementation of the SOPs for case definitions	On-going

1	Annex 4. Item 5.1. – Chapter 1.1.5. Quality management in veterinary testing laboratories
2	MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION
3	Paris, 4–8 September 2023
4	
5	
6	CHAPTER 1.1.5.
7 8	QUALITY MANAGEMENT IN VETERINARY TESTING LABORATORIES

SUMMARY

10 Valid laboratory results are essential for diagnosis, surveillance, and trade. Such results are achieved by the use assured through implementation of good a management practices, valid system that supports accurate 11 and consistent test and calibration methods, proper techniques, quality control and quality assurance, all 12 working together within a quality management system. Laboratory quality management includes technical, 13 14 managerial, and operational elements of testing performing, interpreting and the interpretation of reporting test results. A quality management system enables the laboratory to demonstrate both competency and an 15 16 ability to generate consistent technically valid results that meet the needs of its customers. The need for Mutual recognition and acceptance of test results for international trade, and the acceptance accreditation 17 of tests to international standards such as ISO/IEC²⁵ 17025:2005 (General Requirements for the 18 19 Competence of Testing and Calibration Laboratories) (ISO/IEC, 2005-2017b) requires good suitable laboratory quality management systems. This chapter is not intended to reiterate the requirements of 20 21 ISO/IEC 17025, nor has it been endorsed by accreditation bodies. Rather, it outlines the important issues and considerations a laboratory should address in the design and maintenance of its quality management 22 23 system, whether or not it has been formally accredited regardless of formal accreditation status. Chapter 24 1.1.1 Management of veterinary diagnostic laboratories gives an introduction to veterinary diagnostic 25 laboratories introduces the components of governance and management of veterinary laboratories that are necessary for the effective delivery of diagnostic services, and highlights the critical elements that should be 26 established as minimum requirements. 27

28 29

9

A. KEY CONSIDERATIONS FOR THE DESIGN AND MAINTENANCE OF A LABORATORY QUALITY MANAGEMENT SYSTEM

To ensure that the quality management system is appropriate and effective, the design must be carefully thought out <u>planned</u> and, where accreditation is sought, must address all criteria of the appropriate quality standard. The major categories of considerations and the <u>their associated</u> key issues and activities within each of these categories are outlined in the following eight sections of this chapter.

²⁵ ISO/IEC: International Organization for Standardization/International Electrochemical Commission.

1. The work, responsibilities, and goals of the laboratory

- Many factors affect the necessary elements and requirements of a quality management system. These factors include. including:
- 37 i) Type of testing done-performed, e.g. research versus diagnostic work;
- ii) Purpose and requirements of the-test results, e.g. for-import-or-<u>/</u>export quarantine testing, surveillance, emergency
 disease exclusion, declaration of freedom from disease post-outbreak;
- iii) Potential impact of a questionable-or, erroneous or unfavourable result, e.g. detection of foot and mouth disease
 (FMD) in an FMD-free country;
- 42 iv) The tolerance level of Risk and liability tolerance, e.g. vaccination vs-versus culling-or /slaughter;
- v) Customer needs (requirements, e.g. sensitivity and specificity of the test method, cost, turnaround time, strain or genotype level of characterisation), e.g. for surveillance, or declaration of freedom after outbreak;
- vi) The role of the laboratory <u>Role</u> in legal work or in regulatory programmes, e.g. for disease eradication and declaration
 of disease freedom to the WOAH;
- vii) The role of the laboratory <u>Role</u> in assisting with, confirming, or overseeing the work of other laboratories (e.g. as a reference laboratory);
- 49 viii) Business goals-of the laboratory, including the need for any third-party recognition or accreditation.

50 **2.** Standards, guides, and references

51 The laboratory should choose reputable and accepted follow globally recognised standards and guides to assist in designing the quality management system. For laboratories seeking accreditation formal recognition of testing competency, 52 and for all WOAH Reference Laboratories, the use of ISO/IEC 17025 (ISO/IEC, 2005-2017b) or equivalent will be is 53 54 essential. This standard includes specifies managerial and technical requirements and accredited laboratories that are compliant are regarded as competent. Further information on standards may be obtained from the national standards body 55 56 of each country, from the International Laboratory Accreditation Cooperation (ILAC)²⁶, and from accreditation bodies, e.g. the National Association of Testing Authorities (NATA), Australia, the United Kingdom Accreditation Service (UKAS), the 57 American Association for Laboratory Accreditation (A2LA), etc. Technical and international organisations such as AOAC 58 International (The Scientific Association Dedicated to Analytical Excellence; formerly the Association of Official Analytical 59 Chemists) and the International Organization for Standardization (ISO) publish useful references, guides, application 60 documents and standards that supplement the general requirements of ISO/IEC 17025. Other relevant documents may 61 62 include guides and application documents providing interpretative criteria and recommendations for the application of 63 ISO/IEC 17025 in the field of veterinary testing for both applicant and accredited facilities, e.g. 64 au/phocadownload/spe Animal-Health-ISO-IEC-17025 https://www.nata guidance/animal-health Appendix.pdf -https://nata.com.au/files/2021/05/Animal-Health-ISO-IEC-17025-Appendix-effective-March2021.pdf; 65 66 Newberry & Colling, 2021.

The ISO International Standard 9001 (ISO, 2015), is a certification standard specifies the requirements for quality management systems and while it may be a useful supplement framework to <u>a-underpin a laboratory</u> quality system, <u>fulfilment of</u> its requirements does not necessarily ensure or imply assure technical competence (in the areas listed in Section 3 *Accreditation*). <u>Conformance to the requirements of</u> ISO 9001 is assessed by a certification body that is accredited to <u>undertake such assessments</u> by the national accreditation body <u>to undertake such assessments</u>. When a laboratory meets the requirements of ISO 9001, the term *registration* or *certification* is used to indicate conformity, not *accreditation*.

 With the advent of stronger alliances between medical and veterinary diagnostic testing under initiatives such as "One Health", some laboratories may wish to choose to follow other ISO standards such as ISO 15189 Medical Laboratories – Requirements for Quality and Competence (ISO/IEC, 2012), which include 2022), for testing of human samples, e.g. for zoonotic diseases. It should be noted that for veterinary laboratories, limited availability of suitable material may render validation difficult; under these circumstances it is necessary to highlight the limited validation status when reporting results and their interpretation (Stevenson *et al.*, 2021).

80 3. Accreditation

81 If the laboratory decides to proceed with formal recognition of its-a laboratory's quality management system and testing,

82 then is sought, third party verification of its conformity with the selected standard(s) will be is necessary. ILAC has published

specific requirements and guides for laboratories and accreditation bodies. Under the ILAC system, ISO/IEC 17025 is to

26 ILAC: The ILAC Secretariat, PO Box 7507, Silverwater, NSW 2128, Australia; http://ilac.org/

- 84 be used for laboratory accreditation of testing or calibration activities. Definitions regarding laboratory accreditation may
- be found in ISO/IEC International Standard 17000: Conformity Assessment Vocabulary and General Principles (ISO/IEC,
- 86 2004a-2020). Accreditation is tied to dependent on demonstrated competence, which is encompasses significantly more
- 87 than having and following documented procedures. Providing a competent and customer-oriented service also means that
- 88 the laboratory <u>requires</u>:
- 89 i) Adequate facilities and environmental controls;
- Has Appropriately qualified and trained personnel with a depth of technical knowledge commensurate with
 appropriate level of authority;
- 92 iii) Has appropriate Equipment with planned that is appropriately verified and managed in accordance with the relevant
 93 maintenance and calibration schedule;
- 94 iv) Has adequate facilities and environmental control;
- 95 v) Has procedures and specifications that ensure accurate and reliable results;
- 96 vi) Implements continual improvements in testing and quality management;
- 97 vii) Can assess the need for and implement appropriate corrective or preventive actions, e.g. customer satisfaction;
- 98 viii) Accurately assesses and controls uncertainty in testing;
- 99 ix) Appropriate sample and materials management processes;
- x) Has-Technically valid and validated test methods, procedures and specifications-that are documented in accordance
 with the requirements of the applicable standard or guidelines, e.g. Chapter 1.1.6 *Principles and methods of validation* of diagnostic assays for infectious diseases and chapters 2.2.1 to 2.2.8 *Recommendations for validation of diagnostic* tests and Special Issue of the *Scientific and Technical Review* (2021)²⁷;
- 104 xi) Demonstrates Demonstrable proficiency in the <u>applicable</u> test methods used (e.g. by <u>regular</u> participation in 105 proficiency tests on a regular basis testing schemes);
- 106 xii) Accurate assessment and control of the measurement of uncertainty in testing:
- xiii) <u>Good documentation practices, e.g. ALCOA+ principles (i.e. Attributable, Legible, Contemporaneous, Original,</u>
 <u>Accurate, Complete, Consistent, Enduring, Available);</u>
- xiv) <u>Non-conformance management process, including detection, reporting, risk-assessment and implementation of</u>
 <u>effective corrective and preventive actions;</u>
- 111 xv) Complaints management;
- 112 xvi) Adequate control of data and information;
- 113 xvii) Appropriate reporting and approval process:
- 114 xviii) <u>Culture of continual improvement.</u>
- 115 xix) Has demonstrable competence to generate technically valid results.

4. Selection of an accreditation body

117 To facilitate the acceptance of the laboratory's test results for trade, the accreditation standard used must be recognised 118 by the international community and the accreditation body recognised as competent to accredit laboratories. Programmes 119 for the recognition of accreditation bodies are, in the ILAC scheme, based on the requirements of ISO/IEC International 120 Standard 17011: <u>Conformity Assessment –</u> General Requirements for Accreditation Bodies Accrediting Conformity 121 Assessment Bodies (ISO/IEC, <u>2004b-2017a</u>). Information on recognised accreditation bodies may be obtained from the 122 organisations that recognise them, such as the Asia-Pacific Accreditation Cooperation (APAC), the Inter-American 123 Accreditation Cooperation (IAAC), and the European Co-operation for Accreditation (EA).

Accreditation bodies may also be signatory to the ILAC and regional (e.g. APAC) mutual recognition arrangements (MRAs).
 These MRAs are designed to reduce technical barriers to trade and further facilitate the acceptance of a laboratory's test
 results in foreign markets. Further information on the ILAC MRA may be obtained from the www.ilac.org.

127 5. Determination of the scope of the quality management system or of the laboratory's 128 accreditation

²⁷ Available at: https://doc.woah.org/dyn/portal/index.xhtml?page=alo&aloId=41245

- 129 The-scope of the quality management system should cover all areas of activity affecting all include all activities that impact
- 130 testing that is done at performed by the laboratory. Whilst only accredited laboratories are obliged to meet the requirements
- 131 of the relevant standard as detailed below, these, the guiding principles should be considered best practise and are relevant

132 to all testing laboratories.

133 Laboratories accredited A laboratory's accreditation to ISO/IEC 17025 have includes a specific list of those accredited tests 134 that are accredited, called, referred to as the schedule or scope of accreditation or the scope. Veterinary testing facilities include government and private facilities, veterinary practices, university veterinary schools, and other laboratories for the 135 testing of animals and animal products for the diagnosis, monitoring and treatment of disease. In principle, if new testing 136 methods are introduced these must be assessed and accredited before they can be added to the scope, however a flexible 137 scope can be implemented that assesses the laboratory as competent to add tests to scope, which are then formally added 138 at the next accreditation visit. The quality management system should ideally cover all areas of activity affecting all testing 139 that is done at the laboratory. However, it is up to the laboratory to decide which tests are to be accredited and included in 140 the scope. If an accredited laboratory also offers unaccredited non-accredited tests, these must be clearly indicated as 141 such on any reports that claim or make-reference to-accreditation. Factors It is ultimately the decision of the laboratory to 142 decide which tests require inclusion in the scope of accreditation, and factors that might affect the laboratory's choice of 143 144 tests for scope of accreditation this decision include:

- 145 i) The impact of initial accreditation on resources within a given deadline;
- 146 ii) Associated risks and opportunities;
- 147 iii) Initial investment required (e.g. time, resources);
- 148 iv) A-Contractual requirement for accredited testing (e.g. for international trade, research projects);
- 149 v) The-Importance of the test and the potential impact of an incorrect result;
- 150 vi) The cost of maintaining an accredited test <u>versus frequency of use;</u>
- 151 vii) Availability of personnel, facilities and equipment;
- viii) Availability of <u>appropriate materials and</u> reference standards (e.g. standardised-reagents, internal quality control samples-controls, reference cultures)-and
- 154 ix) <u>Access to</u> proficiency testing schemes;
- 155 x) The quality assurance <u>control processes</u> necessary for materials, reagents and media;
- xi) The validation <u>status, e.g. access to field samples from infected and non-infected animals</u>, technical complexity and
 reliability of the test method;
- 158 xii) The Potential for subcontracting of accredited tests.

6. Quality assurance, quality control and proficiency testing

160 Quality assurance (QA) is the <u>part-element</u> of quality management focused on providing confidence that <u>quality_defined</u> 161 requirements <u>will be are</u> fulfilled. The requirements may be internal or defined in an accreditation or certification standard. 162 QA is process-oriented and <u>ensures-provides</u> the right things are being done in the right way appropriate inputs to prevent 163 problems arising.

Quality control (QC) is the systematic and planned monitoring of outputs to ensure the minimum levels of quality requirements have been met. For a testing laboratory, this is to ensure test processes ensures tests are working correctly performing consistently and reliably, and results are within the expected acceptable parameters and limits. QC is test orientated and ensures the results are as expected_oriented and ensures detection of any problems that arise.

Proficiency testing (PT), sometimes referred to as external quality assurance or <u>(EQA)</u>, is the <u>determination assessment</u> of a laboratory's performance <u>by when</u> testing <u>a standardised panel of</u> specimens of undisclosed content. Ideally, PT schemes should be <u>run managed</u> by an external independent provider. Participation in proficiency testing <u>schemes</u> enables the laboratory to assess and demonstrate <u>the their testing</u> reliability <u>of results by in</u> comparison with those from other participating laboratories.

All laboratories should, where possible, participate in external proficiency testing schemes appropriate to their testing. Participation the suite of tests provided; participation in such schemes is a requirement for accredited laboratories. This provides an independent assessment of the testing methods used and as well as the level of staff competence. If such schemes are not available, valid alternatives may be used, such as ring trials organised by reference laboratories, interlaboratory testing, use of certified reference materials or internal quality control samples, replicate testing using the same or different methods, retesting of retained items, and or correlation of results for different characteristics of a specimen. Providers and operators of proficiency testing programmes should be accredited to ISO/IEC 17043 – Conformity Assessment – General Requirements for Proficiency Testing (ISO/IEC, 2010).

Proficiency testing material from accredited providers has been is well characterised and any spare material, once the proficiency testing has been completed, can be useful to demonstrate staff competence or for test validation. Information about selection and use of reference samples and panels is available in Chapter 2.2.6 *Selection and use of reference samples and panels*. Proficiency testing and reproducibility scenarios are described by Johnson & Cabuang (2021) and Waugh & Clark (2021), respectively.

186 **7. Test methods**

ISO/IEC 17025 requires the use of appropriate test methods and has requirements for their selection, development, and
 validation to show-demonstrate fitness for purpose.

This *Terrestrial Manual* provides recommendations on the selection of test methods for trade, diagnostic and surveillance purposes in the chapters on specific diseases. Disease-specific chapters include, or will include in the near future, a table of the tests available for the disease, graded against the test's fitness for purpose; these purposes are defined in the WOAH Validation Template (chapter 1.1.6), which identifies six main purposes for which diagnostic tests may be carried out. The table is intended to be as a general guide to test application.—: the fact that a test is recommended does not necessarily mean that a laboratory is competent to perform it. The laboratory quality system should incorporate provision of evidence of competency.

In the-veterinary <u>profession_laboratories</u>, other standard methods (published in international, regional, or national standards) or fully validated methods (having undergone a full collaborative study and that are published or issued by an authoritative technical body such as the AOAC International) may be preferable to use, but may_not be_available. Many veterinary laboratories develop or modify methods, and most laboratories have test systems that use non-standard methods, or a combination of standard and non-standard methods. In veterinary laboratories, even with the use of standard and methods, some in-house evaluation, optimisation, or validation is generally must be done required to ensure valid results.

202 Customers and laboratory staff must have a clear understanding of the performance characteristics of the test, and 203 customers should be informed if the method is non-standard. Many veterinary testing laboratories will therefore need to 204 demonstrate competence in the development, adaptation, <u>verification</u> and validation of test methods.

This *Terrestrial Manual* provides more detailed and specific guidance on test selection, optimisation, standardisation, and validation in chapter 1.1.6. Chapter 1.1.6 refers to chapters 2.2.1–2.2.8 *Recommendations for validation of diagnostic tests* that deal with the development and optimisation of fundamentally different assays such as antibody, antigen and nucleic acid detections tests, measurement uncertainty, statistical approaches to test validation, selection and use of reference samples and panels, validation of diagnostic tests for wildlife, and comparability experiments after changes in a validated test method.

211 The following are key test method issues for those involved in the quality management of the laboratory.

212 **7.1.** Selection of the test method

216

217

218 219

220

221

222 223

225

Valid results begin with the selection of a test method that meets the needs of the laboratory's customers in
 addressing their specific requirements (fitness for purpose). Some issues relate directly to the laboratory, others to
 the customer.

- 7.1.1. Considerations for the selection of a test method
 - i) International acceptance;
 - ii) Scientific acceptance;
 - iii) Appropriate or current technology;
 - iv) Suitable performance characteristics (e.g. analytical and diagnostic sensitivity and specificity, repeatability, reproducibility, isolation rate, limits of detection, precision, trueness, and uncertainty);
 - v) Suitability of the test in the species and population of interest;
- vi) Sample type (e.g. serum, tissue, milk) and its expected quality or state on arrival at the laboratory;
 - vii) Test target (e.g. antibody, antigen, live pathogen, nucleic acid sequence);

226	viii) Test turnaround time;
227	ix) Resources and time available for development, adaptation, evaluation;
228	x) Intended use (e.g. export, import, surveillance, screening, diagnostic, confirmatory);
229	xi) Safety factors and biocontainment requirements;
230	xii) Customer expectations;
231	xiii) Throughput of test Sample numbers and required throughput (automation, robot);
232	xiv) Cost of test, per sample;
233 234	xv) Availability of reference standards, reference materials and proficiency testing schemes. (See also chapter 2.2.6.).
235	7.2. Optimisation and standardisation of the test method
236 237 238	Once the method has been selected, it must be set up at the laboratory. Additional optimisation is necessary, whether the method was developed in-house (validation) or imported from an outside source (verification). Optimisation establishes critical specifications and performance standards for the test process as used in a specific laboratory.
239	7.2.1. Determinants of optimisation
240 241	 Critical specifications for equipment, instruments consumables, and reagents (e.g. chemicals, biologicals), reference standards, reference materials, and internal controls;
242 243	 Robustness – critical control points and acceptable ranges, attributes or behaviour at critical control points, using statistically acceptable procedures;
244	iii) Quality control activities necessary to monitor critical control points;
245	iv) The type, number, range, frequency, and arrangement of test run controls;
246	v) Criteria for non subjective objective acceptance or rejection of a batch of test results;
247	vi) Criteria for the interpretation and reporting of test results;
248	vii) A-Documented test method and reporting procedure for use by laboratory staff;
249 250	 viii) Evidence of technical competence for those who performing the test processes methods, authorising test results and interpreting results.
251	7.3. Validation of the test method
252 253 254 255 256 257 258 259 260 261 262 263 264	<u>Test method</u> validation evaluates the test for its-fitness for a given use purpose by establishing test-performance characteristics such as sensitivity, specificity, and isolation rate; and diagnostic parameters such as positive or negative cut-off, <u>repeatability, reproducibility</u> and titre of interest or significance. Validation should be done performed using an optimised, documented, and fixed procedure. The extent and depth of the validation process will depend on logistical and risk factors. It and may involve any number of activities and amount of data, with subsequent data analysis using appropriate statistical methods (<u>Chapter 1.1.6.</u>). Acknowledging diagnostic test validation science as a key element in the effective detection of infectious diseases, WOAH recently published a Special Issue representing an up-to-date compilation of the relevant standards (WOAH and non-WOAH) and guidance documents for all stages of diagnostic test validation studies in the peer-reviewed literature (Colling & Gardner, 2021). It is important to note that the current version of ISO 17025:2017 specifies that personnel must be authorised to perform validation and related activities, which means that training in validation and verification methods, including results interpretation, is likely to become more important to prove competence (Colling & Gardner, 2021).
265	7.3.1. Activities that validation might include
266 267	 Field or epidemiological studies, including disease outbreak investigations <u>and testing of samples</u> from infected and non-infected animals;
268 269	ii) Development of testing algorithms for specific purposes, e.g. surveillance, outbreak investigations, etc.;
	iii) Repeat testing in the same laboratory to establish the effect of variables such as operator,
270 271	reagents, equipment;

- 273v)Collaborative studies with other laboratories using the same documented method. Ideally274organised by a reference laboratory and including testing a panel of samples of undisclosed275composition or titre with expert evaluation of results and feedback to the participants to estimate276reproducibility;
 - vi) Reproduction of data from an accepted standard method, or from a reputable peer-reviewed publication (verification);
 - vii) Experimental infection or disease outbreak-studies;
 - viii) Analysis of internal quality control data.
- 281Validation is always a balance between cost, risk, and technical possibilities. There may be cases where282quantities such as only basic accuracy and precision can only be given determined, e.g. when the disease is283not present in a simplified way country or region. Criteria and procedures for the correlation of test results for284diagnosis of disease status or for regulatory action must be developed. The criteria and procedures developed285should account for screening methods, retesting and confirmatory testing.
- 286 Test validation is covered in chapter 1.1.6.

287 **7.4. Uncertainty of the test method**

288 <u>Statistically relevant numbers of samples from infected and non-infected animals are discussed in chapter 1.1.6. test</u> 289 validation and chapter 2.2.5 statistical approaches to validation.

290 7.4. Estimation of Measurement Uncertainty

- 291 Measurement of Uncertainty (MU) is "a parameter associated with the result of a measurement that characterises the 292 dispersion of values that could reasonably be attributed to the measure" (Eurachem, 2012). Uncertainty of 293 measurement does not imply doubt about a result but rather increased confidence in its validity. It is not the equivalent 294 to *error*, as it may be applied to all test results derived from a particular procedure.
- Laboratories must estimate the MU for each test method resulting in a <u>quantitative</u> measurement-included in their scope of accreditation, and for any methods used to calibrate equipment, included in their scope of accreditation (ISO/IEC 17025, 2005-2017b).
- Tests can be broadly divided into two groups: quantitative (<u>e.g.</u> biochemical assays, enzyme-linked immunosorbent assays [ELISA], titrations, real-time polymerase chain reaction [PCR], pathogen enumeration, etc.); and qualitative (bacterial culture, parasite identification, virus isolation, endpoint PCR, immunofluorescence, etc.).
- The determination of MU is well established in *quantitative* measurement sciences (ANSI, 1997). It may be given as a numeric expression of reliability and is commonly shown as a stated range. Standard deviation (SD) and confidence interval (CI) are examples of the expression of MU, for example the optical density result of an ELISA expressed as $\pm n$ SD, where *n* is usually 1, 2 or 3. The confidence interval (usually 95%) gives an estimated range in which the result is likely to fall, calculated from a given set of test data. For quantitative measurements, example for a top-down or control-sample approach are provided for an antibody ELISA in chapter 2.2.4, and by the Australian government webpage²⁸. An example for a quantitative PCR (TaqMan) assay is provided by Newberry & Colling (2021).
- 308The ISO/IEC 17025 requirement for "quality control procedures for monitoring the validity of tests" implies that the309laboratory must use quality control procedures that cover all major sources of uncertainty. There is no requirement310to cover each component separately. Laboratories may establish acceptable specifications, criteria, ranges, etc., at311critical control points for each component of the test process. The laboratory can then implement appropriate quality312control measures at these critical points, or seek to reduce or eliminate the uncertainty effect of each component.
- 313

315

977

278 279

280

7.4.1. Potential sources of uncertainty include:

- 314 i) <u>Sampling;</u>
 - ii) <u>Contamination;</u>
- 316 iii) <u>Sample transport and storage conditions;</u>
- 317 iv) <u>Sample processing;</u>

²⁸ Australian Government, Department of Agriculture, Fisheries and Forestry. Worked examples of measurement uncertainty. Measurement uncertainty in veterinary diagnostic testing – DAFF (agriculture.gov.au) (accessed 15 March 2023).

318	v) <u>Reagent quality, preparation and storage;</u>						
319	vi) <u>Type of reference material;</u>						
320	vii) <u>Volumetric and weight manipulations;</u>						
321	viii) <u>Environmental conditions;</u>						
322	ix) Equipment effects;						
323	x) <u>Analyst or operator bias;</u>						
324	xi) <u>Biological variability;</u>						
325	xii) <u>Unknown or random effects.</u>						
326	Systematic errors or bias determined by validation must be corrected by changes in the method,						
327	adjusted for mathematically, or have the bias noted as part of the report statement.						
328	If an adjustment is made to a test or procedure to reduce uncertainty or correct bias then a new						
329 330	source of uncertainty is introduced (the uncertainty of the correction). This must be assessed as part of the MU estimate.						
000							
331 332	The application of the principles of MU to <i>qualitative</i> testing is less well defined. The determination and expression of MU has not been standardised for veterinary (or medical, food, or environmental)						
332	testing laboratories, but sound guidance exists and as accreditation becomes more important,						
334	applications are being developed. The ISO/IEC 17025 standard recognises that some test methods						
335 336	may preclude metrologically and statistically valid calculation of uncertainty of measurement. In such cases the laboratory must attempt to identify and estimate all the components of uncertainty based						
337	on knowledge of the performance of the method and making use of previous experience, validation						
338	data, internal control results, etc.						
339	Many technical organisations and accreditation bodies (e.g. AOAC International, ISO, NATA, A2LA,						
340	Standards Council of Canada, UKAS, Eurachem, the Cooperation of International Traceability in						
341 342	Analytical Chemistry) teach courses or provide guidance on MU for laboratories seeking accreditation.						
343	The ISO/IEC 17025 requirement for "quality control procedures for monitoring the validity of tests"						
344	implies that the laboratory must use quality control procedures that cover all major sources of						
345	uncertainty. There is no requirement to cover each component separately. Laboratories may						
346 347	establish acceptable specifications, criteria, ranges, etc., at critical control points for each component of the test process. The laboratory can then implement appropriate quality control measures at these						
348	critical points, or seek to reduce or eliminate the uncertainty effect of each component. Measurement						
349	Uncertainty is covered in chapter 2.2.4.						
350	7.4.1. Components of tests with sources of uncertainty include:						
351	i) Sampling;						
352	ii) Contamination;						
353	iii) Sample transport and storage conditions;						
354	iv) Sample processing;						
355	v) Reagent quality, preparation and storage;						
356	vi) Type of reference material;						
357	vii) Volumetric and weight manipulations;						
358	viii) Environmental conditions;						
359	ix) Equipment effects;						
360	x) Analyst or operator bias;						
361	xi) Biological variability;						
362	xii) Unknown or random effects.						

- 363Systematic errors or bias determined by validation must be corrected by changes in the method,
adjusted for mathematically, or have the bias noted as part of the report statement.365If an adjustment is made to a test or procedure to reduce uncertainty or correct bias then a new
source of uncertainty is introduced (the uncertainty of the correction). This must be assessed as part
of the MU estimate.368Additional information on the analysis of uncertainty may be found in the Eurachem Guides to
 - Additional information on the analysis of uncertainty may be found in the Eurachem Guides to Quantifying Uncertainty in Measurement (Eurachem, 2012) and Use of uncertainty information in compliance assessment <u>Uncertainty Information in Compliance Assessment</u> (Eurachem, 2007).

7.5. Implementation and use of the test method

Training should be a planned and structured activity with steps to ensure adequate supervision is maintained while analysts are being trained. <u>Depending on the complexity of the test and the experience of the analyst, training may</u> include any combination of reading and understanding the documented test method, initial demonstration, performance of the test under supervision and independent performance. Analysts should be able to demonstrate proficiency in using the test method prior to producing being authorised to produce reported results, and on an ongoing basis.

The laboratory must be able to demonstrate traceability for all accredited tests and the principle should apply to all tests whether accredited or not. This covers all activities relating to test selection, development, optimisation, standardisation, validation, <u>verification</u>, implementation, reporting, personnel, quality control and quality assurance (see also Section 7.3.1. point vi). Traceability is achieved by using appropriate documented project management, record keeping, data management and archiving systems.

383 8. Strategic planning

369

370

Laboratories should have evidence of continual improvement, which is an obligatory requirement for accredited laboratories. The laboratory must be knowledgeable of and stay maintain current with knowledge of the relevant quality and technical management standards and with methods used to demonstrate laboratory competence and establish and maintain technical validity. Evidence of this may be provided by include:

- i) Attendance at conferences, organisation of in-house or external meetings on diagnostics and quality
 management;
- 390 ii) Participation in Membership of local and international organisations;
- 391 iii) Participation in writing Contribution to national and international standards (e.g. on ILAC and ISO committees);
- iv) <u>Maintenance of</u> current awareness of <u>publications</u>, <u>writing through review of</u> and <u>reviewing publications about</u>
 diagnostic methods <u>contribution to relevant literature</u>;
- 394 v) <u>Participation in</u> training programmes, including visits to other laboratories;
- 395 vi) Conducting research;
- 396 vii) Participation in cooperative programmes (e.g. Inter-American Institute for Cooperation in Agriculture);
- 397 viii) Exchange of procedures, methods, reagents, samples, personnel, and ideas;
- 398 ix) Planned, continual professional development and technical training;
- 399 x) Management reviews;
- 400 xi) Analysis of customer feedback;
- 401 xii) Root cause analysis of anomalies and implementation of corrective, preventive and improvement actions<u>. as</u> 402 <u>well as effectiveness reviews</u>.

REFERENCES

- 404 ANSI (AMERICAN NATIONAL STANDARDS INSTITUTE) (1997). ANSI/NCS²⁹ Z540-2-1997 (R2012), US Guide to the Expression 405 of Uncertainty in Measurement, First Edition. American National Standards Institute, <u>www.ansi.org</u>.
- COLLING A.& GARDNER I.A., EDS (2021). Diagnostic test validation science: a key element for effective detection and control 406 407 infectious animal diseases. Rev. Sci. Tech. Off. Int. Epiz.. **40**. Available of 408 https://doc.woah.org/dyn/portal/index.xhtml?page=alo&aloId=41245
- EURACHEM (2007). Eurachem/CITAC Guide: Use of Uncertainty Information in Compliance Assessment, First Edition.
 Eurachem Secretariat. <u>www.eurachem.org</u>.
- 411 EURACHEM (2012). Eurachem/CITAC³⁰ Guide (CG4): Quantifying Uncertainty in Analytical Measurement, Third Edition.
 412 Eurachem Secretariat. <u>www.eurachem.org</u>.
- ISO (2015). ISO 9001:2015. Quality Management Systems Requirements. International Organization for Standardization
 (ISO), <u>www.iso.org</u>.
- ISO/IEC (2004a 2020). ISO/IEC 17000:2004 2020. Conformity Assessment Vocabulary and General Principles.
 International Organization for Standardization (ISO), www.iso.org.
- ISO/IEC (2004b-2017a). ISO/IEC 17011:20042 017. Conformity Assessment General Requirements for accreditation
 Bodies Accrediting Conformity Assessment Bodies. International Organization for Standardization (ISO), <u>www.iso.org</u>.
- ISO/IEC (2005-2017b). ISO/IEC 17025:2005-2017. General Requirements for the Competence of Testing and Calibration
 Laboratories. International Organization for Standardization (ISO), <u>www.iso.org</u>.
- ISO/IEC (2010). ISO/IEC 17043:2010. Conformity Assessment General Requirements for Proficiency Testing.
 International Organization for Standardization (ISO), <u>www.iso.org</u>
- ISO/IEC (<u>2022-2012</u>). ISO/IEC 15189:<u>2012-2022</u>. Medical Laboratories Requirements for Quality and Competence.
 International Organization for Standardization (ISO), <u>www.iso.org</u>
- 425 JOHNSON P. & CABUANG L. (2021). Proficiency testing and ring trials. Rev. Sci. Tech. Off. Int. Epiz., 40, 189–203.
- <u>NEWBERRY K. & COLLING A. (2021). Quality standards and guidelines for test validation for infectious diseases in veterinary</u>
 <u>Iaboratories. Rev. Sci. Tech. Off. Int. Epiz.</u>, **40**, 227–237.
- <u>STEVENSON M., HALPIN K. & HEUER C. (2021). Emerging and endemic zoonotic diseases: surveillance and diagnostics. *Rev.* <u>Sci. Tech. Off. Int. Epiz.</u>, 40, 119–129.
 </u>
- WAUGH C. & CLARK G. (2021). Factors affecting test reproducibility among laboratories. *Rev. Sci. Tech. Off. Int. Epiz.*, 40,
 <u>131–141.</u>
- 432 433

434 435

403

NB: FIRST ADOPTED IN 1996 AS GOOD LABORATORY PRACTICE, QUALITY CONTROL AND QUALITY ASSURANCE. MOST RECENT UPDATES ADOPTED IN 2017.

²⁹ NCSL: The National Conference of Standards Laboratories.

³⁰ CITAC: The Cooperation of International Traceability in Analytical Chemistry.

 intended for veterinary use
 MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION
 Paris, 4–8 September 2023
 CHAPTER 1.1.9.
 TESTS FOR STERILITY AND FREEDOM FROM CONTAMINATION OF BIOLOGICAL MATERIALS INTENDED FOR VETERINARY USE

1

10

Annex 5. Item 5.1. – Chapter 1.1.9. Tests for sterility and freedom from contamination of biological materials

INTRODUCTION

11 The international trade-related movements of biological materials intended for veterinary use are subject to 12 restrictions imposed to minimise the spread of animal and human pathogens. Countries may impose 13 requirements for proof-of-freedom testing before allowing the regulated importation of materials of animal derivation and substances containing such derivatives. Where chemical or physical treatments are 14 15 inappropriate or inefficient, or where evidence is lacking of the effectiveness of the treatment is lacking, there 16 may be general or specific testing requirements imposed by authorities of countries receiving such materials. 17 This chapter provides guidance on the approach to such-regulated testing, particularly as might be applied 18 to the movement of vaccine master seed and master cell stocks, and to related biological materials used in 19 manufacturing processes. The term seed stocks is used when testing live products, for killed products the preferred reference is master cell stocks. While the onus for ensuring safety of a product remains with the 20 21 manufacturer and may be regulated by therapeutic guidelines, this chapter provides procedures that are 22 designed in particular to minimise the risk of undetected contaminants in veterinary therapeutics and 23 biological reagents causing the cross-border spread of agents of concern to particular-importing countries. 24 In their review "Extraneous agent detection in vaccines" Farsang & Kulcsar, 2012 reported the following examples of contamination of vaccines with extraneous agents: a) Foamy virus (Spumaretroviridae) was 25 identified as a contaminant of primary monkey kidney cultures used for vaccine production in the early 1950, 26 27 b) In the 1960s it was shown that vellow fever live attenuated vaccines prepared in chicken embryo fibroblasts were infected with avian leukosis virus (ALV). c) Calicivirus was found in Chinese hamster ovary 28 29 (CHO) cells, d) Newcastle disease vaccine strains were found in different live poultry vaccines, e) in 1990 a 30 live attenuated multi component canine vaccine was contaminated with a serotype of Bluetongue virus 31 causing abortions and death in pregnant bitches, f) Fetal calf serum transmitted Pestiviruses (BVDV types 1 and 2) are one of the most common extraneous agents in veterinary and human vaccines, g) RD114 is a 32 33 replication- competent feline endogenous gamma retrovirus which contaminated canine corona and 34 parvovirus vaccines . h) a notable case of human vaccine contamination may have been when in the 20th 35 century tens of millions of people worldwide were immunised with polio vaccines containing simian virus 40 (SV40). SV40 was found to cause cancer in animals and is associated with human brain, bone and lung 36 37 cancers, however, a clear connection was not found between this certain vaccine and any human tumour 38 case, i) a porcine circovirus 1 (PCV1) was found in a rotavirus vaccine widely used worldwide for children. 39 Farsang & Kulcsar (2012) and WHO (2015) describe case studies of veterinary and human vaccines 40 contaminated with extraneous agents and findings support the need of accurate and validated amplification 41 and detection methods as key elements for effective detection and control. Further examples are given in 42 Protocol examples below. Control of contamination with transmissible spongiform <u>Section G.</u> 43 encephalopathy (TSE) agents is not covered in this chapter because <u>standard</u> testing and physical

treatments cannot be used to ensure freedom from these agents. <u>Detection methods are described in</u>
 <u>Chapter 3.4.5. Bovine spongiform encephalopathy.</u>

46 Sterility is defined as the absence of viable microorganisms, which for the purpose of this chapter, includes 47 viruses. It should be achieved using aseptic techniques and validated sterilisation methods, including heating, 48 filtration, chemical treatments, and irradiation that fits the intended purpose. Freedom from contamination is 49 defined as the absence of specified viable microorganisms. This may be achieved by selecting materials 50 from sources shown to be free from specified microorganisms and by conducting subsequent procedures 51 aseptically. Adequate assurance of sterility and freedom from contaminating microorganisms can only be 52 achieved by proper control of the primary materials used and their subsequent processing. Tests on 53 intermediate products are necessary throughout the production process to check that this control has been 54 achieved.

- 55 Biological materials subject to contamination that cannot be sterilised before or during use in vaccine 56 production, such as ingredients of animal origin, e.g. serum and trypsin, primary and continuous cells and 57 cell lines, and viral or bacterial seed stocks, etc., should be tested for viable extraneous agents before use. 58 Assays to detect viral contaminants, if present, can be achieved by various culture methods, including use 59 of embryonated eggs, which are supported by cytopathic effects (CPE) detection/embryo death, fluorescent 60 antibody techniques and other-suitable <u>(fit for purpose),</u> methods such as polymerase chain reaction (PCR) and antigen detection ELISA (enzyme-linked immunosorbent assay). As is explained in more detail in this 61 62 chapter care must be taken when using PCR and ELISA techniques for detection as such tests do not 63 distinguish viable from non-viable agent detection. <u>Specific assays to detect other contaminants, such as</u> 64 <u>fungi, protozoa and bacteria (including rickettsia and mycoplasma) are also described.</u>
- 65Avian materials and vaccines are required to be inoculated on to primary avian cell cultures or eggs for the66detection of avian viruses. A combination of general tests, for example to detect haemadsorbing,67haemagglutinating and CPE-causing viruses and specific procedures aimed at the growth and detection of68specific viruses is recommended to increase the probability of detection. Assays to detect other69contaminants, such as bacteria, fungi, protozoa, rickettsia and mycoplasma are also described.
- Procedures applied <u>Testing procedures</u> should be validated and found to be "fit for purpose" following
 Chapter 1.1.6. Validation of diagnostic assays for infectious diseases of terrestrial animals, where possible.
- It is a requirement of many regulators, that a laboratory testing report notes the use of validated procedures
 and describes the validated procedures in detail including acceptance criteria. This gives the regulator
 transparency in the procedures used in a testing laboratory.
- 75The validation assessment of an amplification process in cell culture should include documentation of the
history of permissive cell lines used, reference positive controls and culture media products used in the
process of excluding adventitious agents, to ensure the process is sound and is not compromised. The
validation assessment should give information (published or in-house) of the limitations that may affect test
outcomes and an assessment of performance characteristics such as analytical specificity and sensitivity of
each cell culture system, using well characterised, reference positive controls.
- *It is <u>the</u> responsibility of the submitter to <u>assure ensure</u> a representative selection and number of items to
 <i>be tested. The principles of Appendix 1.1.2.1* Epidemiological approaches to sampling: sample size
 calculations of Chapter 1.1.2 Collection, submission and storage of diagnostic specimens apply <u>describes</u>
 <u>the principles to be applied</u>. Adequate transportation is described in Chapter 1.1.2 and Chapter 1.1.3
 Transport of biological materials <u>describe transportation requirements</u>.
- 86

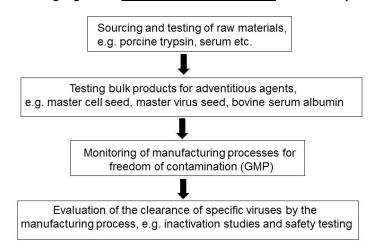
A. AN OVERVIEW OF TESTING APPROACHES

Although testing is seen as a key component of biosafety in biological products intended for veterinary use, testing is not
 enough to ensure a given product is free of viable infectious contaminants, and so a holistic, multifaceted approach must
 be taken. Such an approach includes risk assessment, risk mitigation and management strategies (Barone *et al.*, 2020).
 In general:

- Primary materials must be collected from sources shown to be free from contamination and handled in such a way as
 to minimise contamination and the opportunities for any contaminants to multiply (Figure 1).
- Materials that are not sterilised and those that are to be processed further after sterilisation must be handled aseptically.
 Such materials will require further assessment of freedom of contaminants at certain stages of production to assure freedom of adventitious agents.
- Materials that can be sterilised without their biological activities being affected unduly must be sterilised by a method effective for the pathogens concerned of concern. The method must reduce the level of contamination to be undetectable, as determined by an appropriate sterility test study. (See Section D.1. below). If a sterilisation process is used, it shall be validated to demonstrate that it is fit for purpose. Suitable controls will be included in each sterilisation process to monitor efficiency.
- The environment in which any aseptic handling is carried out must be maintained in a clean state, protected from external sources of contamination, and controlled to prevent internal contamination. Rules governing aseptic preparation of vaccines are documented in Chapter 2.3.3 *Minimum requirements for the organisation and management* of a vaccine manufacturing facility.

105

Figure 1. Testing algorithm Risk assessment flowchart for vaccine production.



106

107 Some procedures have been properly validated and found to be "fit for purpose", whilst others may have undergone only limited validation studies. For example, methods for bacterial and fungal sterility may have not been formally validated 108 although they have been used for many years. In particular, the in-vivo and cell culture in-vitro methods have essentially 109 unknown sensitivity and specificity (Sheets et al., 2012) though there is an accepted theoretical sensitivity, regarding cell 110 culture of 1 colony plaque-forming unit (CFU-PFU). For example, an evaluation of methods to detect bovine and porcine 111 viruses in serum and trypsin based on United States (of America) Code of Federal Regulations, Title 9 (9CFR) revealed 112 113 gaps in sensitivity, even within virus families (Marcus-Secura et al., 2011). It is therefore important to interpret, and report results in the light of specific conditions of cultures employed and considering sensitivity and specificity of detection 114 systems. 115

Newer, more sensitive methods such as molecular assays may afford the ability to detect contaminants, which may not be successfully amplified in traditional culturing systems. The detection range can be broadened by using family specific primers and probes if designed appropriately. However, most, if not all <u>such new_molecular-based</u> tests are also able to detect evidence for non-infectious contaminants, such as traces of nucleic acid from inactivated contaminants. Follow up testing would be required to determine the nature of the contaminant, for example, non-infectious nucleic acid or infectious virus. Attempts at virus isolation or sequencing may remedy this. Note: molecular assays if not designed as fit for purpose may miss detection of contaminating agents or lack sensitivity to do so (Hodinka, 2013).

More recently metagenomic high throughput sequencing (HTS) workflows have shown potential for quality control of 123 biological products (van Borm et al., 2013) and vaccines (Baylis et al., 2011; Farsang & Kulcsar, 2012; Neverov & 124 Chumakov, 2010; Onions & Kolman, 2010; Victoria et al., 2010) in particular for the identification and characterisation of 125 126 unexpected highly divergent pathogen variants (Miller et al., 2010; Rosseel et al., 2011) that may remain undetected using targeted diagnostic tests. Nevertheless, targeted assays, e.g. amplification in cell culture followed by polymerase chain 127 reaction (PCR) may be superior to HTS for specific agent detection (Wang et al., 2014) due to lack of sensitivity of HTS at 128 129 this time. Chapter 1.1.7. gives an overview of the standards for high throughput sequencing, bioinformatics and 130 computational genomics. Similarly, recent improvements in protein and peptide separation efficiencies and highly accurate 131 mass spectrometry have promoted the identification and quantification of proteins in a given sample. Most of these new 132 technologies are broad screening tools, limited by the fact that they cannot distinguish between viable and non-viable 133 organisms.

organisms.

Given the availability of new technologies, there will be future opportunities and challenges to determine presence of
 extraneous agents in biologicals intended for veterinary use for industry and regulators. Problems can arise when the
 presence of genome positive results are interpretated as evidence for the presence of contamination (Mackay & Kriz,
 2010). When using molecular technologies, it is important to understand the correlation between genome detection and
 detection of live virus agent. It cannot be assumed that detection of genome corresponds to the presence of an infectious
 agent.

140B. LIVING VIRAL VACCINES FOR ADMINISTRATION BY INJECTION, OR THROUGH141DRINKING WATER, SPRAY, OR SKIN SCARIFICATION

- Materials of animal origin shall should be (a) sterilised, or (b) and obtained from healthy animals that, in so far as is possible, should be shown to be free from pathogens that can be transmitted from the species of origin to the species to be vaccinated, or any species in contact with them by means of extraneous agents testing.
- Seed lots of virus, any continuous cell line and biologicals used for virus growth shall should be shown to be free from viable-bacteria, fungi, mycoplasmas, protozoa, rickettsia, and extraneous viruses and other pathogens-that can be transmitted from the species of origin to the species to be vaccinated or any species in contact with them. There may be some exceptions for a limited number of non pathogenic bacteria and fungi to be present in live viral vaccines produced in eggs and administered through drinking water, spray, or skin scarification.
- For the production of vaccines in embryonated chicken eggs and the quality control procedures for these vaccines, it is recommended (required in many countries) that eggs from specific pathogen-free birds should be used.
- Each batch of vaccine shall-should pass tests for freedom from extraneous agents that are consistent with the importing country's requirements for accepting the vaccine for use. Some examples of document acceptable testing procedures-processes in various countries include: (US) Code of Federal Regulations (2015); European Pharmacopoeia (2014); European Commission (2006); World Health Organization (WHO) (1998; 2012) and Department of Agriculture (of Australia) (2013).
- 157 Code of Federal Regulations, Title 9 (9CFR) (of the United States of America) (2015).
 - Department of Agriculture, Forest and Fisheries (Australia) (2013).
 - Department of Agriculture, Forest and Fisheries (Australia) (2021b) Live Veterinary Vaccines.
 - European Medicines Agency Sciences Medicines Health (2016).
- 161 European Pharmacopoeia, 10th Edition (2021).
- 162 World Health Organization (WHO) (1998; 2012).

Tests for sterility-freedom of contamination shall should be appropriate to prove that the vaccine is free from viable 163 4 extraneous viruses, bacteria including rickettsia and mycoplasmas, fungi, and protozoa. Each country will have 164 165 particular-requirements as to what agents are necessary to exclude should be tested for and what by which procedures are acceptable. Such tests will include amplification of viable extraneous agents using cell culture that is 166 susceptible to particular known viruses of the species of concern, tests in embryonated eggs, bacterial, mycoplasma 167 and fungal culturing techniques and, where necessary and possible there is no alternative le, tests involving animal 168 inoculation. PCR, fluorescence antibody test (FAT), presence of colonies or cytopathic effects (CPE) and antigen 169 detection ELISA will-can be used for detection purposes after amplification using culturing techniques to improve 170 specificity and sensitivity. If in-vitro or in-vivo amplification of the target agent is not possible, direct PCR may be 171 useful if validated for this purpose. 172

173C. LIVING VIRAL VACCINES FOR ADMINISTRATION THROUGH DRINKING WATER,174SPRAY, OR SKIN SCARIFICATION

175 1. Section B applies.

158

159

160

 A limited number of contaminating, non pathogenic bacteria and fungi may be permitted (see Section 1.2.2 General Procedure for testing live viral vaccines produced in eggs and administered through drinking water, spray, or skin scarification for the presence of bacteria and fungi).

179

D-C. INACTIVATED VIRAL AND BACTERIAL VACCINES

- 180 1. Each batch of vaccine shall pass a test for inactivation of the vaccinal virus seed and should include inactivation studies on representative extraneous agents if the virus or bacterial seed has not already been tested and shown to be free 181 182 from extraneous agents. An example of a simple inactivation study could include assessment of the titre of live vaccine before and after inactivation and assessing the log₁₀ drop in titre during the inactivation process. This would give an 183 indication of the efficacy of the inactivation process. There is evidence that virus-titration tests may not have sufficient 184 185 sensitivity to ensure complete inactivation. In these circumstances, a specific innocuity test would need to be developed and validated to be fit for increased sensitivity. To increase sensitivity more than one passage would be required 186 depending on the virus or bacteria of concern. An example of this approach can be found at: 187 188 https://www.aphis.usda.gov/animal health/vet biologics/publications/memo 800 117.pdf (accessed 25 July 2023).
- If studies on representative extraneous agents are required, then spiking inactivated vaccine with live representative agents and following the example of an inactivation study as in D.1 above would <u>could</u> be useful. The inactivation process and the tests used to detect live <u>virus-agent</u> after inactivation must be validated and shown to be suitable for their intended purpose.
- 193 In addition, each country may have <u>particular_its own</u> requirements for sourcing or tests for sterility as detailed in 194 Section B above.

E. D. LIVING BACTERIAL VACCINES

196 1. <u>See</u> Section B applies.

195

- Seed lots of bacteria shall be shown to be free from other bacteria as well as fungi and mycoplasmas, protozoa, rickettsia, and extraneous viruses. Agents required for exclusion will be dependent on the country accepting the vaccine for use. Use of antibiotics to 'inactivate' the living bacterial seed or vaccine prior to exclusion of viruses and fungi is recommended to ensure testing in culture is sensitive. Interference testing is recommended to ensure that the antibiotics used do not affect the growth of the extraneous virus or fungi that is being excluded. Sonication may also be useful
- Interference testing is required to ensure that antibiotics used (or sonication) does not affect the growth of extraneous
 virus or fungi being excluded, compromising the test outcome.
- 205 Due to the difficulties and reduced sensitivity in exclusion of extraneous bacteria and some mycoplasma, protozoa, and 206 rickettsia from high-titred seed lots of bacteria, the use of narrow-range antibiotics aimed specifically at reducing seed 207 lot bacteria is recommended useful if antibiotics do not affect the growth of bacteria being excluded. The optimal concentration of antibiotics can be determined in a dilution experiment such as documented in 9CFR Section 113.25(d). 208 209 Other methods of exclusion of extraneous bacteria from bacterial seeds may include filtering for size exclusion such as 210 removing bacteria seed to look for mycoplasma contamination and use of selective culturing media. Such processes would require validation verification to ensure the process does not affect the sensitivity of exclusion of extraneous 211 212 agents of concern.
- 213 <u>3. Sonication of a living bacterial seed may be useful when excluding specific viral agents. Once again, the inactivation</u>
 214 <u>procedure would require a verification process to ensure the adventitious virus being excluded is not affected by the</u>
 215 <u>treatment. Use of a suitable reference virus control during the exclusion process would be required.</u>
- Direct PCR techniques may be useful when culturing processes fail to be sensitive successful in detecting extraneous bacteria from live bacterial seeds or vaccines.

218

F. INACTIVATED BACTERIAL VACCINES

 Section D applies. It should not be necessary to test for extraneous viruses that would not grow in bacteriological culture media as long as freedom from contamination of all starting materials can be assured. Complete inactivation of the vaccinal bacteria should be demonstrated by means of titration and innocuity tests – in some cases general bacterial sterility testing (Section I.2.1) may suffice.

223 G-<u>E</u>. SERA, <u>PLASMA</u> AND DIAGNOSTIC AGENTS FOR ADMINISTRATION TO 224 ANIMALS

- Section B.1 <u>applies for sera/diagnostic agents that are not inactivated</u>. Section C applies for non-inactivated sera/diagnostic agents.
- Some countries require quarantine, health certification, and tests for specific diseases to be completed for all serum and plasma donor animals, for example, 9CFR (2015) and Australian Quarantine Policy and Requirements for the Importation of Live and Novel Veterinary Bulk and Finished Vaccines (1999). For some diseases, for example equine infectious anaemia, the product (plasma) must be stored until the seroconversion period has been exceeded and the donors tested negative.
- 3. It is recommended that each batch of non inactivated serum be assessed for viable extraneous agents, including
 mycoplasma. Each batch of serum shall pass a test for freedom from extraneous agents. Suitable test methods have
 been published for various countries, for example, European Pharmacopoeia (2014); 9 CFR (2015) and Australian
 Quarantine Policy and Requirements for the Importation of Live and Novel Veterinary Bulk and Finished Vaccines
 (1999) and Department of Agriculture (of Australia) (2013).
- 237 4. Inactivated serum, Section D applies.
- 238 5. Section B or D may apply if a virus is used in the production of the diagnostic agent; Section E or F may apply if a
 239 bacterium is used.

H. F. EMBRYOS, OVA, SEMEN

Special precautions must be taken with relation to the use of embryos, ova, semen (Hare, 1985). Most countries will have
 regulatory guidelines for import of these biologicals for veterinary use. Such guidelines can be found at various websites
 such as the European Commission (2015), FAO and Department of Agriculture Forest and Fisheries (2021a; 2021b),
 though many such some guidelines may give more detail in regard-to the food safety aspect.

245

240

J. G. PROTOCOL EXAMPLES

246 1. General procedures <u>Introduction</u> to protocol examples

247 This section provides some examples to illustrate scope and limitations of testing protocols. It is not intended to be 248 prescriptive or exhaustive. Examples are based on standards and published methods to increase the sensitivity for 249 exclusion of live adventitious agents, using general and specific techniques.

250 In principle, proposed testing represents-an attempted isolation of viable agents in culturing systems normally considered supportive of the growth of each specified agent or group of general agents. After amplification, potential pathogens can 251 be detected further, by sensitive and specific diagnostic tests such as FAT or PCR if as required. General detection systems 252 can include haemabsorbance and CPE by immunohistochemistry staining methods. The example procedures for sterility 253 254 detection of contamination testing and general detection of viable-virus, fungi, protozoa and bacteria (including rickettsia 255 and mycoplasma, fungi and viruses) described below are derived from standards such as the 9CFR (2015), European Pharmacopoeia, (2014)-10th Edition (2021), European Commission (2006), WHO-Medicines Agency Sciences Medicines Health (2016), Department of Agriculture, Forest and Fisheries (Australia) (2013) and World Health Organization (1998; 256 257 258 2012).

Individual countries or regions should adopt a <u>holistic</u>, risk-based approach to determine the appropriate testing protocols
 based on their animal health status. As well as applying general testing procedures documented in national or regional
 standards as mentioned above, it may be necessary to apply rigorous exclusion testing for specific agents that are exotic
 to the particular country or region of concern.

General procedures <u>will_do</u> not necessarily detect all extraneous agents that may be present in biological material; however,
 they are useful as screening tests. Some examples of agents that may require specific methods for detection in biologicals
 refer to Table 1 below. Procedures documented in the Review of Published Tests to Detect Pathogens in Veterinary
 Vaccines Intended for Importation into Australia (2013) available from the Department of Agriculture. <u>Forest</u> and Water
 Resources, Australia Eisheries are able to address such agents in offering sensitive testing approaches based on reputable
 publications. <u>A CVMP reflection paper published written by the European Medicines Agency Sciences Medicines Health</u>

269 <u>Committee of Veterinary Medicinal Products (CVMP) in (2016), adopted in May 2017, documents-lists specific test method</u> 270 <u>approaches for a number of agents, listed in Table 1, that cannot be excluded using general test procedures (Table 1).</u>

Exclusion of specific agents requires procedures that maximise sensitivity by providing ideal amplification and detection of
 the pathogen in question. Extraneous agents, for example, Maedi Visna virus, bovine immunodeficiency virus, (and other
 retroviruses), *Trypanosoma evansi* and porcine respiratory coronavirus are difficult to culture even using the most sensitive
 approaches. In these circumstances, application of molecular assays directly to the biological material-in question to
 assess, assessing for the presence of nucleic acid from adventitious agents offers an alternative. Refer to Table 1.
 Consideration must be noted as described in Section A.6 as, though detection of the presence of non-viable and host
 associated agents may-is also be detected using this procedure-possible.

278 Table 1 gives examples of causative infectious agents that may be present in animal biologicals intended for veterinary use, for example PCV-1 in a rotavirus vaccine (WHO, 2015). BVDV is well known for its presence in many bovine 279 associated biologicals, including cell culture. More recently, non-CPE pestivirus, BVD type 3 (HoBi-like) are found in foetal 280 calf serum and cell culture. Classical Swine fever has contaminated various porcine cell lines used for African swine fever 281 and FMDV diagnosis, and thus the potential for contamination of porcine based vaccines. PEDV is linked to spray-dried 282 porcine plasma used for feed. This is not an exhaustive list of agents of concern or by any means required for exclusion 283 by every country based on risk, they are just examples of infectious agents that are not culturable using general culturing 284 procedures and require a more use of specialised culturing processes and specific detection process by means of the 285 indirect fluorescent antibody test, PCR or ELISA, where applicable processes. Notably, some subtypes of an agent type 286 287 may be detectable by general methods, and some may require specialised testing for detection. For example, bovine adenovirus subgroup 1 (serotypes 1, 2, 3 and 9) can be readily isolated using general methods (Vero cells) however bovine 288 adenovirus subgroup 2 (serotypes 4, 5, 6, 7, 8 and 10) are not readily isolated and required specialised methods for 289 isolation. 290

291 292

294

305

Table 1. Some Examples of infectious agents of veterinary importance

 that require specialist specialised

 culturing and detection techniques

Rotaviruses	Pestiviruses (non-CPE)	Turkey rhinotracheitis
Porcine epidemic diarrhoea virus	Bluetongue virus	Brucella abortus
Porcine circoviruses (PCV 1, 2)	Swine pox virus	Rickettsias
Swine/equine influenza, some strains	Some adenoviruses	Protozoa
Bovine respiratory syncytial virus	<u>Rhabdoviruses (e.g.</u> rabies virus <u>)</u>	Some fungi (e.g. <i>Histoplasma</i>)

293 2. <u>Example of detection of bacteria and fungi contamination</u>

2.1. General procedure for assessing the sterility of viable bacteria and fungi

295 Standard tests for detecting extraneous bacteria and fungi (sterility testing) in raw materials, master cell stocks, or 296 final product are the membrane filtration test or the direct inoculation sterility test.

For the membrane filtration technique, a filter having a nominal pore size not greater than 0.45 µm and a diameter of at least 47 mm should be used. Cellulose nitrate filters should be used if the material is aqueous or oily; cellulose acetate filters should be used if the material is strongly alcoholic, oily or oil-adjuvanted. Immediately before the contents of the container or containers to be tested are filtered, the filter is moistened with 20–25 ml of Diluent A or B.

301	2.1.1.	Diluent A

302Diluent A is for aqueous products or materials. Dissolve 1 g peptic digest of animal tissue in water to303make 1 litre, filter, or centrifuge to clarify, adjust the pH to 7.1 ± 0.2, dispense into containers in 100 ml304quantities, and sterilise by steam.

2.1.2. Diluent B

306Diluent B is for oil-adjuvanted products or materials: Add 1 ml polysorbate 80 to 1 litre Diluent A, adjust
the pH to 7.1 ± 0.2, dispense into containers in 100 ml quantities, and sterilise by steam.308If the biological being tested has antimicrobial properties, the membrane is washed three times after
sample application with approximately 100 ml of the appropriate diluent (A or B). The membrane is then
transferred whole to culture media, aseptically cut into equal parts and placed in media, or the media is

311transferred to the membrane in the filter apparatus. If the test sample contains merthiolate as a312preservative, fluid thioglycolate medium (FTM) is used and the membranes are incubated at both 30–31335°C and 20–25°C. If the test sample is a killed biological without merthiolate preservative, FTM is used314at 30–35°C and soybean casein digest medium (SCDM) at 20–25°C. If the sample tested is a live viral315biological, SCDM is used at both incubation temperatures. It has been suggested that sulfite-polymyxin-316sulfadiazine agar be used to enhance the detection of *Clostridium* spp. when the membrane filtration317technique is used (Tellez *et al.*, 2005).

If direct inoculation of culture media is chosen, a sterile pipette or syringe and needle are used to aseptically transfer the biological material directly into liquid media. If the biological being tested has antimicrobial properties, the ratio of the inoculum to the volume of culture medium must be determined before the test is started, for example as explained in 9CFR 113.25(d) and detailed testing procedures can be found for example in supplemental assay method USDA SAM 903 https://www.aphis.usda.gov/animal_health/vet_biologics/publications/sam903.pdf_(accessed_24 July 2023) (SAM) 903 USDA SAM 003 ممك https://www.aphis.usda.gov/animal_health/vet_biologics/publications-2022). (accessed Julv То determine the correct medium volume to negate antimicrobial activity, 100 CFU of the control microorganisms listed in Table 2 are used. If the test sample contains merthiolate as a preservative, FTM is used in test vessels incubated at both 30-35°C and 20-25°C. Growth should be clearly visible after an appropriate incubation time (see Section I.2.1.3 Growth promotion and test interference). If the test sample is a killed biological without merthiolate, or a live bacterial biological, FTM is used at 30-35°C and SCDM at 20-25°C. If the test sample is a live viral biological, SCDM is used at both incubation temperatures. If the inactivated bacterial vaccine is a clostridial biological, or contains a clostridial component, the use of FTM with 0.5% added beef extract (FTMB) in place of FTM is preferred. It may also be desirable to use both FTM and SCDM for all tests.

Table 2. Some American Type Culture Collection ³¹ strains with their respective
medium and incubation conditions

Medium	Test missographism	Incubatio	Incubation			
wealum	Test microorganism	Temperature (°C)	Conditions			
FTM	Bacillus subtilis ATCC # 6633	30–35	Aerobic			
FTM	Candida krusei ATCC # 6258	20–25	Aerobic			
SCDM	Bacillus subtilis ATCC # 6633	30–35	Aerobic			
SCDM	Candida krusei ATCC # 6258	20–25	Aerobic			
FTMB	Clostridium sporogenes ATCC # 11437	30–35	Anaerobic			
FTMB	Staphylococcus aureus ATCC #6538	30–35	Aerobic			

337For both membrane filtration and direct inoculation sterility tests, all media are incubated for no fewer338than 14 days. At intervals during incubation, and after 14 days' incubation, the test vessels are339examined for evidence of microbial growth. Microbial growth should be confirmed by subculture and340Gram stain.

318

319 320

321

322

323

324

325

326

327 328

329

330

331

332 333

334

335 336

341

342

343

344

345

346

347

348

349

2.1.3. Example of growth promotion and test interference

The sterility of the media should be confirmed by incubating representative containers at the appropriate temperature for the length of time specified for each test.

The ability of the culture media to support growth in the presence and absence of product, product components, cells, seeds, or other test material should be validated for each product to be tested, and for each new batch or lot of culture media for example as outlined in 9CFR 113.25(b). Detailed testing procedures can be found for example in USDA SAMs 900-902, See USDA APHIS | Supplemental Assay Methods - 900 Series (accessed 22 July 2023) https://www.aphis.usda.gov/animal_health/vet_biologics/publications (accessed 4 July 2022).

Report of the Meeting of the Biological Standards Commission / September 2023

³¹ American Type Culture Collection, 10801 University Boulevard, Manassas, Virginia 20110-2209, USA.

350 To test for ability to support growth in the absence of the test material, media should be inoculated 351 with 10-100 viable control organisms of the suggested ATCC strains listed in Table 2 and incubated according to the conditions specified. 352 To test for ability of the culture media to support growth in the presence of the test material, containers 353 354 should be inoculated simultaneously with both the test material and 10-100 viable control organisms. 355 The number of containers used should be at least one-half the number used to test the product or 356 product component. The test media are satisfactory if clear evidence of growth of the control organisms appears in all inoculated media containers within 7 days. In the event that growth is 357 evident, the organism should be identified to confirm that it is the organism originally added to the 358 359 medium. The sterility test is considered invalid if any of the media show inadequate growth response, 360 or if the organism recovered, is not the organism used to inoculate the material. If the material being tested renders the medium turbid so that the presence or absence of microbial 361 growth cannot be readily determined by visual examination, 14 days after the beginning of incubation 362 transfer portions (each not less than 1 ml) of the medium to fresh vessels of the same medium and 363 then incubate the original and transfer vessels for not less than 4 days. 364

365 2.2. General procedure for testing live viral vaccines produced in eggs and administered 366 through drinking water, spray, or skin scarification for the presence of bacteria and fungi

367 Each batch of final container biological should have an average contamination of not more than one bacterial or fungal colony per dose for veterinary vaccines. From each container sample, each of two Petri dishes are inoculated 368 with vaccine equal to ten doses if the vaccine is recommended for poultry, or one dose if recommended for other 369 animals. To each plate 20 ml of brain heart infusion agar are added containing 0.007 IU (International Units) of 370 penicillinase per ml. One plate should be incubated at 30-35°C for 7 days and the other at 20-25°C for 14 days. 371 Colony counts are made at the end of each incubation period. An average colony count of all the plates representing 372 373 a batch should be made for each incubation condition. If the average count at either incubation condition exceeds one colony per dose in the initial test, one retest to rule out faulty technique may be conducted using double the 374 number of unopened final containers. If the average count at either incubation condition of the final test for a batch 375 exceeds one colony per dose, the batch of vaccine should be considered unsatisfactory. 376

377 2.3<u>2</u>. Example of general procedure for testing seed lots of bacteria and live bacterial biologicals for purity

- Each seed lot of bacteria or batch of live bacterial biological should be tested for purity by inoculation of SCDM, which is incubated at 20–25°C for 14 days, and FTM, which is incubated at 30–35°C for 14 days. <u>Using good practices in</u> sterile technique to avoid laboratory contamination, a sterile pipette or syringe and needle is used to aseptically transfer the quantity of biological directly into the two types of culture medium. The minimum ratio of inoculum to culture medium is 1/15. <u>Both positive and negative controls are set up as well</u>.
- If the inoculum or growth of the bacterial vaccine renders the medium turbid so that the absence of atypical microbial growth cannot be determined by visual examination, subcultures should be made from all turbid tubes on day 3 through until day 11. Subculturing is done by transferring 0.1–1.0 ml to differential broths and agar and incubating for the balance of the 14-day period. Microscopic examination by Gram stain should also be done.
- 388 If no atypical growth is found in any of the test vessels when compared with a positive control included in the test, the lot of biological may be considered satisfactory for purity. If atypical growth is found but it can be demonstrated by a 389 negative control that the media or technique were faulty, then the first test may should be repeated. If atypical growth 390 391 is found but there is no evidence invalidating the test, then a retest may should be conducted. Twice the number of biological containers and test vessels of the first test are used in the retest. If no atypical growth is found in the retest, 392 the biological could be considered to be satisfactory for purity but the results from both the initial and retest should 393 be reported for assessment by the individual countries relevant regulatory agency if the laboratory is sure that the 394 first test result was not due to in-laboratory contamination. If atypical growth is found in any of the retest vessels, the 395 396 biological is considered to be unsatisfactory for purity. If, however, it can be demonstrated by controls that the media 397 or technique of the retest were faulty, then the retest may should be repeated.

398 2.4<u>3</u>. An Example of a specific test procedure for exclusion of *Brucella sp.* including *B. abortus* 399 (where general testing is not sufficient) for detection of *Brucella abortus*

It should be confirmed that each batch of culture medium supports the growth of *B. abortus* by inoculating plates and
 flasks of biphasic medium with a known number of cells (around 100) of the fastidious *B. abortus* biovar 2. If the
 media supports the growth of this biotype it will support all other biovars.

- Inoculate 1.0 ml of prepared master or working viral-live agent or cell seed material (not containing antibiotics) by
 inoculating 50 µl of the test product into each of 10 flasks containing biphasic medium. At the same time 10 plates of
 serum dextrose agar (SDA) are inoculated with 50 µl of inoculum and spread with a sterile bent glass Pasteur pipette
 or hockey stick. An un-inoculated serum dextrose agar plate and a biphasic flask are also set up at the same time as
 negative controls.
- For assessment of inhibitory substances 50 μl of previously prepared master or working viral or cell seed material
 and 10–100 CFU of *B. abortus* are inoculated on to duplicate SDA plates. Positive controls are prepared by inoculating
 10–100 CFU of *B. abortus* on to duplicate SDA plates.
- 411 All plates and flasks are incubated at 37° C in a 5–10% CO₂ environment. Plates are incubated with the agar 412 uppermost and flasks with the agar slope vertical. Flasks are incubated with the cap loose.
- Plates are checked for growth of colonies at days 4 and 8 of incubation. The biphasic medium is examined every 4 to
 7 days for 28 days. After each examination of the flasks, they are tilted so that the liquid phase runs over the solid
 phase, then righted and returned to the incubator.
- 416 During the incubation period, SDA plates with positive control and test material are visually compared with plates with 417 the positive control only and if there is no inhibition of growth of the organism in the presence of the test material, the 418 interference testing test is successful, and testing can be assured to be sensitive.
- 419 Any signs of growth of suspicious contaminating microorganisms on SDA plates, cloudiness or colonies in biphasic 420 flasks require follow-up testing by PCR to confirm whether *B. abortus* is present.

421 2.5<u>4</u>. An Example of a general procedure for detection of Salmonella contamination

422 Each batch of live virus biological reagents made in eggs should be free from contamination with Salmonella. This testing must be done before bacteriostatic or bactericidal agents are added. Five samples of each batch should be 423 tested; 5 ml or one-half of the container contents, whichever is the lesser, of the sample should be used to inoculate 424 100 ml of tryptose broth and tetrathionate broth. The inoculated broths should be incubated for 18-24 hours at 35-425 37°C. Transfers from these broths should be made on to MacConkey and Salmonella-Shigella agar, incubated for 426 427 18-24 hours, and examined. If no growth typical of Salmonella is noted, the agar plates should be incubated an additional 18-24 hours and again examined. If colonies typical of Salmonella are observed, further subculture on to 428 suitable differential media should be made for positive identification. Sensitive PCR tests are available for the 429 detection of Salmonella spp. in cultured material. If Salmonella is detected, the batch is determined to be 430 431 unsatisfactory.

432 3. Example of detection of Mycoplasma-contamination

4333.1.An example of a general specific procedure for detection exclusion of Mycoplasma434mycoides subsp. mycoides (where general testing is not sufficient)

435 Each batch of live viral vaccine, each lot of master seed virus (MSV), each lot of primary and master cell stock (MCS), 436 and all ingredients of animal origin not steam-sterilised should be tested for the absence of mycoplasmas. Solid and 437 liquid media that will support the growth of small numbers of test organisms, such as typical contaminating organisms Acholeplasma laidlawii, Mycoplasma arginini, M. fermentans, M. hyorhinis, M. orale, and M. synoviae should be 438 439 used. The nutritive properties of the solid medium should be such that no fewer than 100 CFU should occur with each test organism when approximately 100-200 CFUs are inoculated per plate. An appropriate colour change should 440 occur in the liquid media when approximately 20-40 CFUs of each test organism are inoculated. The ability of the 441 culture media to support growth in the presence of product should be validated for each product to be tested, and for 442 each new batch or lot of culture media. 443

444 One sample of each lot of vaccine, e.g. MSV or MCS, should be tested. Four plates of solid medium are inoculated 445 with 0.25 ml of the sample being tested, and 10 ml of the sample inoculated into 100 ml of the liquid medium. An alternative is to inoculate each of the plates with 0.1 ml and to inoculate 100 ml of liquid medium with 1 ml of the 446 sample being tested. Two plates are incubated at 35-37°C aerobically (an atmosphere of air containing 5-10% CO2 447 448 and adequate humidity) and two plates are incubated anaerobically (an atmosphere of nitrogen containing 5-10% CO2- and adequate humidity) for 14 days. On day 3 or day 4 after inoculation, 0.25 ml from the liquid media are 449 subcultured on to two plates of solid media. One plate is incubated aerobically and the second anaerobically at 35-450 37°C for 14 days. The subculture procedure is repeated on day 6, 7, or 8 and again on day 13 or 14. An alternative 451 452 method is to subculture on days 3, 5, 10, and 14 on to a plate of solid medium. All the subculture plates are incubated for 10 days except for the 14-day subculture, which is incubated for 14 days. Liquid media is observed every 2-3 453 days and, if any colour change occurs, has to be subcultured immediately. 454

455 **3.2. Interpretation of Mycoplasma test results**

- At the end of the incubation period (total 28 days), examine all the inoculated solid media microscopically for the presence of mycoplasma colonies. The test sample passes the test if the growth of mycoplasma colonies has occurred on the positive controls, and if growth has not occurred on any of the solid media inoculated with the test material. If at any stage of the test, more than one plate is contaminated with bacteria or fungi, or is broken, the test is invalid and should be repeated. If mycoplasma colonies are found on any agar plate, a suitable confirmatory test on the colonies should be conducted, such as PCR. Some mycoplasmas cannot be cultivated, in which case the MSV and MCS have to be tested using an indicator cell line such as Vero cells, DNA staining, or PCR methods.
- 463 Further detailed procedures can be found in Veterinary Medicinal Products, VICH GL34: 464 http://www.ema.europa.eu/docs/en_GB/document_library/Scientific_guideline/2013/03/WC500140352.pdf
- Prior to beginning testing it is necessary to determine that each batch of media promotes the growth of *M. mycoides* subsp. *mycoides* SC-(*Mmm*SC) type strain PG1. General mycoplasma broth and agar are used but contain porcine serum as a supplement. Each batch of broth and agar is inoculated with 10–100 CFU of *Mmm*SC. The solid medium is suitable if adequate growth of *Mmm*SC is found after 3–7 days' incubation at 37°C in 5–10% CO₂. The liquid medium is suitable if the growth on the agar plates subcultured from the broth is found by at least the first subculture. If reduced growth occurs another batch of media should be obtained and retested.
- 1 ml of cell or virus seed to be tested is inoculated into 9 ml of the liquid medium and 100 µl on to solid mycoplasma 471 agar. The volume of the product is inoculated so that it is not more than 10% of the volume of the medium. The liquid 472 473 medium is incubated at 37°C in 5–10% CO₂ and 100 µl of broth is subcultured on to agar at days 7, 14 and 21. The 474 agar plates are incubated at 37°C in 5–10% CO₂ for no fewer than 14 days, except those corresponding to day 21 subculture, which are incubated for 7 days. An un-inoculated mycoplasma broth and agar plate are incubated as 475 476 negative controls. For assessment of inhibitory substances, inoculate 1 ml of sample to be tested into 9 ml of the 477 liquid medium and 100 µl on to solid medium and add 10-100 CFU of MmmSC to each. Prepare positive control by inoculating 9 ml of mycoplasma broth and a mycoplasma agar plate with 10-100 CFU of MmmSC. Incubate as for 478 samples and negative controls. 479
- During incubation time, visually compare the broth of the positive control with sample present with the positive control broth and, if there is no inhibition of the organism either the product possesses no antimicrobial activity under the conditions of the test, or such activity has been satisfactorily eliminated by dilution. If no growth or reduced growth of *MmmSC* is seen in the liquid and solid medium with test sample when compared with the positive control, the product possesses antimicrobial activity, and the test is not satisfactory. Modifications of the conditions to eliminate the antimicrobial activity and repeat test are required.
- If antimicrobial activity is present it is necessary to dilute the test product further. Repeat the test above using 1.0 ml
 of sample in 39 ml of mycoplasma broth and then inoculate with 10–100 CFU of *MmmSC* and incubate as above. All
 broths and plates are examined for obvious evidence of growth. Evidence of growth can be determined by comparing
 the test culture with the negative control, the positive control, and the inhibition control.
- 490 If evidence of microbial growth is found in the test samples the contaminating bacterium will be identified and 491 confirmed as *Mmm*SC by <u>specific</u> PCR assay.

492 <u>3.2 General testing for exclusion of Mycoplasma sp.</u>

- 493 <u>General testing for exclusion of *Mycoplasma* sp. that are less fastidious may require up to 28 days in culture, using
 494 <u>general mycoplasma media. Some mycoplasmas cannot be cultivated, in which case the live biological sample will</u>
 495 <u>have to be tested using an indicator cell line such as Vero cells, DNA staining, or PCR methods.</u>
 </u>
- 496
 Further
 detailed
 procedures
 can
 be
 found
 in
 Veterinary
 Medicinal
 Products,
 VICH
 GL34:

 497
 http://www.ema.europa.eu/docs/en_GB/document_library/Scientific_guideline/2013/03/WC500140352.pdf
 Image: Constraint of the second second
- 498 <u>and</u>
- 499
 USDA SAM 910: https://www.aphis.usda.gov/animal_health/vet_biologics/publications/910.pdf, (both accessed 25

 500
 July 2023).

501 4. <u>Example of</u> detection of rickettsia and protozoa

502 There are no general test procedures for exclusion of rickettsia or protozoa. Procedures to exclude specific agents of 503 concern such as *Coxiella burnetti* (Q fever), *Ehrlichia canis, Trypanosoma evansi* and *Babesia caballi* can be found for example, in the Review of Published Tests to detect pathogens in veterinary vaccines Intended for Importation into
 Australia (Department of Agriculture [of Australia] [, Forest and Fisheries (2013]). The review is based on the reading and
 interpretation of applicable published papers from reputable journals and are regarded as examples of sensitive methods
 for detection of specified agents.

5084.1. An Example of a specific test protocol based on published methods for exclusion of Babesia509caballi and Theileria equi

Babesia caballi and Theileria equi can be cultured in vitro in 10% equine red blood cells (RBC) in supportive medium 510 supplemented with 40% horse serum and in a micro-aerophilic environment. Culture isolation of T. equi is more 511 sensitive than for B. caballi. Giemsa-stained blood smears are prepared from cultures daily for 7 days (Avarzed et 512 al., 1997; Ikadai et al., 2001). Babesia caballi is characterised by paired merozoites connected at one end. Theileria 513 514 equi is characterised by a tetrad formation of merozoites or 'Maltese cross'. Confirmation of the diagnosis is by PCR (see Chapter 2.5.8 Equine piroplasmosis). Molecular diagnosis is recommended for the testing of biological products 515 516 that do not contain whole blood or organs. Molecular diagnosis by PCR or loop-mediated isothermal amplification (LAMP) assay are the most sensitive and specific testing methods for detection of the pathogens of equine 517 piroplasmosis (Alhassan et al., 2007). 518

519 5. Example of detection of virus viruses in biological materials

In brief, general testing usually includes the use of continuous and primary cell lines of the source species, <u>e.g.</u>; cells of known susceptibility to <u>the-likely</u> viral contaminants, which are inoculated for <u>usually</u> a period of <u>up to 3</u>–4 weeks with weekly subcultures. Virus seeds also require testing on a primary cell line of the species in which the final product is intended. At Day 21 or 28, assessment of the monolayers is done using <u>H&E-appropriate histology</u> staining <u>procedures</u> to assess CPE₁ and haemadsorption with guinea-pig and chicken RBC to assess the presence of haemadsorbing agents. Note that general testing is useful as a screening tool though not sufficiently sensitive enough to detect all viruses of concern <u>to all countries</u>.

527 Specific testing requires test material to be inoculated on to sensitive, susceptible cells lines for the virus to be excluded; 528 the amplification process in cell culture is usually up to 28 days but depending <u>of on</u> the virus, may require longer culturing 529 times. Detection of specific viral contaminants is by recognition of CPE in conjunction with more sensitive antigen detection 530 or molecular tests such as FAT and PCR and ELISA after the amplification process in cell culture <u>is completed</u>.

531 All testing using cell lines to amplify for target viruses is contingent on the sensitivity of the cells for the target agent and 532 the ability to recognise the presence of the agent in the cells. The quality, characteristics, and virus permissibility profile of 533 cell lines in use should be determined as fit for purpose and appropriately maintained. Positive and negative controls should 534 be used at all passages of cell culture to determine sensitivity and specificity. Interference testing should be performed at 535 first pass to ensure that the test sample does not inhibit the growth of the virus being excluded for.

536 **5.1.** An example of general testing for the exclusion of viruses from virus and cell seed stocks 537 used in production of veterinary vaccines

538 If the test virus inoculum is cytopathogenic If a virus seed is known to cause cytopathic effect (CPE) in a permissive 539 cell line, the effect must be specifically neutralised without affecting the likelihood of isolation of the target agent. For 540 affected cell type, 1 ml of the test master (or working) virus seed (MVS) is thawed or reconstituted and neutralised with the addition of 1 ml mono specific antiserum. The serum must be shown to be free from antibodies against any 541 agents for which the test is intended to detect. Antiserum must should be tested for nonspecific inhibiting affects. For 542 a general test, this can be difficult to ascertain. Serum should be of sufficiently high titre to neutralise the seed virus 543 effectively with the use of an approximately equal volume or less of serum. A microplate block titration is used-useful 544 to determine the titre amount of the antiserum required to neutralise the MVS a known amount of concern. The 545 antiserum CPE causing virus seed. This is allowed to neutralise the MVS at 37°C for 1 hour. The MVS and antiserum 546 mixture is then inoculated on to a 75 cm² flask with appropriate cells. If the MVS is known to be high-titred or difficult 547 to neutralise, the blocking antiserum can be added to the growth medium at a final concentration done in the normal 548 549 conditions required of 1-2%.each test system (e.g. time, temperature, cell type etc.).

- 550 Master cell-If a virus seed is known to be high-titred or difficult to neutralise, antiserum can be added to the growth 551 medium in a test system at a final concentration of 1–2%.
- 552 Cell seed stocks do not require a neutralisation process.

555

556

557 558

559

560

561

562

563

564 565

566 567

568

569

570

571

572

573

574

575

576 577

578

579

580

581

582 583

584

585 586

587

588

589 590

591 592

593

594

595

596

597

598

599

5.1. Example of general testing procedures for the exclusion of viruses from virus and cell seed 553 stocks used in production of veterinary vaccines 554

5.1.1 Example of amplification in cell culture

The cells should be passaged weekly up to a 28-day period. Continuous and primary, 75 cm² area monolayers of the source species (and intended species as applicable) are infected with 1 ml of seed stocks and passaged weekly for between up to 21-28-days. Depending on the procedure followed, monolayers can be subcultured between passes or freeze/thawed to disrupt cells. Negative and positive controls should be also set up at each pass using the same cell population. Certain relevant viruses may be selected as indicators for sensitivity and interference (positive controls) but these will not provide validation for the broader range of agents targeted in general testing. The final culture is examined for cytopathology and haemadsorption.

5.1.2 Example of general detection procedures: cvtopathology

May-Grünwald-Giemsa or H&E staining procedures are used to assess for cytopathological changes associated with virus growth. Monolayers must have a surface area of at least 6 cm² and can be prepared on appropriate chambered tissue culture slides and incubated for 7 days. The plastic wells of the slides are removed leaving the rubber gasket attached to the slide. The slides are rinsed in Dulbecco's phosphate buffered saline (PBS), fixed in acetone, methanol or formalin depending on the stain used and placed on a staining rack. For May-Grünwald-Giemsa staining: the slides are stained for 15 minutes at room temperature with May-Grünwald stain diluted 1/5 with absolute methanol. The May-Grünwald stain is removed by inverting the slides. The slides are then stained for 20 minutes with Giemsa stain diluted 1/15 in deionised water. The Giemsa stain is removed by inverting the slides and rinsing them in deionised water for 10-20 seconds. The slides are air-dried and mounted with a coverslip using paraffin oil. The May-Grünwald-Giemsa stain differentially stains ribonucleoprotein (RNP); DNA RNP stains redpurple, while RNA RNP stains blue. The monolayers are examined with a conventional microscope for the presence of inclusion bodies, an abnormal number of giant cells, or other cytopathology attributable to a viral contaminant of the test product. The inoculated monolayers are compared with suitable control non-inoculated monolayers. If specific cytopathology attributable to an extraneous virus is found, results are reported, and additional specific testing may be conducted.

5.1.3 Example of general detection procedures: haemadsorption

Testing for haemadsorption uses requires the use of 75 cm² area monolayers established in tissue culture flasks after the 28-day passage period described above. Guinea-pig, chicken, and any other blood for use in this assay is collected in an equal volume of Alsever's solution and may be stored at 4°C for up to 7 days. Immediately prior to use, the stored erythrocytes are again washed by adding 5 ml of blood in Alsever's solution to 45 ml of calcium and magnesium-free PBS (PBSA) and centrifuging in a 50 ml centrifuge tube at 500 g for 10 minutes. The supernatant is aspirated, and the erythrocytes are suspended in PBSA and re-centrifuged. This washing procedure is repeated at least twice until the supernatant is clear. Erythrocytes from each species are combined by adding 0.1 ml of each type of packed blood cells to 100 ml of PBSA. The erythrocytes from different species may be kept separate or combined, as desired. To each flask, add 5 ml of the erythrocyte suspension, and incubate the flasks at 4°C for 30 minutes. Monolayers are washed twice with PBSA and examined for haemadsorption. If no haemadsorption is apparent, 5 ml of the fresh erythrocyte suspension is added to each flask; the flasks are incubated at 20-25°C (room temperature) for 30 minutes, rinsed as before, and examined for haemadsorption. Separate flasks may be used for each incubation temperature if desired. Monolayers are examined for the presence of haemadsorption using an illuminated light box and microscopically. Non-inoculated monolayers are used as negative controls. The PBSA and fresh erythrocytes should prevent most nonspecific haemadsorption from occurring. If specific haemadsorption attributable to an extraneous agent is found, results are reported, and additional specific testing may be conducted.

Specific testing requires specialised test procedures that are sensitive to amplifying a particular agent in 600 culture and then detection of that agent by means of fluorescence, antigen capture ELISA or PCR; 601 602 whichever is more sensitive. Specific testing is usually required when general procedures are not 603 adequate for effective exclusion of more fastidious, viruses Some examples are listed in Table 1.

6045.2. An Examples of specific virus agent exclusion testing from of biologicals used in the605production of veterinary vaccines

5.2.1. Example of porcine epidemic diarrhoea virus (PEDV)

- Trypsin presence is required at inoculation and in the culture medium for isolation of porcine epidemic 607 diarrhoea virus (PEDV) in Vero cells (CCL81, ATCC) to ensure the virus can enter host cells. Just 608 confluent monolayers (100%) are required; as under confluent monolayers (<90%) are more sensitive 609 to the presence of trypsin and will be destroyed well before the 7 days required for each passage in 610 culture. An over confluent or aging monolayer will not be sensitive for growth of PEDV. Maintenance 611 media (MM) formulation consists of Earle's MEM (minimal essential medium) (with 5.6 M HEPES IN-612 2-hydroxyethylpiperazine, N-2-ethanesulphonic acid] and glutamine) + 0.3% Tryptose phosphate 613 broth, 0.02% yeast extract and 4 µg/ml TPCK treated trypsin. The addition of the trypsin into the MM 614 615 should occur on the day the media is to be used.
 - Prior to inoculation, confluent 75 cm² monolayers are washed twice with the MM (with trypsin added) to remove growth media containing FCS. Virus or cell seed (1 ml) is added with 1 ml of MM to each monolayer; incubate at 37°C for 2 hours, then add 30 ml/flask of MM. Negative control monolayers of the same size are set up prior to inoculation of test material. Positive and interference controls are set up last, and where possible, in a separate laboratory area-to avoid contamination. Assessment for sensitivity and interfering substances requires assessment use of PEDV reference virus of known titre. A control for interference using co-inoculation of test sample and PEDV needs only to be set up on the first pass. Positive controls must should be set up at every pass to ensure each monolayer used gives expected sensitivity. PEDV virus is titrated in log dilutions starting at 10⁻¹ to 10⁻⁶ in MM (depending of <u>on</u> the endpoint titre of reference virus) in duplicate rows of 6 wells of a 24-well tissue culture plate. For the interference test, PEDV is titrated in the same dilution series but using MM spiked with a 10% volume of test material. Decant off the growth media and discard. Wash plates to ensure no FCS is present. Two washes using approximately 400 µl/well MM (with trypsin added) are sufficient.
- 630Add 100 μl of diluted virus on to each of two duplicate wells. Rock inoculated plates to distribute the631inoculum evenly over the surface of the monolayer. Incubate at 37°C with 5% CO2 for 2 hours then632add <u>a further 1</u> ml volumes/well of MM.
- After 7 days, 75 cm² monolayers have cells disrupted using two freeze-thaw cycles at -80°C. Positive 633 control plates are read for end-point titres, and these are compared with virus in the presence of test 634 635 material to ensure titres are comparable and interference has not occurred. Freeze-thaw lysates are clarified at 2000 g for 5 minutes and re-passed on to newly formed monolayers as for the first 636 passage. Passages are repeated until a total of four passages are completed at which point cell 637 lysates are assessed by PCR for detection of PEDV and day 7 monolayers in 24-well plates are fixed 638 639 and stained by IFA-for FAT. If a seed virus is to be tested and requires neutralisation using antiserum, extra care in the isolation of PEDV needs to be considered. Trypsin is rendered inactive in the 640 presence of serum proteins and without trypsin present, PEDV is unable to grow in cell culture grows 641 poorly, or not at all. Washing off the inoculum with two MM washes is required after an extended 642 adsorption time of up to 4 hours to ensure acceptable sensitivity. 643

644 645

606

616

617

618

619 620

621

622

623 624

625

626

627

628 629

$J-\underline{H}$. INFORMATION TO BE SUBMITTED WHEN APPLYING FOR AN IMPORT LICENCE

When undertaking risk analysis for biologicals, Veterinary Authorities should follow the Terrestrial Manual the manufacturer 646 should follow the requirements of the importing country. Requirements for each importing country should be accessible 647 and published online. The manufacturer or the Veterinary Authority of the exporting country should make available detailed 648 649 information, in confidence if-as necessary, on the source of the materials used in the manufacture of the product (e.g. substrates). They should make available details of the method of manufacture (and where appropriate inactivation) of the 650 substrates and component materials, the quality assurance procedures for each step in the process, final product testing 651 652 regimes, and the pharmacopoeia with which the product must conform in the country of origin. They should also make 653 available challenge organisms, their biotypes and reference sera, and other means of appropriate product testing.

- 654 <u>For detailed examples of a risk-based assessment of veterinary biologicals for import into a country refer to:</u>
- European Commission (2015). The Rules Governing Medicinal Products in the European Union. Eudralex. Volume 6.
 Notice to applicants and regulatory guidelines for medicinal products for veterinary use

- Department of Agriculture, Forest and Fisheries of Australia (2021b). Live veterinary vaccines Summary of information
 required for biosecurity risk assessment, Version 6 and Inactivated veterinary vaccines, Version 8.
- <u>Outline of the Regulatory System of Veterinary Drugs in Japan (2015) Assurance of the Quality, Efficacy, and Safety</u> Based on the Law for Ensuring the Quality, Efficacy, and Safety of Drugs and Medical Devices.

661 When applying for an import licence other regulatory requirements may need to be addressed depending on the type of 662 sample and if the sample needs to be shipped out of country to a testing laboratory. For example, cell seeds may come 663 under certain requirements for permits such as the Convention for International Trade in Endangered Species of Wild 664 Fauna and Flora (CITES), where a cell line is derived from an endangered species, e.g. the cell line and its derivatives. 665 Applying for such a permit is time consuming and requires input from both the exporting and importing country.

<u>Genetically modified organisms (GMOs) are becoming more frequent in use with changes in manufacturing technologies</u>
 and specialised, time-consuming procedures need to be in place. A laboratory that accepts a GMO product for testing shall
 follow the procedures of the Office of the Gene Regulator (OGTR) to allow the GMO to be dealt with.

669

I. RISK ANALYSIS PROCESS

670 Risk analysis should be as objective and transparent as possible and should be performed in accordance with Section 2 671 of the *Terrestrial Code*, and certification in line with Section 5 of the *Terrestrial Code*. Of necessity, assessment of the 672 country and commodity factors and risk reduction measures will be based largely on manufacturers' data. These data 673 depend on quality assurance at all stages of manufacture, rather than on testing of the final product alone.

Domestic exposure may be influenced by the approved usage of the product. Veterinary Authorities may place limits on usage of some products (e.g. restricting usage to institutions of appropriate biosecurity).

676 L_J. BIOCONTAINMENT

677 Suitable biocontainment may be necessary for many forms of biologicals. In particular, the importation of exotic micro-678 organisms should be carried out in accordance with Chapter 1.1.4 *Biosafety and biosecurity: standard for managing* 679 *biological risk in the veterinary laboratory and animal facilities.*

Laboratories using high risk agents should have well researched and documented risk assessments in place prior to
 working with such agents to ensure the safety of their staff and laboratory.

682

REFERENCES

ALHASSAN A., GOVIND Y., TAM N.T., <u>THEKISOE O.M.</u>, YOKOYAMA N., INOUE N. & IGARASHI I. (2007). Comparative evaluation of
 the sensitivity of LAMP, PCR and in vitro culture methods for the diagnosis of equine piro-plasmosis. *Parasitol. Res.*, **100**,
 1165–1168.

AUSTRALIAN QUARANTINE POLICY AND REQUIREMENTS FOR THE IMPORTATION OF LIVE AND NOVEL VETERINARY BULK AND FINISHED VACCINES (1999). Available online at:

- https://www.agriculture.gov.au/sites/default/files/sitecollectiondocuments/ba/memos/1999/animal/99-085acleaned.pdf
 (Accessed 25 July 2023).
- Avarzed A., Igarashi I., Kanemaru T., Hirumi, K., Omata T., SaitoY., Oyamada A., Nagasawa H., Toyoda Y. & Suzuki N.
 (1997). Improved *in vitro* cultivation of *Babesia caballi*. J. Vet. Med. Sci., 59, 479–481.
- BARONE P., WIEBE M.E., LEUNG J.C., HUSSEIN I.T.M., KEUMURIAN F.J., BOURESSA J., BRUSSEL A., CHEN D., CHONG M.,
 DEHGHANI H., GERENTES L., GILBERT J., GOLD D., KISS R., KREIL T.R., LABATUT R., LI Y, MÜLLBERG J., MALLET L., MENZEL C.,
 MOODY M., MONPOEHO S., MURPHY M., PLAVSIC M., ROTH N.J., ROUSH D., RUFFING M., SCHICHO R., SNYDER R., STARK D.,
 ZHANG C., WOLFRUM J., SINSKEY A.J. & SPRINGS S.L. (2020). Viral contamination in biologic manufacture and implications
 for emerging therapies. *Nat. Biotechnol.*, 38, 563–572.

BAYLIS S.A., FINSTERBUSCH T., BANNERT N., BLUMEL J. & MANKERTZ A. (2011). Analysis of porcine circovirus type 1 detected in Rotarix vaccine. *Vaccine*, **29**, 690–697. 699 CODE OF FEDERAL REGULATIONS (OF THE UNITED STATES OF AMERICA) (CFR) (2015). Subchapter E. Viruses, serums, toxins 700 and analogous products; organisms and vectors. In: Code of Federal Regulations, Animals and Animal Products. Title 9, 101-124. US Government Printing Washington 701 Parts Office, DC. USA. Available online: 702 https://www.law.cornell.edu/cfr/text/9/part-113 (Accessed 25 July 2023).

DEPARTMENT OF AGRICULTURE, <u>WATER AND ENVIRONMENT</u> (OF AUSTRALIA) (2013). Review of Published Tests to Detect Pathogens in Veterinary Vaccines Intended for Importation into Australia, Second Edition. CC BY 3.0. Available online at:

705 <u>https://www.agriculture.gov.au/biosecurity-trade/policy/risk-analysis/animal/review-of-published-tests-to-detect-pathogens</u>
 706 <u>(Accessed 25 July 2023).</u>

DEPARTMENT OF AGRICULTURE, WATER AND ENVIRONMENT (OF AUSTRALIA) (2021a). Inactivated veterinary vaccines Summary
 of information required for biosecurity risk assessment. Version 8. Available online at:
 https://www.agriculture.gov.au/sites/default/files/sitecollectiondocuments/aqis/importing/bio-products/vet-vaccines/inact vet-vacs.pdf (Accessed 25 July 2023).

- 711 <u>DEPARTMENT OF AGRICULTURE, WATER AND ENVIRONMENT (OF AUSTRALIA) (2021b). Live veterinary vaccines. Summary of</u> 712 information required for biosecurity risk assessment. Version 6. Available online at:
- https://www.awe.gov.au/sites/default/files/sitecollectiondocuments/aqis/importing/bio-products/vet-vaccines/live-vetvaccines.doc (Accessed 25 July 2023).
- EUROPEAN COMMISSION (2006-2015). The Rules Governing Medicinal Products in the European Union. Eudralex. Volumes
 4–9. European Commission, DG Health and Food Safety, Public health, EU Pharmaceutical information, Eudralex.
 Available online at: http://ec.europa.eu/health/documents/eudralex/index_en.htm (accessed 25 July 2023).

EUROPEAN MEDICINES AGENCY SCIENCES MEDICINES HEALTH (2016). EMA/CVMP/IWP/251741/2015. Reflection paper on 718 methods found suitable within the EU for demonstrating freedom from extraneous agents of the seeds used for the 719 720 production of immunological veterinary medicinal products. Available online at: 721 https://www.edgm.eu/en/d/95422?p | back url=%2Fen%2Fsearch-

- 722
 edqm%3Fq%3DEMA%252FCVMP%252FIWP%252F251741%252F2015%26delta%3D40%26category%3D302281

 723
 (Accessed 25 July 2023).
- EUROPEAN PHARMACOPOEIA 8.2. (2014) (2021-2023). European Directorate for the Quality of Medicines and Health Care
 (EDQM), Council of Europe, Strasbourg, France. Available online at http://online.edgm.eu/ (accessed 8 March 2017)
 Current version available online at European Directorate for the Quality (accessed 8 March 2017)
 Current version available online at European Directorate for the Quality (accessed 4 March 2017)
 Current version available online at European Directorate for the Quality (accessed 4 July 2022) Current version available online at European Directorate for the Quality (accessed 4 July 2022) Current version available online at European Directorate for the Quality (accessed 4 July 2022) Current version available online at European Directorate for the Quality (Accessed 4 July 2022) Current version available online at European Directorate for the Quality (Accessed 25 July 2023).
- FARSANG A. & KULCSAR G. (2012). Extraneous agent detection in vaccines a review of technical aspects. *Biologicals*, 40,
 225–230.
- HARE W.C.D. (1985). Diseases Transmissible by Semen and Embryo Transfer Techniques. OIE Technical Series No. 4.
 World Organisation for Animal Health (OIE), Paris, France.
- HODINKA R.L. (2013). Point: is the era of viral culture over in the clinical microbiology laboratory? J. Clin. Microbiol., **51**, 2–4.
- Ikadai H., Martin M.D., Nagasawa H., Fujisaki K., Suzuki N., Mikami T., Kudo N., Oyamada T. & Igarashi I. (2001). Analysis
 of a growth-promoting factor for *Babesia caballi* cultivation. *J. Parasitol.*, 87, 1484–1486.
- MARCUS-SECURA C., RICHARDSON J.C., HARSTON R.K., SANE N. & SHEETS R.L. (2011). Evaluation of the human host range
 of bovine and porcine viruses that may contaminate bovine serum and porcine trypsin used in the manufacture of biological
 products. *Biologicals*, **39**, 359–369.
- 740 MACKAY D. & KRIZ N. (2010). Current challenges in viral safety and extraneous agents testing. Biologicals, 38, 335–337.

MILLER P.J., AFONSO C.L., SPACKMAN E., SCOTT M.A., PEDERSEN J.C., SENNE D.A., BROWN J.D., FULLER C.M., UHART M.M.,
 KARESH W.B., BROWN I.H., ALEXANDER D.J. & SWAYNE D. (2010). Evidence for a new avian paramyxovirus serotype 10
 detected in rockhopper penguins from the Falkland Islands. *J. Virol.*, 84, 11496–11504.

NEVEROV A. & CHUMAKOV K. (2010). Massively parallel sequencing for monitoring genetic consistency and quality control
 of live viral vaccines. *Proc. Natl Acad. Sci.* USA, **107**, 20063–20068.

- ONIONS D. & KOLMAN J. (2010). Massively parallel sequencing, a new method for detecting adventitious agents. *Biologicals*,
 38, 377–380.
- OUTLINE OF THE REGULATORY SYSTEM OF VETERINARY DRUGS IN JAPAN (2015). Assurance of the Quality, Efficacy, and Safety
 Based on the Law for Ensuring the Quality, Efficacy, and Safety of Drugs and Medical Devices.
 https://www.maff.go.jp/nval/english/regulatory/pdf/Outline RegulatorySystem VMPs.pdf (Accessed 25 July 2023).
- ROSSEEL T., LAMBRECHT B., VANDENBUSSCHE F., VAN DEN BERG T. & VAN BORM S. (2011). Identification and complete genome
 sequencing of paramyxoviruses in mallard ducks (*Anas platyrhynchos*) using random access amplification and next
 generation sequencing technologies. *Virol. J.*, 8, 463.
- SHEETS R., LOEWER, J., RAYDCHAUDHURI G. & PETRICCIANI J. (2012). Adventitious agents, new technology, and risk assessment, 19–20 May 2011, Baltimore, MD. *Biologicals*, **40**, 162–167.
- TELLEZ S., CASIMIRO R., VELA A.I., FERNANDEZ-GARAYZABAL J.F., EZQUERRA R., LATRE M.V., BRIONES V., GOYACHE J., BULLIDO
 R., ARBOIX M. & DOMINGUEZ L. (2005). Unexpected inefficiency of the European pharmacopoeia sterility test for detecting
 contamination in clostridial vaccines. *Vaccine*, 24, 1710–1715.
- VAN BORM S., ROSSEEL T., STEENSELS M., VAN DEN BERG T. & LAMBRECHT B. (2013). What's in a strain? Viral metagenomics
 identifies genetic variation and contaminating circoviruses in laboratory isolates of pigeon paramyxovirus type 1. *Virus Res*,
 171, 186–193.
- VICTORIA J.G., WANG C., JONES M.S., JAING C., MCLOUGHLIN K., GARDNER S. & DELWART E.L. (2010). Viral nucleic acids in live-attenuated vaccines: detection of minority variants and an adventitious virus. *J. Virol.*, **84**, 6033–6040.
- WANG J., LUNT R., MEEHAN B. & COLLING A. (2014). Evaluation of the potential role of next-generation sequencing (NGS) in
 innocuity testing. Technical Report CSIRO, Australian Animal Health Laboratory.
- WORLD HEALTH ORGANIZATION (WHO) (1998). WHO Expert Committee on Biological Standardization. World Health Organization Technical Report Series, Report No. 858. World Health Organization, Geneva, Switzerland.
- WORLD HEALTH ORGANIZATION (WHO) (2012). WHO Expert Committee on Biological Standardization. World Health
 Organization Technical Report Series, Report No. 964. World Health Organization, Geneva, Switzerland. Available online
 at: https://apps.who.int/iris/bitstream/handle/10665/89142/9789241209779 eng.pdf (Accessed 25 July 2023).
- 771World Health Organisation (WHO) (2015) Expert Committee on Biological Standardization Sixty-fifth report. Technical
report series 993, 2015. Annex 2 Scientific principles for regulatory risk evaluation on finding an adventitious agent in a
marketed vaccine. Available online at: https://cdn.who.int/media/docs/default-source/biologicals/cell-
773
substrates/annex2 adventious agent in marketed vaccine_eng.pdf?sfvrsn=e5de04a 3&download=true (Accessed25
July 2023)
- 776

FURTHER READING

- 777 Details of methods and culture media <u>will can</u> be found in the following books and also in commercial catalogues.
- BARROW G.I. & Feltham R.K.A., eds. (1993). Cowan and Steel's Manual for the Identification of Medical Bacteria, Third
 Edition. Cambridge University Press, Cambridge, UK.
- COLLINS C.H., LYNE P.M. & GRANGE J.M., eds. (1995). Collins and Lyne's Microbiological Methods, Seventh Edition.
 Butterworth Heinemann, Oxford, UK.
- MURRAY P.R., ED. (2003). Manual of Clinical Microbiology, Eighth Edition. American Society for Microbiology Press,
 Washington DC, USA.
- 784

785

786

NB: FIRST ADOPTED IN 1989. MOST RECENT UPDATES ADOPTED IN 2017.

INTRODUCTION

The WOAH Validation Recommendations provide detailed information and examples in support of the WOAH Validation Standard that is published as Chapter 1.1.6 Principles and methods of Validation of diagnostic assays for infectious diseases of terrestrial animals this Terrestrial Manual, or Chapter 1.1.2 of the Aquatic Manual. The Term "WOAH Validation Standard" in this chapter should be taken as referring to those chapters.

Estimation of measurement uncertainty (MU), sometimes termed measurement imprecision, is a 13 14 requirement for testing laboratories based on international quality standards such as ISO/IEC 17025-2005. 2017 General requirements for the competence of testing and calibration laboratories (ISO/IEC 17025). The 15 measurement process for detection of an analyte in a diagnostic sample is not entirely reproducible and 16 hence there is no exact value that can be associated with the measured analyte. Therefore, the result is 17 18 most accurately expressed as an estimate together-with an associated level of imprecision level. This imprecision is the measurement uncertainty (MU). MU is limited to the measurement process of quantitative 19 tests. The approach described here is known as "top-down" or "control sample" because it uses a weak 20 positive control sample and expresses the MU result at the cut-off, where it most matters. It is not a question 21 of whether the measurement is appropriate and fit for whatever use to which it may be applied. It is not an 22 alternative to test validation but is rightly considered a component of that process (see the WOAH Validation 23 Standard, chapter 1.1.6 Section B.1.1 Repeatability). 24

25

7

A. THE NECESSITY OF DETERMINING MU

To assure compliance with ISO/IEC 17025-2005-2017 requirements, national accreditation bodies for diagnostic testing 26 27 laboratories require laboratories to calculate MU estimates for accredited test methods that produce quantitative results, e.g. optical densities (OD), percentage of positivity or inhibition (PP, PI), titres, cycle threshold (CT) values, etc. This 28 includes tests where numeric results are calculated and then are expressed as a positive or negative result at a cut-off 29 30 value. For the purpose of estimating MU in serology and reverse transcriptase polymerase chain reaction (RT-PCR), suitable statistical measures are mean target values ± 2 standard deviations (SD), which is approximately equal to a 95% 31 confidence interval (CI), relative standard deviation (RSD = SD / mean of replicates) and coefficient of variation (CV = RSD 32 33 × 100%). Examples provided below assume normal distribution of data. The concept of MU does not apply to strictly binary 34 (qualitative) results (positive or negative).

1. Samples for use in determining MU

Repeatability is the level of agreement between results of replicates of a sample both within and between runs of the same 36 test method in a given laboratory. During assay development, repeatability is estimated by evaluating variation in results 37 38 of independent replicates from a minimum of three (preferably five) samples representing analyte activity within the operating range of the assay (see-the WOAH Validation Standard, Chapter 1.1.6 Validation of diagnostic assays for 39 infectious diseases of terrestrial animals, Sections A.2.5 Robustness and B.1.1 Repeatability, and Chapter 2.2.6 Selection 40 and use of reference samples and panels, Section 3.1.A.4.2). Typically, the variation in replicate results is expressed as 41 42 RSD or CV. The significant feature is that repeatability studies can be used to define the expected precision of the assay 43 in the detection of a range of analyte concentrations.

44 The use of internal quality or process controls over a range of expected results has become part of daily quality control and quality assurance operations of accredited facilities (see the WOAH Validation Standard, chapter 1.1.6, Sections A.2.6 45 Calibration of the assay to standard reagents and B.5.1 Monitoring the assay, and Chapter 2.2.6, Section 1.4-C.1). These 46 47 results provide a continuous monitor relative to different aspects of repeatability, e.g. intra- and inter-assay variation, intraand inter-operator variation and intra- and inter-batch variation, which, when subjected to statistical analysis, provide an 48 expression of the level of robustness (precision) of a test procedure. The monitoring of assay quality control parameters 49 50 for repeatability provides evidence that the assay is or is not performing as expected. For control samples to provide valid inferences about assay precision, they should be treated in exactly the same way as test samples in each run of the assay, 51 52 e.g. including sample preparation such as extraction steps or dilution of serum samples for an antibody enzyme-linked immunosorbent assay (ELISA). 53

The variation of the results for control samples can also be used as an estimate of those combined sources of uncertainty and is called the "top-down" approach. This approach recognises that the components of precision will be manifest in the ultimate measurement. So monitoring the precision of the measurement over time will effectively show the combined effects of the imprecision associated with component steps.

58 The imprecision or uncertainty of the measurement process associated with a test result becomes increasingly more important the closer the test value is to the diagnostic cut-off value. This is because an interpretation is made relative to 59 the assay threshold regarding the status of the test result as positive, negative, or inconclusive (as will be described in the 60 following example). In this context, low weak positive samples, like those used in repeatability studies or as the low weak 61 62 positive control, are most appropriate for estimation of MU. The rationale being that MU, which is a function of assay precision, is most critical at decision-making points (i.e. thresholds or cut-offs), which are usually near the lower limit of 63 detection for the assay. In this chapter, the application of MU with respect to cut-off (threshold) values, whether 64 recommended by test-kit manufacturers or determined in the diagnostic laboratory, is described. 65

MU, using the top-down approach, ideally requires long-term accumulated data from a weak positive control sample after
 multiple test runs over time, with multiple operators and variable conditions. The examples given below are based on 10
 data points but higher numbers will increase robustness.

69 2. Example of MU calculations in ELISA serology

For most antibody detection tests, it is important to remember that the majority of tests are measurements of antibody activity relative to a threshold against which a dichotomous interpretation of positive or negative is applied. This is important because it helps to decide where application of MU is appropriate. In serology, uncertainty is frequently most relevant at the threshold between positive and negative determinations. Results falling into this zone are also described as intermediate, inconclusive, suspicious or equivocal (see the WOAH Validation Standard, chapter 1.1.6, Section B.2.4 *Selection of a cut-off (threshold) value for classification of test results*).

A limited data set from a competitive ELISA for antibody to avian influenza virus is used as an example of a "top-down" approach for serology. A low-weak positive control sample was used to calculate MU at the cut-off level $\frac{32}{2}$.

78 **2.1. Method of expression of MU**

As the uncertainty is to be estimated at the threshold, which is not necessarily the reaction level of the <u>low-weak</u> positive control serum, the relative standard deviation (RSD), or coefficient of variation (CV), if expressed as a percentage, provides a convenient transformation:

<u>32 The Australian Government, Department of Agriculture, Fisheries and Forestry, has compiled worked examples for a number of diagnostic tests Available online at: https://www.agriculture.gov.au/agriculture-land/animal/health/laboratories/tests/measurementuncertainty (accessed 22 June 2023)</u>

RSD ((X)	=	SD	(X)	/ (X)

To simplify assessment, the transformed result is regarded as the assay output result, which is the averaged across 82 the number of replicates (\overline{X}). In the case of this example, a competitive ELISA, results are "normalised" (as defined 83 in the WOAH Validation Standard, chapter 1.1.6, Section A.2.7 (Normalising' test results to a working standard) to a 84 85 working standard by forming a ratio of all optical density (OD) values to the OD result of a non-reactive (negative) control (OD_N). This ratio is subtracted from 1 to set the level of antibody activity on a positive correlation scale; the 86 87 greater the level, the greater the calculated value. This adjusted value is expressed as a per cent and referred to as the percentage inhibition or PI value. So for the low-weak positive control serum (ODL), the transformation to obtain 88 the per cent inhibition values for the low-weak positive control (PIL) is: 89

90 $PI_L = 100 \times [1 - {OD_L / OD_N}]$

91 The relative standard deviation becomes:

92

 $RSD(PI_L) = SD(PI_L)/(PI_L)$

93 **2.2. Example**

A limited data set for the AI competitive ELISA example is shown below. In the experiment, the operator tested the low-weak positive control serum ten times in the same run. Ideally in the application of this "top down" method, a larger data set would be used, which would enable accounting for effects on precision resulting from changes in operator and assay components (other than only the control serum).

98

99

Table 1. Top-down or control sample approach for an influenza antibody C-ELISA

Test	PI (%)				
1	56				
2	56				
3	61				
4	64				
5	51				
6	49				
7	59				
8	70				
9	55				
10	42				

Mean PI = 56.3; Std Dev (SD) = 7.9; Assays (n) = 10

100 2.3. Calculating uncertainty

- 101 From the limited data set,
- 102 RSD (PI_L) = SD/Mean 7.9/56.3 = 0.14 (or as coefficient of variation = 14%)

Expanded uncertainty (U) is the statistic defining the interval within which the value of the measure and is believed to lie within a specified level of confidence, usually 95%. Expanding the uncertainty is done by multiplying the RSD (PI_L) by a factor of 2; this allows the calculation of an approximate 95% confidence interval around the threshold value (in this case at PI = 50%), assuming normally distributed data.

- 107 U (95%CI) = 2 × RSD = 0.28
- 108 This estimate can then be applied at the threshold level

109 95% CI = 50 ± (50 × 0.28) = 50 ± 14%

110 **2.4. Interpretation**

111 Any positive result (PI > 50%) that is less than 64% is not positive with 95% confidence. Similarly, a negative result 112 (PI < 50%) that is higher or equal to a PI of 36 is not negative at the 95% confidence level. This zone of lower 113 confidence may correlate with the "grey zone" or "inconclusive/suspect zone" for interpretation that should be 114 established for all tests (Greiner *et al., 1995*).

115 3. Example of MU calculation in molecular tests

116 <u>3.1. Example</u>

117For real-time PCRs, replicates of positive controls with their respective cycle threshold (CT) values can be used to118estimate MU using the top-down approach (Newberry & Colling, 2021). The method of expression follows the same119formula as for the ELISA example above. This example uses data from replicate runs of a weak positive control120sample (10 runs) of an equine influenza hydrolysis probe assay.

<u>Table 2. Top-down or control sample approach for an equine influenza TaqMan A assay</u>

<u>Test</u>	<u>Ct value</u>
<u>1</u>	<u>33.60</u>
2	<u>33.20</u>
<u>3</u>	<u>33.96</u>
<u>4</u>	<u>33.18</u>
<u>5</u>	<u>33.96</u>
<u>6</u>	<u>32.72</u>
<u>7</u>	<u>33.57</u>
<u>8</u>	<u>33.45</u>
<u>9</u>	<u>32.80</u>
<u>10</u>	<u>33.20</u>

122

121

<u>Mean = 33.36; Std Dev (SD) = 0.43; Assay n=10</u>

- 123 <u>3.2. Calculating uncertainty</u>
- 124 <u>From the limited data set,</u>
- 125 RSD (PlL) = SD/Mean 0.43/33.36 = 0.0128 (or as coefficient of variation = 1.28%)
- Expanded uncertainty (U) is the statistic defining the interval within which the value of the measure and is believed to lie within a specified level of confidence, usually 95%. Expanding the uncertainty is done by multiplying the RSD (PL) by a factor of 2; this allows the calculation of an approximate 95% confidence interval around the threshold value (in this case at Ct value = 37), assuming normally distributed data.
- 130 <u>U (95%CI) = 2 × RSD = 0.0255</u>
- 131 This estimate can then be applied at the threshold level
- 132 95% CI = $37 \pm (37 \times 0.0255) = 37 \pm 0.94$
- 133The mean cycle threshold (Ct) value after 10 runs is 33.36 and the standard deviation is 0.43. The relative standard
deviation is 0.0128. The expanded uncertainty (95% Cl) is 2 × the relative standard deviation = 0.0255. Measurement
of uncertainty (MU) is most relevant at the cut-off (Ct = 37) and can be applied by multiplication (37 × 0.0255 = 0.94).
136136Subtraction from the threshold (37-0.94) provides the lower 95% confidence limit (Ct = 36.06) and addition (37+0.94)
the upper 95% confidence limit (Ct = 37.94).

138 <u>3.3. Interpretation of the results</u>

- 139Any positive result (Ct < 37) that is higher than 36 Ct is not positive with 95% confidence. Similarly, any negative</th>140result (Ct > 37) that is less than 38 Ct is not negative with 95% confidence.
- 141

152

171

B. OTHER APPLICATIONS

The top-down approach should be broadly applicable forto a range of diagnostic tests including molecular tests. For the 142 calculation of tests using a typical two-fold dilution series for the positive control such as virus neutralisation, complement 143 fixation and haemagglutination inhibition tests geometric mean titre (i.e. mean and SD of log base 2 titre values) of the 144 positive control serum should be calculated. Relative standard deviations based on these log scale values may then be 145 applied at the threshold (log) titre, and finally transformed to represent the uncertainty at the threshold. However, in all 146 cases, the approach assumes that the variance about the positive control used to estimate the RSD is proportionally similar 147 at the point of application of the MU, for example at the threshold. If the RSD varies significantly over the measurement 148 scale, the positive control serum used to estimate the MU at the threshold should be selected for an activity level close to 149 that threshold. The Australian Government, Department of Agriculture, Fisheries and Water Resources Forestry, has 150 151 compiled worked examples for a number of diagnostic tests (see footnote 1), (DAFF, 2010), which are available online at:

http://www.agriculture.gov.au/animal/health/laboratories/tests/worked-example-measurement

For quantitative real-time PCRs (qPCR) replicates of positive controls with their respective cycle threshold (CT) values can
 be used to estimate MU using the top-down approach.

155 Other approaches and variations have been described, i.e. for serological tests (Dimech *et al.*, 2006; Goris *et al.*, 2009; 156 Toussaint *et al.*, 2007). Additional work and policy <u>Central</u> documents are available from the National Pathology 157 Accreditation Advisory Group and Life Science. The central document to MU isare the Guide to the expression of 158 uncertainty in measurement (GUM), ISO/IEC Guide, (1995) and Eurachem/CITAC Guide, 2012 CG 4: Quantifying 159 uncertainty in analytical measurement.

160 Scope and limitations of the top-down approach

Methods for quantifying uncertainty (addressing MU) for tests vary. When estimating MU for quantitative, biologically based 161 162 diagnostic tests, where variations in the substrate or matrix have large and unpredictable effects, a top-down approach is recommended (Dimech et al., 2006; Eurachem 2012; Goris et al., 2009; ISO/IEC Guide 98-3:2008; Newberry & Colling, 163 2021; Standards Council of Canada, 2021; and footnote 1). The advantage of this method is that quality control data are 164 165 generated during normal test runs and can be used to estimate the precision of the assay and express it at the cut-off. The 166 application at the cut-off depends on the performance of the test at different analyte concentrations, e.g. variation is likely 167 to increase at higher diluted samples. The top-down approach does not identify individual contributors to measurement 168 uncertainty but rather provides an overall estimate. Measurement uncertainty does not replace test validation; however, 169 the validation process includes assessments of repeatability through quality control samples which facilitate calculation of 170 <u>MU.</u>

172 AMERICAN ASSOCIATION FOR LABORATORY ACCREDITATION (A2LA). Policy on estimating measurement uncertainty for life 173 science testing labs https://portal.a2la.org/policies/A2LA_P103b.pdf (accessed 22 November 2018).

REFERENCES

- 177
 AUSTRALIAN GOVERNMENT DEPARTMENT OF HEALTH AND ACEING (2007). Requirements for the estimation of measurement 178
 uncertainty,
 National
 Pathology
 Accreditation
 Advisory
 Group.

 179
 http://www.health.gov.au/internet/main/publishing.nsf/Content/B1074B732F32282DCA257BF0001FA218/\$File/dhaeou.p
 180
 df (accessed 22 November 2018).
- 181 DIMECH W., FRANCIS B., KOX J. & ROBERTS G. (2006). Calculating uncertainty of measurement for serology assays by use 182 of precision and bias. *Clin. Chem.*, **52**, No. 3, 526–529.

Australian Government, Department of Agriculture and WATER RESOURCES.<u>.</u> Worked <u>examples of measurement uncertainty</u>
 <u>http://www.agriculture.gov.au/animal/health/laboratories/tests/worked-example-measurement</u> (accessed 22 November
 2018).

- 183 <u>ELLISON S.L.R. & WILLIAMS A. (EDS) (2012). Eurachem/CITAC Guide CG 4: Quantifying uncertainty in analytical</u>
 184 <u>measurement, 3rd Edition. Eurachem, Teddington, United Kingdom, Section 3. Available at:</u>
 185 <u>www.eurachem.org/images/stories/Guides/pdf/QUAM2012 P1.pdf (Accessed 10 June 2023).</u>
- GORIS N., VANDENBUSSCHE F., HERR, C., VILLERS, J., VAN DER STEDE, Y. & DE CLERCQ K. (2009). Validation of two real-time
 PCR methods for foot-and-mouth disease diagnosis: RNA-extraction, matrix effects, uncertainty of measurement and
 precision. *J. Virol. Methods*, **160**, 157–162.
- 189 GREINER M., SOHR D. & GOEBEL P.A. (1995). Modified ROC analysis for the selection of cut-off values and the definition of 190 intermediate results of serodiagnostic tests, *J. Immunol. Methods*, **185**, 123–132.
- ISO/IEC (1995). Guide to the expression of uncertainty in measurement (GUM), ISO/IEC Guide 98:1995. International
 Organization for Standardization (ISO), www.iso.org.
- 193 ISO/IEC (2005). ISO/IEC 17025:20052017. General Requirements for the Competence of Testing and Calibration
 194 Laboratories. International Organization for Standardization (ISO), <u>www.iso.org</u>.
- ISO/IEC GUIDE 98-3:2008 (E). Uncertainty of measurement Part 3 Guide to the expression of uncertainty in measurement
 (GUM:1995), International Organization for Standardization (ISO), <u>www.iso.org</u> (accessed 10 June 2023).
- 197 <u>SCC- STANDARDS COUNCIL OF CANADA. Requirements and Guidance for method Verification and Validation in Testing</u>
 198 <u>Laboratories (RG-MVVT), 2021-05-24. http://www.scc.ca/ (accessed 10 June 2023).</u>
- TOUSSAINT J.F., ASSAM P., CAIJ B., DEKEYSER F., KNAPEN K., IMBERECHTS H., GORIS N., MOLENBERGHS G., MINTIENS K., & DE CLERCQ K. (2007). Uncertainty of measurement for competitive and indirect ELISAs. *Rev. sci. tech. Off. int. Epiz.*, **26**, 649– 656.
- 202 NEWBERRY K. & COLLING A (2021). Quality standards and guidelines for test validation for infectious diseases in veterinary 203 laboratories. Rev. Sci. Tech. Off. Int. Epiz., 40, 227-237. Available at 204 https://doc.woah.org/dyn/portal/index.xhtml?page=alo&aloId=41245 (accessed 22 June 2023)
- 205
 206
 **
 207
 208
 209
 209
 209
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200
 200

1	Annex 7. Item 5.1. – Chapter 2.2.6. Selection and use of reference samples and panels						
2	MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION						
3	Paris, 4–8 September 2023						
4							
F	 CHAPTER 2.2.6.						
5			UNAFIER 2.2.0.				
6	_		ECTION AND US	-	-		
7	REFERE	N (CE SAMPLES A	NI	D PANELS		
8			INTRODUCTION				
9 10 11 12 13	The WOAH Validation Recommendations provide detailed information and examples in support of the WOAH Validation Standard that is published as Chapter 1.1.6 Principles and methods of Validation of diagnostic assays for infectious diseases of terrestrial animals this Terrestrial Manual, or Chapter 1.1.2 of the Aquatic Manual. The Term "WOAH Validation Standard" in this chapter should be taken as referring to those chapters.						
14 15 16 17 18 19	through to the maintenance and monitoring <u>of</u> assay performance in the diagnostic laboratory and all of the stages in between. The critical importance of reference samples and panels cannot be over-emphasised. The wrong choice of reference materials can lead to bias and flawed conclusions right from development through to validation and use. Therefore, care must be exercised in selecting reference samples and						
20 21 22 23	topics and alphanumeric subheadings (e.g. Proof of concept, A.2.1) refer to the relevant section in the WOAH Validation Standard, <u>Chapter 1.1.6 Validation of diagnostic assays for infectious diseases of</u>						
	Group A		Group B		Group D		
	Proof of concept, A.2.1.		Asp, B.1.2.		Standard method comparison, B.2.6.		
	Operating range, A.2. 2 <u>3.</u>		Analytical accuracy, <u>ancillary</u> <u>tests</u> B.1.4.		Provisional recognition, B.2.6-7		
	<u>ASe, B.1.3.</u>		Reference samples and panels		Biological modifications, B.5.2.2.		
	Optimisation, A.2. 3 2.		Group C		Group E		
	Robustness, A.2.5. Preliminary repeatability, A.2.8.		Repeatability B.1.1.		DSp and DSe Gold standard, B.2.1.		
	Calibration <u>and process control</u> , A.2.6.		Preliminary reproducibility, B.2.6 $\underline{\underline{7}}$.		Group F		
	Process control, A.2.6.		Reproducibility, B.3.		DSp and DSe no gold standard B.2.2.		
	ASe, B.1.3.		Proficiency testing, B.5.1.				

ASp = Analytical specificity; ASe = Analytical sensitivity; DSp = diagnostic specificity; DSe = diagnostic sensitivity

Technical modifications, B.5.2.1. Reagent replacement, B.5.2.3.

25 As can be seen in Figure 1, Reference samples and/or panels are mentioned throughout the WOAH 26 Validation Standard, chapter 1.1.6. As defined in the glossary of the OIE Quality Standard and Guidelines 27 for Veterinary Laboratories: Infectious Diseases, 'Reference materials are "substances whose properties are 28 sufficiently homogenous and well established to be used for the calibration of an apparatus, the assessment 29 of a measurement method, or for assigning values to materials<u>"³³.</u> In the context of test method validation, 30 reference materials or samples contain the analyte of interest in varying concentrations or activities 31 reactivities and are used in developing and evaluating the candidate assay's analytical and diagnostic 32 performance-characteristics. In our case, Analyte means the specific component of a test sample that is 33 detected or measured by the test method, e.g. antibody, antigen or nucleic acid. These Reference samples 34 may be sera, fluids, tissues, excreta, feed and or environmental samples that contain the analyte of interest 35 and are usually harvested from infected animals and their environments. However, in some cases, they may 36 be prepared in the laboratory from an original starting material (e.g. a dilution of a high positive serum in 37 negative serum) or perhaps created by spiking the chosen matrix with a derived analyte (e.g. a bacterial or 38 viral culture, a recombinant/expressed protein, or a genomic construct). Whether natural or prepared, they 39 are used in experiments throughout the development process, carry over into the validation pathway and 40 can be used to monitor performance throughout the lifespan of the assay.

- In Figure 1, reference samples and panels are grouped based on similar characteristics and composition
 and these groupings will be the basis for the following descriptions. As a cross-reference, the appropriate
 Section of the OIE Validation Standard is indicated under each particular application of the reference sample
 or panel.
- 45 Reference samples may be used for multiple purposes from the initial stages of development and 46 optimisation, through Stage 1 and into continual monitoring and maintenance of the assay. Wherever possible, large quantities of these-reference samples should be collected or prepared and preserved for 47 48 long-term use. Switching reference samples during the validation process introduces an intractable variable that can severely undermine interpretation of experimental data and therefore, the integrity of the 49 50 development and validation process. For assays that may target multiple species, the samples should be 51 representative of the primary species of interest. It is critical that these samples reflect both the target analyte 52 and the matrix in which it is found in the population for which the assay is intended. The reference materials should appropriately represent the range of analyte concentration to be detected by the assay. 53
- 54 It is important to emphasise that, no matter Whether reference samples are selected from natural sources 55 or prepared in the laboratory, all selection criteria <u>or and</u> preparation procedures, as well as testing 56 requirements, need to be fully described and put into document control. Not only is this good quality 57 management practice, but it will provide both an enhanced level of continuity and confidence throughout the 58 lifespan of the assay. <u>Summaries of the data to be collected and documented for reference material can be</u> 59 found in Figure 2. For more detail on best practice and quality standards for the documentation of 60 provenance of reference material refer to Watson et al. (2021).

³³ https://www.techlab.fr/Commun/UK Def MRC.asp

61 62

63 64

65

Figure. 2. Documentation of reference material should be thorough to ensure i) transparency of intended purpose during assay development; ii) the correct sample types are used in all stages of assay development and validation; iii) accurate replacement of depleted reagents; and iv) appropriate choice of reference material during assay modification and re-validation. Minimum descriptive metadata are listed for pathogen, animal host, tissue type and phase of infection.

Pathogen data	Animal host and sample type data	Phase of Infection data
 <u>Strain/isolate</u> <u>Serotype</u> <u>Genotype</u> <u>Lineage</u> <u>Tests used for</u> <u>characterisation</u> 	 <u>Natural infection</u> <u>Experimental infection and protocol used</u> <u>Species</u> <u>Breed</u> <u>Age</u> <u>Sex</u> <u>Reproductive status</u> <u>Vaccination history</u> <u>Herd history</u> 	 <u>Clinical signs</u> <u>Antibody profiles</u> <u>Pathogen loading and</u> <u>shedding</u> <u>Tests used to determine</u> <u>status of disease/infection</u> (case definition)
	 <u>Tissue type/s (matrix) used</u> <u>For spiked samples – detail source of analyte</u> <u>and diluent (matrix) used</u> <u>Details relating to pooling of samples</u> 	

66

A. GROUP A

67 The question of pooling of samples to create a reference sample is often asked. If reference material is harvested from a 68 single animal, it is important to ascertain whether or not it is representative of a typical course and stage of infection within the context of the population to be tested. If not, this could lead to bias and flawed conclusions related to validation. Pooling 69 is a good alternative but it is imperative to pool from animals that are in a similar phase of infection. This is particularly 70 important for antibody detection systems. Pooling also addresses the issue of the larger quantities of reference material to 71 72 be stored for long term use, especially when dealing with smaller host species. Before pooling any samples, it is preferable that they be independently tested to demonstrate that they are similar with respect to analyte concentration and/or 73 reactivity. There should be an assessment following pooling to ensure that unforeseen interference is not introduced by 74 the pooling of multiple samples, for example differing blood types or antibody composition within the independent samples 75 76 could cross-react within the pool, thus causing the pooled sample to behave differently in the specified assay than the individual samples when tested independently. 77

It is often difficult to obtain individual samples that truly represent analyte concentrations or reactivities across the spectrum 78 of the expected range. Given the dynamics of many infections or responses to pathogens, intermediate ranges are often 79 very transient. In the case of antibody responses, early infection phases in individual animals often result in highly variable 80 and heterogeneous populations of antibody isotypes and avidities. In general, these do not make good reference samples 81 for assessing the analytical characteristics of an assay. They are nonetheless important for different types of reference 82 panels as will be discussed later. For most applications in Group A, it is acceptable to use prepared samples that are 83 spiked with known concentrations of analyte or a dilution series of a high positive in negative matrix to create a range of 84 85 concentrations.

86 Whether natural or prepared, reference samples should represent the anticipated range of analyte concentrations, from 87 low-weak to high strong positive, which would be expected during a typical course of infection. A negative reference sample 88 should be included as a background monitor. If a negative (matrix) is used as diluent for preparation of a positive reference 89 sample (e.g. a negative serum used to dilute a high positive serum or tissue spiked with a construct), that negative should 90 definitely be included as the negative reference sample.

91 As mentioned above, all reference samples should be well characterised. This includes documentation on both the 92 pathogen and donor host. For pathogens, this may include details related to strain, serotype, genotype, lineage, etc. The source of the host material should be well described with respect to species, breed, age, sex, reproductive status, 93 vaccination history, herd history, etc. Wherever possible, the phase of infection should be noted. This could include details 94 related to clinical signs, antibody profiles, pathogen load or shedding, etc. Equally important, tests that are used to 95 determine disease/infection status need to be well documented (see Section E of this chapter for further explanation). In 96 some cases, experimental infection/exposure may be the only viable option for the production of reference material. In this 97 98 case, all of the above considerations plus the experimental protocol should be detailed.

Above all else, natural or prepared, reference materials must be unequivocal with respect to their status as representing either a true positive or a true negative sample. This may require that the status be confirmed using another test or battery of tests. For example, many antibody reference sera are characterised using multiple serological tests. This provides not only confidence but additional documented characteristics that may be required when attempting to replace or duplicate

103 this reference material in the future.

Recommendations regarding stability and storage of reference materials are available: <u>https://www.woah.org/en/what-we-</u>
 <u>offer/veterinary-products/#ui-id-4</u>

106 **1.** Proof of concept (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section A.2.1)

The WOAH Validation Standard, Chapter 1.1.6 states that test methods and related procedures must be appropriate for 107 108 specific diagnostic applications in order for the test results to be of relevance. In other words, the assay must be 'fit for purpose'. Many assays are developed with good intentions but without a specific application in mind. At the very outset, it 109 is critical that the diagnostic purpose(s) should be defined with respect to the population(s) to be tested. The most common 110 111 purposes are listed in broad terms in Section A of the WOAH Validation Standard, chapter 1.1.6. As such, they are inclusive of more narrow and specific applications. However, these specific purpose(s) need to be clearly defined from the outset 112 and are critically important in the context of a fully validated assay. As will be seen in the following descriptions, clearly 113 defining the application will have impact on both the selection of reference samples and panels and the design of analytical 114 and diagnostic evaluations. 115

116 **2.** Operating range (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section A.2.<u>2</u><u>3</u>) and 117 analytical sensitivity (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.1.3)

118 2.1. Analytical approaches Operating range and analytical sensitivity

The operating range of the assay is defines the lower and upper analyte detection limits and the interval of analyte 119 120 concentrations (amounts) over which the method provides suitable accuracy and precision. It also defines the lower and upper detection limits the assay. To establish this range. The operating range is established by serial dilution, to 121 122 extinction, of replicates of a high strong positive reference sample is selected. This high positive sample, either natural 123 or prepared, is serially diluted to extinction. Dilutions of the strong positive are made in a-negative matrix representative of the typical sample matrix of samples type taken from animals in the population targeted by the 124 assay. This includes antibody assays where a high replicates of a strong positive reference serum should be diluted 125 126 in a negative reference serum to create the dilution series. Analytical sensitivity (ASe) is measured by replicates of 127 the lower limit of detection (LOD) of an analyte in an assay. The same high strong positive reference sample may be 128 used to determine both the operating range and the analytical LOD.

129 2.2. Comparative approaches to analytical sensitivity

130 If the intended purpose is to detect low levels of analyte or subclinical infections, it may be difficult to obtain the 131 appropriate reference materials from early stages of the infection process. In some cases, it may be useful to 132 determine a comparative ASe by running a panel of samples on the candidate assay and on another independent 133 assay. Ideally this panel of samples would be serially collected from either naturally or experimentally infected animals 134 and should represent infected animals early after infection, on-through to the development of clinical or fulminating 135 disease, if possible. This would provide a relative comparison of ASe between the assays, as well as, and a temporal 136 comparison of the earliest point of detection relative to the pathogenesis of the disease.

An experiment like the one described above, provides a unique opportunity to collect reference samples representing 137 138 a natural range of concentrations that would be useful for other validation purposes. Care must be taken to avoid use of such samples when inappropriate (consult Group D below). Wherever possible serial samples should be collected 139 from at least five a statistically sound number of animals throughout the course of infection. In cases where sampling 140 141 is lethal (e.g. requiring the harvest of internal organ tissues), the number of animals required would be a minimum depends on need and fitness of five per sampling event the experimental approach. In all cases approval from an 142 ethics committee is required. For smaller host species, this the number may need to be increased in order to collect 143 sufficient reference material. Given that experiments like this require a high commitment of resources, it would be 144 wise to maximise the collection of not only the currently targeted reference samples but additional materials (e.g. 145 multiple tissues, fluids, etc.) that may be useful as reference materials in the future. 146

Optimisation (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section A.2.3<u>2</u>) and preliminary repeatability (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section A.2.6<u>8</u>)

149 Optimisation is the process by which the most important physical, chemical and biological parameters of an assay are evaluated and adjusted to ensure that the performance characteristics of the assay are best suited to the intended 150 application. At least three reference samples representing negative, low weak and high strong positive may be chosen 151 from either natural or prepared reference samples. Optimisation experiments are rather exhaustive especially when assays 152 153 with multiple preparatory and testing steps are involved. It is very important that a sufficient quantity of each reference 154 sample be available to complete all optimisation experiments. Changing reference samples during the course of optimisation is not recommended as this will result in the addition of an uncontrolled variable and a disruption in the 155 continuity of optimisation evidence. 156

Assessment of repeatability should begin during assay development and optimisation stages. Repeatability and is further
 verified during Stage 1 of assay validation (Section B.1.1 <u>of chapter 1.1.6</u>). The same reference samples should be used
 for both processes, again throughout to provide continuity of evidence.

4. Calibration and process controls (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section A.2.6)

162 **4.1.** International, national or in-house analyte reference standards

163 International reference standards are highly characterised, contain defined concentrations of analyte, and are usually prepared and held by international reference laboratories. They are the reagents to which all assays and/or other 164 reference materials should be standardised. National reference standards are calibrated by comparison with an 165 international standard reagent whenever possible. In the absence of an international standard, a national reference 166 167 standard may be selected or prepared and it then becomes the standard of comparison for the candidate assay. In the absence of both of the above, an in-house standard should be selected or prepared by the development laboratory 168 within the responsible organisation. In all cases, thorough documentation of reference material should be observed 169 as summarised in Figure 2. All of the standard reagents, whether natural or prepared, must be highly characterised 170 through extensive analysis, and preferably the methods for their characterisation, preparation, and storage have been 171 published in peer-reviewed publications (Watson et al., 2021). These reference standards should also be both stable 172 173 and innocuous.

Reference standards, especially antibody, are usually provided in one of two formats. They may be provided as a 174 175 single positive reagent of given titre with the expectation that the candidate assay will be standardised to give an equivalent titre. This is a straight forward analytical approach but many of these 'single' standards have been prepared 176 from highly positive samples as a pre-dilution in a negative matrix in order to maximise the number of aliquots 177 available. The drawback here is that there is no accounting for any potential matrix effect in the candidate assay as 178 there is no matrix control provided. The other approach is to provide a negative and a low-weak and high-strong 179 positive set of reference standards that are of known concentrations or reactivities and are within the operating range 180 181 of the standard method that was used to prepare them. The negative provided in the set must be the same as the negative diluent used to prepare the weak and strong positive reference standard, if the positive standards were 182 diluted. This compensates for any potentially hidden matrix effect. In addition, this set of three acts as a template for 183 the selection and/or preparation of process controls (discussed below). 184

185 Classically, the above standards usually have been polyclonal antibody standards and to a lesser extent, conventional 186 antigen standards used for calibration of serological assays. However, today, reference standards could also be 187 monoclonal antibodies or recombinant/expressed proteins or genomic constructs, if they are to be used to calibrate 188 assays to a single performance standard.

1894.2.Working standards or process controls

Working standard reagent(s), commonly known as quality or process controls, are calibrated to international, national, 190 or in-house standard reagents. They are selected or prepared in the local matrix which is found in the population for 191 which the assay is intended. Ideally, negative and low weak and high strong positive working standards should be 192 selected or prepared. Concentrations and/or reactivities should be within the normal operating range of the assay. 193 Large quantities should be prepared, aliquoted and stored for routine use in each diagnostic run of the assay. The 194 195 intent is that these controls should mimic, as closely as possible, field samples and should be handled and tested like routine samples. They are used to establish upper and lower control limits of assay performance and to monitor 196 random and/or systematic variability using various control charting methods. Their daily performance will determine 197 whether or not an assay is in control and if individual runs may be accepted. As such, these working reference 198 samples are critically important from a quality management standpoint. 199

5. Technical modifications (WOAH Validation Standard, Chapter 1.1.6, Section B.5.2.1)

Technical modifications to a validated assay such as changes in instrumentation, extraction protocols, and conversion of an assay to a semi-automated or fully automated system using robotics will typically not necessitate full revalidation of the assay. Rather, a methods comparison study may be done to determine if these minor modifications to the assay protocol will affect the test results. Consult <u>See</u> chapter 2.2.8 *Comparability* of assays after changes in a validated test method for description of experiments and statistical approaches to assay precision in the face of technical modifications that are appropriate for comparability testing (Bowden & Wang, 2021; Reising *et al.*, 2021).

207 In general, these approaches require the use of three reference samples, a negative, a weak and a low and high strong positive. Again these samples to represent the entire operating range of both assays. Samples may be either natural or 208 prepared. The important point to re-iterate here is that the same reference samples that were used in the developmental 209 210 stages of the assay may be used to assess modifications after the method has been put into routine diagnostic use. This provides a higher level of confidence assessing potential impacts because the performance characteristics of these 211 212 reference samples have been well characterised. At the very least, if new reference samples are to be used, they should 213 be selected or prepared using the same criteria or preparation procedures established for previous materials. Again as this 214 enhances the continuity of evidence.

6. Reagent replacement (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.5.2.3)

When a reagent such as a process control sample is nearing depletion, it is essential to prepare and repeatedly test a replacement before such a control is depleted. The prospective replacement should be included in multiple runs of the assay in parallel with the original control to establish their proportional relationship. It is important to change only one control reagent at a time to avoid the compound problem of evaluating more than one variable.

Again, it cannot be over emphasised that any-Replacement reference reagent should be selected or prepared using the same criteria or preparation procedures established for previous materials. Again as this enhances the continuity of evidence and confidence in the assay and underlines the importance of documentation of reference material data (Figure 2).

B. GROUP B

1. Analytical specificity (WOAH Validation Standard, Chapter 1.1.6, Section B.1.2)

Analytical specificity (ASp) is the degree to which the assay distinguishes between the target analyte and other components
 that may be detected in the assay. This is a relatively broad definition that is often not well understood. ASp may be broken
 down into different elements as described below.

The choice of reference samples that are required to assess ASp is highly dependent on the <u>specific intended</u> purpose or application that was originally envisaged <u>defined</u> at the development stage of the assay. Assessment of ASp is a crucial element in proof of concept and verification of fitness for purpose <u>and may be broken down into three elements: selectivity</u>. <u>exclusivity and inclusivity</u>.

<u>Selectivity:</u> an important element is the extent to which a method can accurately detect and or quantify the targeted analyte
 in the milieu of nucleic acids, proteins and/or antibodies in the test matrix. This is sometimes termed 'selectivity'. An
 example is the use of reference samples for tests that are designed to differentiate infected from vaccinated animals (DIVA
 tests).

Reference samples need to be selected and tested from i) non-infected/non-vaccinated, ii) non-infected/vaccinated, iii) 236 infected/non-vaccinated, and iv) infected/vaccinated animals. These samples may be collected under field conditions but 237 238 it is important that an accurate history be collected, ideally with respect to the animals, but at least to the herds involved, including vaccination practices and disease occurrences (Figure 2). Alternatively, it may be necessary to produce this 239 material in experiments like those described in Section A.2.2 of this chapter, but including a combination of experimentally 240 vaccinated and challenged animals. It Application of the 3 Rs (replacement, reduction and refinement) aims to avoid or 241 242 minimise the number of animals used in experiments. For enzyme-linked immunosorbent assays (ELISAs), it is important 243 to avoid use of the vaccine as capture antigen in the assay (e.g. indirect ELISA enzyme linked immunosorbent assay [I-ELISA]), because carrier proteins in the vaccine may stimulate non-specific antibody responses in vaccinated animals that 244 245 may be detected in ELISA-leading to false positives in the assay. Similarly to the comparative approach described above 246 with respect to ASe, at least five animals in each group should be considered. For smaller host species, this number may need to be increased in order to collect sufficient reference material., leading to false positives in the assay. Depending on 247 the DIVA test, a single experiment could be designed to assess aspects of both ASe and ASp. 248

249 A second element, sometimes termed 'exclusivity', Exclusivity is the capacity of the assay to detect an analyte or genomic 250 sequence that is unique to a targeted organism, and excludes all other other known organisms that are potentially cross-251 reactive. This is especially true in serological assays where there are many examples of antigens expressed by other organisms that are capable of eliciting cross-reacting antibody. An attempt should be made to obtain reference samples 252 from documented cases of infections and/or organisms that may be cross-reactive. Depending on the type of assay, these 253 reference materials may represent the organism itself, host-derived samples, or genomic sequences. A profile for the 254 exclusivity of the assay should be established, and expanded on a continual basis as potentially cross-reactive organisms 255 256 arise.

257 Thirdly, a critical design consideration_Inclusivity relates to the capacity of an assay to detect one or several strains or 258 serovars of a species, several species of a genus, or a similar grouping of closely related organisms-viruses, bacteria or 259 antibodies. This defines the scope of detection and thus the fitness for purpose. Reference samples are required to define 260 the scope of the assay The scope of the assay defines the choice of reference samples and the results will determine proof 261 of fitness. If for example an assay is developed as a screening test to detect all known genotypes or serotypes of a virus, 262 then reference samples from each representative type should be tested. As new lineages or serotype variants arise, they 263 too should be tested as part of the test profile, which should be updated on an ongoing basis.

264 2. Analytical accuracy of adjunct ancillary tests (WOAH Validation Standard, Chapter 1.1.6, 265 Section B.1.4)

Some test methods or procedures are solely analytical tools and are usually applied used to further characterise an analyte 266 that has been detected in a primary assay, for example assays like. Examples are the virus neutralisation tests used to 267 type an isolated virus or characterise an antibody response and subtyping of haemagglutinin genes by polymerase chain 268 reaction of avian influenza virus. Such adjunct ancillary tests must be validated for analytical performance characteristics, 269 but and differ from to routine diagnostic tests because they do not require validation for diagnostic performance 270 characteristics. The analytical accuracy of these tests is often dependant on the use of reference reagents material. These 271 272 reagents, whether they are antibody for typing strains of organisms or reference strains of the organism, etc., should be 273 thoroughly documented, as required for any other reference material (Figure 2), with respect to their source, identity and 274 performance characteristics.

275

C. GROUP C

Reference samples in Group C may be used for a number of purposes. In the initial development stages, they may be 276 277 used in the assessment of assay repeatability and both preliminary reproducibility in Stage 1 and the more in depth assessment of reproducibility in Stage 3 of the Validation Pathway. However, these samples have a number of other 278 potential uses once the assay is transferred to the diagnostic laboratory. They may be used as panels for training and 279 280 qualifying of analysts, and for assessing laboratory proficiency in external ring testing programmes. Ideally, 20 or more individual samples should be prepared in large volumes. About a quarter (25%) should be negative samples and the 281 remainder (75%) should represent a collection of positives spanning the operating range of the assay. They should be 282 aliquoted into individual tubes in sufficient volumes for single use only and stored for long term use (Chapter 1.1.2 283 284 Collection, submission and storage of diagnostic specimens). The number of aliquots of each that will be required will depend on how many laboratories will be using the assay on a routine diagnostic basis and how often proficiency testing 285 is anticipated. Ideally, they should be prepared in an inexhaustible quantity, but this is seldom feasible. At a minimum, 286 287 several hundred or more aliquots of each should be prepared at a time if the assay is intended for use in multiple 288 laboratories. This allows assessment of laboratory proficiency by testing the same sample over many testing intervals - a useful means of detecting systematic error (bias) that may creep into long term use of an assay. 289

290 These samples may be natural or prepared from either single or pooled starting material. The intent is that they should 291 mimic as closely as possible a true test sample. Because mass storage is always a problem, it may be necessary to store these materials in bulk and prepare working aliquots from time to time. However, if storage space is available, it is 292 preferable to prepare and store large numbers of aliquots at one time because bulk quantities of analyte, undergoing 293 294 freeze-thaw cycles to prepare a few aliquots at a time, may be subject to degradation. Because this type of reference material is consumed at a fairly high rate, they will need to be replaced or replenished on a continual basis. As potential 295 296 replacement material is identified during routine testing or during outbreaks, it is advisable to work with field counterparts 297 to obtain bulk reference material and store it for future use. Alternatively, it may be necessary to produce this material in 298 experiments like those described in Section A.2.2 of this chapter. Similar to the comparative approach described above with respect to ASe, at least five animals in each group should be considered. For smaller host species, this number may 299 need to be increased in order to collect sufficient reference material. 300

Repeatability (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.1.1) and preliminary reproducibility provisional assay recognition (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.2.6)

Repeatability is the level of agreement between results of replicates of a sample both within and between runs of the same test method in a given laboratory. Repeatability is estimated by evaluating variation in results of replicates from a minimum of three (preferably five) samples representing analyte activity within the operating range of the assay. Consult Chapter 2.2.4 *Measurement uncertainty* for statistical approaches for measures of uncertainty for assessments of repeatability.

308 Reproducibility is the ability of a test method to provide consistent results, as determined by estimates of precision, when applied to aliquots of the same samples tested in different laboratories. However, preliminary reproducibility estimates of 309 310 the candidate assay should be determined during developmental stages. A small panel of three (but preferably five) representing negative, weak and both low and high strong positives, like those described above, would be adequate. This 311 type of panel could also be used for a limited evaluation of reproducibility to enhance provisional acceptance status for the 312 assay. The test method is usually assessed in one two or more laboratories with a high level of experience and proficiency 313 314 in assays similar to the candidate assay. The panel of 'blind' samples is evaluated using the candidate assay in each of these laboratories, using the same protocol, same reagents and comparable equipment. This is a scaled-down version of 315 Stage 3 of assay validation. Consult Chapter 2.2.4 for further explanation of the topic and its application. 316

317 2. Reproducibility (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.3)

318 Reproducibility is an important measure of the precision of an assay when used in a cross-section of laboratories located 319 in distinct or different regions or countries using the identical assay (protocol, reagents and controls). As the number of laboratories increases, so does the number of variables encountered with respect to laboratory environments, equipment 320 differences and technical expertise. These An overview of the factors affecting testing reproducibility is provided in Waugh 321 & Clark (2021). Reproducibility studies are a measure of an assay's capacity to remain unaffected by substantial changes 322 323 or substitutions in test conditions anticipated in multi-laboratory use (e.g. shipping conditions, technology transfer, reagents batches, equipment, testing platforms and/or environments). Each of At least three laboratories should test the same panel 324 of 'blind' samples containing a minimum of 20 samples, representing negative and a range of positive samples. If selected 325 326 negative and/or positive samples in the panel are duplicated, in the panel then it may be possible to assess both assay 327 reproducibility and within-laboratory repeatability estimates may be augmented by replicate testing of these samples when used in the reproducibility studies. 328

329 3. Proficiency testing (WOAH Validation Standard, Chapter 1.1.6, Section B.5.1)

330 A validated assay in routine use in multiple laboratories needs to be continually monitored to ensure uniform performance and provide overall confidence in test results. This is assessed through external quality assurance programmes. Proficiency 331 testing is one measure of laboratory competence derived by means of an inter-laboratory comparison; implied is that 332 participating laboratories are using the same (or similar) test methods, reagents and controls. Results are usually 333 expressed qualitatively, i.e. either negative or positive, to determine pass/fail criteria. However, for single dilution assays, 334 where semi-quantitative results provide are provided, additional data for assessment of analysis may assess non-random 335 336 error among the participating laboratories. Refer to Johnson & Cabuang (2021) for an overview of proficiency testing and 337 ring trials.

338 Proficiency testing programmes are varied depending on the type of assay in use. For single dilution type assays, panel sizes also vary but a minimum of five samples, representing negative and both low and high positives, like those described 339 above, would be adequate. Proficiency testing is not unlike a continuous form of reproducibility assessment. However, 340 341 reproducibility, by definition, is a measure of the assay's performance in multiple laboratories; whereas proficiency testing is an assessment of laboratory competence in the performance of an established and validated assay. Measurements of 342 precision can be estimated for both the reproducibility and repeatability data if replicates of the same reference sample are 343 included in this 'blind' panel. Consult Chapter 2.2.4 for further explanation of the topic and its application. vary but a 344 minimum of five samples, representing negative weak and strong positives, would be adequate. 345

346

D. GROUP D

Reference samples in Group D differ from the previous Groups in that each sample in the panel should be from a different individual animal. As indicated in Chapter 2.2.8 <u>Comparability of assays after changes in a validated test method</u>, experimental challenge studies often include repeated sampling of individual animals to determine the progression of disease, but this is a different objective than to comparing performance characteristics that would be associated with diagnostic sensitivity (DSe) and diagnostic specificity (DSp) of a test method. Serially drawn samples, taken on different days from the same animal, cannot be used as representative of individual animals in populations targeted by the assay,
 because such samples violate the rule of independence of samples required for such studies.

Care must be taken in choosing the reference samples and the standard (independent) method used in this type of comparison to ensure that the analytes being detected (if different) demonstrate the same type of pathogenic profile in terms of time of appearance after exposure to the infectious agent, and relative abundance in the test samples chosen.

Standard method comparison and provisional recognition (WOAH Validation Standard, Chapter 1.1.6, Sections B.2.6-5 and B.2.6)

359 There are situations where it is not possible or desirable to fulfil Stage 2 of the Validation Pathway because appropriate samples from the target population are scarce and animals are difficult to access (such as for exotic diseases). However. 360 361 a small but select panel of highly characterised test samples representing the range of analyte concentration should be run in parallel in the candidate assay method and by a WOAH standard method, as published in the WOAH Manuals. 362 363 Biobanks may be a useful resource in this context, providing well-characterised samples supported with metadata to 364 enhance transparency and provenance of samples used in method comparisons (Watson et al., 2021). If the methods are 365 deemed to be comparable (Chapter 2.2.8), and depending on the intended application of the assay, the choice may be 366 made that further diagnostic validation is not required. For example, if the intended application is for screening of imported animals or animal products for exotic pathogens or confirmation of clinical signs, full validation beyond a test method 367 368 comparison may not be feasible or warranted.

Experience has shown that the greatest obstacle to continuing through Stage 2 of the Validation Pathway is the number of defined samples required to estimate diagnostic performance parameters with a high degree of certainty (WOAH Validation Standard, <u>chapter 1.1.6</u>, Section B.2). In some cases, provisional recognition by international, national or local authorities may be granted for an assay that has not been completely evaluated past analytical stages. The different rationales for provisional acceptance are well explained in the WOAH Validation Standard, <u>chapter 1.1.6</u>. In all cases however, sound evidence must exist for comparative estimates of DSp and DSe based on a small select panel of well-characterised samples containing the targeted analyte.

376 Ideally, for both comparison with a standard method or provisional recognition, a panel of, for example, 60 samples could 377 be assembled to ensure sufficient sample size for statistical analysis of the resulting data. This would include 30 'true' 378 negatives and 30 'true' positives. Wherever possible, the positives should reflect the range of analyte concentrations or 379 activities expected in the target population. As mentioned above, each sample in this panel must represent an individual animal. Consult Chapter 2.2.5 for statistical approaches to determining methods comparability using diagnostic samples.

381 2. Biological modifications (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.5.2.2)

382 There may be situations where changes to some of the biologicals used in the assay may be necessary and/or warranted. 383 This may include changes to reagents themselves or a change to a different type of specimen which contains the same analyte as targeted in the original validated assay (e.g. from serum to saliva). At the very least, all of the analytical criteria 384 of the validation pathway must be re-assessed before proceeding. If the analytical requisites are met, the remaining 385 386 question relates to whether or not a full diagnostic validation is required. A similar approach to the above using a panel of 387 60 individual reference samples may be considered. However, in this case the original test method would be considered 388 as the standard (independent) test and the modified method would be considered the candidate. Consult Chapter 2.2.5 for statistical approaches to determining methods comparability using diagnostic samples. 389

390

E. GROUP E

Reference animals and reference samples in this Group E are well described in the WOAH Validation Standard, <u>chapter</u>
 <u>1.1.6</u>, Section B.2.1). However, there are a few points that are worth re-iterating here.

Gold standard'³⁴ – diagnostic specificity and diagnostic sensitivity (WOAH Validation Standard, Chapter 1.1.6, Section B.2.1)

For conventional estimates of DSp, negative reference samples refer to true negative samples, from animals that have had no possible infection or exposure to the agent. In some situations, where the disease has never been reported in a country or limited to certain regions of a country, identification of true negative reference samples is usually not a problem. However, where the disease is endemic, samples such as these may be difficult to locate. It is often possible to obtain

³⁴ The term "Gold Standard" is limited to a perfect reference standard as described in the WOAH Validation Standard, chapter 1.1.6, Section B.2.1.2, and Chapter 2.2.5 Statistical approaches to validation, Introduction and Figure 1.

these samples from regions within a large country or perhaps different countries where the disease in question <u>does not</u> <u>occur or</u> has either been eradicated or has never had the disease in question.

Again-For conventional estimates of DSe, positive reference samples refer to true positives. Care must be taken to ensure that the sample population is representative of the population that will be the target of the validated assay. It is generally problematic to find sufficient numbers of true positive reference animals, as determined by isolation of the organism. It may be necessary to resort to samples from animals that have been tested by a combination of methods that unequivocally classify animals as infected/exposed as discussed in the WOAH Validation Standard, chapter 1.1.6.

406 The important point here is that All samples, irrespective of origin, must be documented as they would for any other 407 reference sample so as to unequivocally to classify animals as infected or exposed, dependent on the fitness for purpose 408 and proposed use of the test. As mentioned in Section A, and summarised in Figure 2, of this chapter, all reference samples should be well characterised. This includes documentation on both the pathogen and donor host. For pathogens, this may 409 include details related and data documented to strain, serotype, genotype, lineage, etc. The source of the host material 410 should be well described with respect to species, breed, age, sex, reproductive status, vaccination history, herd history, 411 etc. Wherever possible, the phase of infection should be noted. This could include details related clinical signs, antibody 412 413 profiles, pathogen load or shedding, etc. In some cases, experimental infection/exposure may be the only viable option ensure appropriate sample selection for the production of reference material (see the OIE Validation Standard, Section 414 B.2.3). In this case, all of the above and the experimental protocol should be detailed intended purpose. 415

Particularly relevant to these reference samples, the tests that are used to determine their so called 'true' disease/infection status need to be well documented in order to assess potential errors in estimates that may be carried over into the estimates for the candidate assay. Indeed, when using imperfect standard assays to define reference animal or sample status, the DSe and DSp performance estimates of the candidate assay may be flawed and often overestimated. Consult Chapter 2.2.5 for statistical considerations. <u>Situations where a perfect reference is available for either positive or negative</u> animals, and one where the reference is perfect for both are described for diagnostic test validation by Heuer & Stevenson (2021).

423

F. GROUP F

Animals of unknown status – diagnostic specificity and diagnostic sensitivity (WOAH Validation Standard, <u>Chapter 1.1.6</u>, Section B.2.2)

Latent-class models are introduced in the WOAH Validation Standard, <u>chapter 1.1.6</u>. They do not rely on the assumption of a perfect reference (standard or independent) test but rather estimate the accuracy of the candidate test and the reference standard with the combined test results. Because these statistical models are complex and require critical assumptions, statistical assistance should be sought to help guide the analysis and describe the sampling from the target population(s), the characteristics of other tests included in the analysis, the appropriate choice of model and the estimation methods based on peer-reviewed literature. Consult Chapter 2.2.5 for statistical considerations.

Reference populations, not individual reference samples, used in latent-class studies need to be well described. This
includes documentation on both the pathogen and donor host. For pathogens, this may include details related to strain,
serotype, genotype, lineage, etc., that may be circulating in the population. The source of the host material should be well
described with respect to species, breed, age, sex, reproductive status, vaccination history, herd history, etc.
summarised in Figure 2. Wherever possible, the phase of infection in the populations should be noted with respect to

As a special note, if latent class models are to be used to ascribe estimates of DSe and DSp and include multiple laboratories in the design, it is possible to incorporate an assessment of reproducibility into the assessment. As stated above, statistical advice should be sought in this respect. Bayesian latent class models are complex and require adherence to critical assumptions. Statistical assistance should be sought to help guide the analysis and describe the sampling from the target population(s), the characteristics of other tests included in the analysis, the appropriate choice of model and the estimation methods (based on peer-reviewed literature). See chapter 2.2.5 for details and Cheung *et al.*, 2021.

444	FURTHER READING
445	BOWDEN T.R., CROWTHER J.R. & WANG J. (2021). Review of critical factors affecting analytical characteristics of serological
446	and molecular assays. Rev. Sci. Tech. Off. Int. Epiz., 40, 53–73. doi:10.20506/rst.40.1.3208.
447	CHEUNG A., DUFOUR S., JONES G., KOSTOULAS P., STEVENSON M.A., SINGANALLUR N.B. & FIRESTONE S.M. (2021). Bayesian
448	latent class analysis when the reference test is imperfect. <i>Rev. Sci. Tech. Off. Int. Epiz.</i> , 40 , 271–286.
449	doi:10.20506/rst.40.1.3224
450	<u>JOHNSON P. & CABUANG L. (2021). Proficiency testing and ring trials. <i>Rev. Sci. Tech. Off. Int. Epiz.</i>, 40, 189–203.</u>
451	https://doi.org/10.20506/rst.40.1.3217
452	HEUER C. & STEVENSON M.A. (2021). Diagnostic test validation studies when there is a perfect reference standard. <i>Rev. Sci.</i>
453	Tech. Off. Int. Epiz., 40, 261–270. doi:10.20506/rst.40.1.3223
454	REISING M.M., TONG C., HARRIS B., TOOHEY-KURTH K.L., CROSSLEY B., MULROONEY D., TALLMADGE R.L., SCHUMANN K.R., LOCK,
455	A.B. & LOIACONO C.M. (2021). A review of guidelines for evaluating a minor modification to a validated assay. <i>Rev. Sci.</i>
456	<i>Tech. Off. Int. Epiz.</i> , 40 , 217–226. doi:10.20506/rst.40.1.3219
457 458 459	WATSON J.W., CLARK G.A. & WILLIAMS D.T. (2021). The value of virtual biobanks for transparency purposes with respect to reagents and samples used during test development and validation. <i>Rev. Sci. Tech. Off. Int. Epiz.</i> , 40 , 253–259. doi:10.20506/rst.40.1.3222.
460	WAUGH C. & CLARK G. (2021). Factors affecting test reproducibility among laboratories. Rev. Sci. Tech. Off. Int. Epiz., 40,
461	131–143. doi:10.20506/rst.40.1.3213
462 463	* *
464	NB: There is a WOAH Collaborating Centre for
465	Diagnostic Test Validation Science in the Asia-Pacific Region (please consult the WOAH Web site:
466	https://www.woah.org/en/what-we-offer/expertise-network/collaborating-centres/#ui-id-3).
467	Please contact the WOAH Collaborating Centre for any further information on validation.
468 469	NB: FIRST ADOPTED IN 2014.

Annex 8. Item 3.1.1. – Chapter 3.1.5. Crimean–Congo haemorrhagic fever

MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023

CHAPTER 3.1.5.

CRIMEAN-CONGO HAEMORRHAGIC FEVER

B. DIAGNOSTIC TECHNIQUES

. . .

Table 1. Diagnostic test formats for Crimean–Congo haemorrhagic fever virus infections in animals

	Purpose						
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases in animals	Prevalence of infection – surveillance	Immune status in individual animals or populations post- vaccination	
Detection and identification of the agent ^(a)							
Real-time RT-PCR	-	+++	-	+++(<u>b)</u>	+(c)	-	
Virus isolation in cell culture			-				
	Detection of immune response						
IgG ELISA	+++	+	_	+ <u>+(d)</u>	+++	-	
Competitive ELISA	+++	+	_	+ <u>+(d)</u>	+++	-	
IgM ELISA	-	++	-	++ <u>(e)</u>	_	-	

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; – = not appropriate for this purpose.

RT-PCR = reverse-transcription polymerase chain reaction; ELISA = enzyme-linked immunosorbent assay.

^(a)A combination of agent identification methods applied on the same clinical sample is recommended.

^(b)Molecular testing/isolation can be used to confirm acute infection in rare cases in animals showing

clinical signs as viraemia tends to be transient.

^(c)RT-PCR is used for the screening of tick populations in the context of surveillance studies.

^(d)Serological evidence of active infection with CCHFV has been demonstrated by seroconversion based on a rise in total or IgG antibody titres on samples taken at 2-4 weeks apart.

(e)Serological evidence of active infection with CCHEV has been demonstrated by the detection of IgM antibodies specific to CCHEV using two different ELISAs based on two different antigens.

. . .

1	Annex 9. Item 5.1. – Chapter 3.3.6. Avian tuberculosis
2	MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION
3	Paris, 4–8 September 2023
4	
5	CHAPTER 3.3.6.
6	AVIAN TUBERCULOSIS

- SUMMARY
- <u>Description of the disease:</u> Avian tuberculosis, or avian mycobacteriosis, is <u>an important a significant</u> disease that affects companion, captive exotic, wild, and domestic birds and mammals. The disease is most often caused by Mycobacterium avium subsp. avium (M. a. avium), <u>a member of the M. avium complex</u>. However, more than ten other mycobacterial species have been reported to infect birds. The most significant cause of poultry disease is M. a. avium.

7

8

9

10

11 12

- 13 Clinical signs of the disease vary depending on the organs involved. The classical presentation is 14 characterised by chronic and progressive wasting and weakness. Diarrhoea is common and joint swelling 15 are standard features in infected flocks. Some birds may show respiratory signs, and occasionally, sudden 16 death occurs. Some birds may develop granulomatous ocular lesions.
- Mycobacterium tuberculosis, the agent that most commonly causes human tuberculosis (gene IS61101) is
 *less commonly <u>rarely</u> the cause of infection in birds, <u>and it is</u> often as a the result of transmission from pet
 bird owners.*
- Members of M. avium complex: M. <u>a. avium (serotypes 1–3; containing gene segments IS901 and IS1245).</u>
 <u>M.</u> avium subsp. hominissuis (serotypes 4–6, 8–11, and 21; lacking gene segment IS901 and containing segment IS1245) and M. intracellulare (serotypes 7, 12–20, and 22–28; lacking both IS901 and IS1245) can also infect an extensive range of mammals such as swine, cattle, deer, sheep, goats, horses, cats, dogs, and exotic species. In humans, all members of <u>the</u> M. avium complex and M. genavense are capable of inducing can induce a progressive disease that is refractory to treatment, mostly mainly in immunocompromised patients.
- All manipulations involving <u>Due to</u> the <u>contagious nature of this group of organisms</u>, handling of open live
 cultures or of-material from infected birds must <u>only</u> be carried out <u>with <u>after an</u> appropriate biorisk</u>
 management <u>risk assessment and the implementation of biosafety measures designed to avoid infection</u>.
- Diagnosis of avian tuberculosis in birds depends on the demonstration of the above-mentioned <u>a</u> mycobacterial species in live or dead birds or the detection of an immune response, cellular or humoral, culture examination, or gene segments-IS6110, IS901 and IS1245 by polymerase chain reaction (PCR) in the excretions or secretions of live birds.
- Detection of the agent: Where clinical signs of avian tuberculosis are seen in the flock, or typical 34 tuberculous lesions are present in birds at necropsy, the demonstration of acid-fast bacilli in smears or 35 36 sections made from affected organs is sufficient for a quick positive diagnosis. If acid-fast bacilli are not found but typical tuberculous signs or lesions are present in the birds, <u>a</u> culture of the organism <u>or PCR</u> must 37 38 be attempted. PCR could also be carried out directly on tissue samples. Any acid-fast organism isolated should be identified by nucleic-acid-based tests or chromatographical (e.g. high-performance liquid 39 chromatography [HPLC]) criteria; serotyping of isolates of M. avium complex members or PCR for 16S rRNA 40 gene followed by sequencing, or the presence of an amplicon for the insertion sequences IS6110, IS901, 41 42 and IS1245 could <u>also</u> be performed.

Tuberculin test and serological tests: These tests are normally typically used to determine the disease
 prevalence of disease in a flock or to-detect infected birds. When used to detect the presence of avian
 tuberculosis in a flock, they should be supported by the necropsy of any birds that give positive reactions.

In domestic fowl, the tuberculin test in the wattle is the test of choice. This test is less useful in other species
 of bird. A better test, especially in waterfowl, is The whole blood stained-antigen agglutination test is better.
 <u>especially in</u> waterfowl. It is more reliable and has the advantage that it will can give a result within a few
 minutes while the bird is still being held.

50 **Requirements for vaccines and diagnostic biologicals:** No vaccines are available for use in birds. Avian 51 tuberculin purified protein derivative (*PPD*)-is the standard preparation for use in the tuberculin test of 52 domestic poultry. Avian PPD is also used as a component in the comparative intradermal tuberculin test in 53 cattle (see Chapter 3.1.13 Mammalian tuberculosis [infection with Mycobacterium tuberculosis complex]).

54

A. INTRODUCTION

Several mycobacterial species can be involved in the aetiology of avian tuberculosis-and, also known as avian 55 mycobacteriosis. Avian tuberculosis is most commonly produced caused by infection with Mycobacterium avium subsp. 56 57 avium (serotypes 1, 2, and 3: containing specific gene segment IS901 and nonspecific segment IS1245) and less frequently by M. genavense (Guerrero et al., 1995; Pavlik et al., 2000; Salamatian et al., 2020; Sattar et al., 2021; Tell et al., 2001). 58 Avian mycobacteriosis is also caused by other two members of the M. avium complex: M. avium subsp. hominissuis 59 (serotypes 4-6, 8-11, and 21: lacking gene segment IS901 and containing segment IS1245 and mainly infecting humans 60 and pigs) and M. intracellulare (serotypes 7, 12-20, and 22-28: lacking both gene segments IS901 and IS1245) and by 61 M. intracellulare, M. scrofulaceum, M. fortuitum, and other potentially pathogenic mycobacterial species including 62 63 M. scrofulaceum and M. fortuitum. Under some circumstances, an extensive range of mammalian species, such as swine, cattle, deer, sheep, goats, horses, cats, dogs, and exotic animals, can be infected by these mycobacterium species 64 (Dvorska et al., 2004; Kunze et al., 1992; Mijs et al., 2002; Shitaye et al., 2009; Tell et al., 2001; Thorel et al., 1997; 2001). 65 66 Mycobacterium tuberculosis and M. bovis are less common as causal rarely the causative agents of tuberculosis in birds (Hoop, 2002; Lanteri et al., 2011; Peters et al., 2007; Schmidt et al., 2022; Tell et al., 2001). 67

Mycobacterium avium species with standing in nomenclature as of 2023³⁵ (Arahal et al., 2023) consists of four-three 68 subspecies: M. avium subsp. avium, M. avium subsp. hominissuis, M. avium subsp. silvaticum, and M. avium subsp. 69 paratuberculosis (Mijs et al., 2002; Thorel et al., 1990). The latter is the causal agent of Johne's disease, or 70 71 paratuberculosis, in ruminants and other mammalian species (see Chapter 3.1.16 Paratuberculosis [Johne's disease]). 72 Mycobacterium a. silvaticum, which like M. avium subsp. paratuberculosis grows in-vitro only on media with Mycobactin, 73 which can cause avian tuberculosis in wood pigeons (Thorel et al., 1990). With the widespread use of whole genome 74 sequencing (WGS) and bioinformatics, some studies have investigated the classification of species belonging to the genus 75 Mycobacterium and have proposed that M. avium be three subspecies M. avium subsp. avium, M. avium subsp. paratuberculosis, and M. avium subsp. lepraemurium. Further subdividing M. avium subsp. avium into three variants M. 76 77 avium subsp. avium var. avium, M. avium subsp. avium var. silvaticum, and M. avium subsp. avium var. hominissuis (Riojas 78 et al., 2021; Tortoli et al., 2019).

79 All M. a avium isolates from birds and mammals, including humans, have a multiple repetitive sequence IS901 in their genome and produce a characteristic three-band pattern in IS1245 restriction fragment length polymorphism (RFLP) as 80 described and standardised previously (Dvorska et al., 2003; Ritacco et al., 1998). This repetitive sequence is also present 81 in M. a. silvaticum and RFLP analysis can help with identification. IS 901 has only been detected in M. avium strains with 82 serotypes 1, 2 and 3 (Pavlik et al., 2000; Ritacco et al., 1998) that are apparently more pathogenic to birds than other 83 serotypes (Tell et al., 2001). On the basis of genetic and phenotypic differences it has recently been proposed to 84 differentiate M. a. avium into two subspecies based on the target organism: M. a. hominissuis for human and porcine 85 isolates and M. a. avium for bird-type isolates (Mijs et al., 2002). Mycobacterium a. hominissuis has polymorphic multiband 86 IS1245 RFLP patterns and is able to grow between 24 and 45°C (Mijs et al., 2002; Van Soolingen et al., 1998). It is worth 87 noting that the typical features of bird-isolates, the three-band pattern in IS1245 RFLP and presence of IS901, have also 88 89 been found in cervine and bovine isolates of M. a. avium.

Avian tuberculosis in birds is most prevalent in gallinaceous poultry and in-wild birds raised in captivity. Turkeys are quite susceptible, but ducks, geese, and other water birds are comparatively resistant. The practices of allowing poultry to roam at large on the farm (free range) and of keeping the breeders for several years are conducive to the spread of the causal agent of avian tuberculosis among them. Infected individuals and contaminated environments (water and soil) are the main primary sources of infection. The above-mentioned mycobacterial species causing avian tuberculosis can survive for several months in the environment (Dvorska *et al.*, 2007; Kazda *et al.*, 2009; Shitaye *et al.*, 2008; Tell *et al.*, 2001).

³⁵ https://lpsn.dsmz.de/species/mycobacterium-avium

96 In most cases, Infected birds usually show no clinical signs but they may eventually become lethargic and emaciated.

97 Many affected birds show diarrhoea <u>and swollen joints</u>, and comb and wattles may regress and become pale. Affected 98 birds, especially gallinaceous poultry, are usually older than 1 year. Some show respiratory signs-and, including sudden

death may occur, dyspnoea is less common, and granulomatous ocular lesions (Pocknell *et al.*, 1996) as well as and skin

lesions have been reported. Under intensive husbandry conditions, sudden death may occur, often associated with severe

losions in the liver; such lesions are easily observed at post-mortem examination (Salamatian et al., 2020; Tell et al., 2001).

The primary lesions of avian tuberculosis in birds-poultry (chickens and turkeys) are nearly always in the intestinal tract. Such lesions take the form of deep ulcers filled with caseous material containing many mycobacterial cells, and these are discharged into the lumen and appear in the faeces. Before the intestinal tract is opened, the ulcerated areas appear as tumour-like masses attached to the gut wall, but. Still, when the intestine is opened, the true nature of the mass becomes evident. Typical caseous lesions are nearly always found in the liver and spleen, and; these organs <u>are</u> usually-are greatly enlarged because of the formation of new tuberculous tissue. The lungs and other tissues are ordinarily free from lesions even in advanced cases (Salamatian et al., 2020; Tell et al., 2001; Thorel et al., 1997).

109 Among domestic animals (mammals), domestic pigs (Sus scrofa f. domesticus) are the most susceptible to avian 110 tuberculosis. Usually, no clinical manifestations are observed in such animals. Avian tuberculosis is suspected when tuberculous lesions are found in the head and mesenteric lymph nodes on meat inspection after slaughter. Findings of 111 tuberculous lesions that involve other organs (liver, spleen, lungs, etc.) are rare, usually occurring at the advanced stage 112 of the disease. Mycobacterium a. avium accounted for up to 35% of the Mycobacteria isolated from such tuberculous 113 114 lesions (Dvorska et al., 1999; Pavlik et al., 2003, 2005; Shitaye et al., 2006). Unlike the other species mentioned previously, cattle are highly resistant to the causative agent of avian tuberculosis, and tuberculous lesions are detected in head lymph 115 nodes, or occasionally in liver lymph nodes, only on meat inspection. Mycobacterium a. avium can be successfully isolated 116 from tuberculous lesions in mesenteric lymph nodes from juvenile cattle: the isolation rate from cattle under 2 years of age 117 was 34.4% in contrast to 13.0% from cattle over 2 years of age (Dvorska et al., 2004). 118

Pet and wild birds with avian mycobacteriosis have clinical presentations often exacerbated by age and viral and fungal 119 120 co-infections (Schmidt et al., 2022; Schmitz et al., 2018b). The presence of nonspecific clinical signs and the absence of 121 gross finds during necropsy in psittacine and passeriform birds may confound diagnosis. Furthermore, differences in body 122 condition and gross pathology are observed, where psittacines have more severe lesions than passeriform birds. These 123 differences could also be attributed to the fact that they are often more likely infected with M. genavense than M. avium 124 (Schmitz et al., 2018a). The advent of more affordable WGS has allowed the study of M. avium and M. genavense and their epidemiology in a large captive population of birds belonging to multiple taxa for over 22 years. In this large bird 125 126 population, 68% of all birds at necropsy had isolates that were infected with M. avium or M. genavense. The WGS study 127 of these mycobacterium isolates demonstrated strong evidence of disease clustering among those birds infected with M. 128 avium but not among those harbouring M. genavense (Witte et al., 2021). This works sheds light on the epidemiology of 129 mycobacterium among captive birds, and future studies are necessary to understand these pathogens' epidemiology better 130 and to help identify its reservoirs.

It is essential to bear in mind that all members of *M. avium-complex* and *M. genavense* are capable of giving rise to a 131 progressive disease in humans that is refractory to treatment, especially in immunocompromised individuals (Narsana et 132 al., 2023; Pavlik et al., 2000; Tell et al., 2001). Members of Mycobacterium avium complex are classed in Risk Group 2 for 133 human infection and should be handled with appropriate measures All Mycobacterium species can cause infection in 134 people (Cowman et al., 2019). Caution should be exercised by those working with birds in environments infected with 135 Mycobacterium, especially those immunosuppressed. All laboratory manipulations with live cultures or potentially 136 137 infected/contaminated material must be performed at an appropriate biosafety and containment level determined by conducting a thorough risk assessment as described in Chapter 1.1.4 Biosafety and biosecurity: Standard for managing 138 biological risk in the veterinary laboratory and animal facilities. Biocontainment measures should be determined by risk 139 analysis as described in Chapter 1.1.4. The CDC's online Manual for Biosafety in Microbiological and Biomedical 140 141 Laboratories is also a good reference³⁶.

36 https://www.cdc.gov/labs/pdf/SF 19 308133-A BMBL6 00-BOOK-WEB-final-3.pdf

142

143

144

145

146

147

B. DIAGNOSTIC TECHNIQUES

			Ρι	ırpose		
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	Prevalence of infection – surveillance	Immune status in individual animals or populations post- vaccination
	Detection and identification of the agent ^(a)					
Ziehl–Neelsen staining	_	_	_	++	_	_
Culture	-	-	-	++	-	-
Haemagglutination (stained antigen)	ŧ	+++	÷	-	++	-
PCR	++ <u>+</u>	<u>_</u> +	<u>++</u> _	+++	<u>+</u>	-
Detection of immune response						
Haemagglutination (stained antigen)	ŧ	<u>+++</u>	<u>+</u>	Ξ	<u>++</u>	Ξ
Tuberculin test	++	+++	+	_	++	-

Table 1. Test methods available for the diagnosis of avian tuberculosis and their purpose

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; - = not appropriate for this purpose.

PCR = polymerase chain reaction.

^(a)A combination of agent identification methods applied to the same clinical sample is recommended.

148 **1. Identification of the agent**

If there is a characteristic history of avian tuberculosis in a flock and typical lesions are found in birds at post-mortem, the 149 detection of acid-fast bacilli (AFB) in smears or sections from affected organs, stained by the Ziehl-Neelsen method usually 150 151 is normally sufficient to establish a diagnosis. Confirmation of *M. avium* subspecies should be carried out by PCR or other molecular techniques (Kaevska et al., 2010; Slana et al., 2010). Occasionally a case will occur, presumably as a result of 152 due to large infecting doses giving rise to acute overwhelming disease, in which affected organs, most obviously the liver, 153 have a 'morocco leather' appearance with fine greyish or yellowish mottling. In such cases AFB may not be found in such 154 155 cases, but careful inspection will reveal parallel bundles of brownish refractile bacilli. Prolongation of the hot carbol-fuchsin stage of Ziehl-Neelsen staining to 10 minutes will usually reveal that these are indeed AFB, with unusually high resistance 156 to penetration of the stain. Recently, DNA probes and, polymerase chain reaction (PCR), and WGS techniques have been 157 used to identify the agent at the species and subspecies level specifically. Traditionally, *M. a. avium* is separated from common nonchromogenic slow-growing organisms by their ability to grow at 42°C (*M. a. avium*). *Mycobacterium* 158 159 genavense is particularly fastidious and has special unique requirements for growth and identification (Shitaye et al., 2010). 160

161 **1.1. Culture**

If there is a characteristic flock history and suggestive lesions are found at necropsy, but no AFB are seen in smears 162 or sections, an attempt must be made to isolate the causative organism from the necropsy material. The liver or 163 spleen is usually the best organ to use, but if the carcass is decomposed, bone marrow may prove more satisfactory 164 as it could be less contaminated. As with the culture of M. bovis, non-sterile specimens need to be processed with 165 166 detergent, alkali, or acid to eliminate rapidly growing microorganisms before culture (see Chapter 3.1.13 Mammalian tuberculosis [infection with Mycobacterium tuberculosis complex]). Mycobacterium a. avium grows best on media 167 such as Lowenstein-Jensen, Herrold's medium, Middlebrook 7H10-and, 7H11, or Coletsos, with 1% sodium pyruvate 168 added. It may occasionally be is necessary to incorporate mycobactin J, as it is used for the isolation of to isolate 169 170 M. a. paratuberculosis genavense and M. a. silvaticum. Growth may be confined to the edge of the condensation water. Cultures should be incubated for at least 8-12 weeks, less if using liquid media. Typically, M. a. avium produces 171 'smooth' colonies within 2-4 weeks; rough variants do-occur. Shorter incubation times can be achieved using the 172 liquid culture BACTEC system or the automated fluorescent MGIT 960 culture system. Mycobacterium a. avium can 173 also be detected in massively infected tissue by a conventional PCR, which also allows acceleration of the accelerates 174

175pathogen detection and identification (Moravkova et al., 2008). Currently, Direct detection and quantification of176M. a. avium using IS901 quantitative real-time PCR can be considered as the best fast and inexpensive method177(despite its rather high cost per test) (Kaevska et al., 2010; Slana et al., 2010).

178For *M. genavense*, the optimal medium is liquid media supplemented with Mycobactin J (an iron chelator) and then179plated onto a solid medium is such as180charcoal (Realini *et al.*, 1999). The incubation period at 37°C with 5–7% CO2 should be extended for at least 6 months18142 days. If samples are directly plated onto solid media, plates should be held for at least 12 weeks. Bacterial growth182should be prepared in a smear and stained using an acid-fast stain. All acid-fast organisms should be identified using183MALDI-TOF (matrix assisted laser desorption ionisation-time of flight [mass spectrometry]) or PCR (Buckwalter *et al.*, 2016; Hall *et al.*, 2003; Shitaye *et al.*, 2010).

Typing of mycobacteria to the species and subspecies level requires a specialised laboratory. Conventional 185 biochemical tests for species identification are lengthy and fail to distinguish between M. avium and M. intracellulare. 186 Thus, a miscellaneous group of mycobacteria that includes both species is usually classified under the denomination 187 of M. avium complex denomination. Seroagglutination, which is based on the sugar residue specificity of surface 188 glycopeptidolipids, allows classification the parsing of M. avium complex organisms into 28 serovars (Wolinsky & 189 Schaefer, 1973). More sophisticated typing methods directed at cell-wall-specific targets are currently available, such 190 as enzyme-linked immunosorbent assays with monoclonal antibodies to major serovars, and high-performance liquid 191 chromatography (HPLC), and WGS. Based on DNA-rRNA hybridisation serovars 1 to 6, 8 to 11, and 21 are currently 192 have been ascribed to M. a. avium and M. a. hominissuis, and serovars 7, 12 to 20, and 25 to M. intracellulare. 193 194 However, no consensus was achieved on other serovars, and some isolates cannot be serotyped (Inderlied et al., 1993). For final species and subspecies identification, the current methods are WGS and bioinformatic analysis of 195 isolates obtained from sick birds. Avian tuberculosis in birds is commonly caused by M. a. avium types 1, 2, or 3. If 196 the isolate is not one of these three serotypes, further molecular identification tests (IS901 PCR) must be carried out 197 conducted in a specialised laboratory. However, it should be borne in mind noted that superficial tuberculous lesions 198 in caged pet birds, especially psittacines, may be caused by M. tuberculosis, and IS6110 PCR should be used for 199 200 precise identification should always be attempted (Hoop, 2002; Lanteri et al., 2011; Peters et al., 2007; Schmidt et al., 2008; Tell et al., 2001). 201

202 **1.2.** Nucleic acid recognition methods

203 Specific and reliable genetic tests for speciation are currently have been available (Saito et al., 1990) ..., including commercial nucleic acid hybridisation probes have become a 'gold standard' reference method for distinction between 204 205 distinguishing M. avium, and M. intracellulare cultures. and M. genavense can also be distinguished with these tests. 206 A further probe that covers the whole *M. avium* complex was also developed, as genuine *M. avium* complex strains have been described that fail to react with specific M. avium and M. intracellular probes (Soini et al., 1996). 207 Nevertheless, identification errors were reported due to the cross-reactivity, which may have serious consequences 208 209 (van Ingen et al., 2009). Various in-house molecular methods have been reported for the identification of to identify mycobacterial cultures, including MAC-members of the Mycobacterium avium complex. The following gene segments 210 could be used to identify Mycobacterium isolates as M. avium in one multiplex PCR reaction: IS900, IS901, IS1245. 211 212 The isolates of M. a. avium/M. a. silvaticum are IS900-, IS901+, IS1245+, the isolates of M. a. hominissuis are 213 IS900-, IS901-, IS1245+, and the isolates of M. a. paratuberculosis are IS900+, IS901-, IS1245- (Kaevska et al., 2010; Moravkova et al., 2008). A multiplex 16S rRNA PCR and sequencing method for differentiating M. avium from 214 M. intracellulare and M. tuberculosis complex has some advantages (Cousins et al., 1996). 16S rRNA is currently 215 commercially available. Similarly, many veterinary diagnostic laboratories commonly perform in-house PCR and 216 sequencing (Kirschner et al., 1993)-may also be used. Culture-independent in-house molecular tests have been 217 developed for the detection to detect and identification of identify species belonging to the M. avium complex directly 218 from samples (Hall et al., 2003; Kaevska et al., 2010). WGS of isolates has recently become the go-to molecular 219 method to identify mycobacterium isolates from birds with great accuracy. It enables, with the use of bioinformatic 220 221 tools, not only an accurate identification of species and subspecies, but also helps to determine the organism relatedness within a flock or environment (Witte et al., 2021). In recent years, veterinary diagnostic laboratories have 222 223 extensively adopted real-time PCR methods to detect M. a. avium directly from different specimens (faeces, tissues, formalin-fixed tissues, and environmental samples). The technique rapidly detects fastidious and slow-growing 224 225 microorganisms, such as M. a. avium (Tell et al., 2003a; 2003b).

226Several commercial diagnostic PCR tests for detecting *M. a. avium* are available. Still, users should consider the skill227set and equipment necessary to perform such tests. Furthermore, it is important to determine the fitness for the228purpose of these tests before implementation. The interpretation of the results of these molecular tests also requires229veterinary expertise.

Mycobacterium a. avium, the causative agent of avian tuberculosis (Thorel et al., 1990), previously designated as
 M. avium species only, is assigned to serotypes 1 to 3 within the M. avium complex of 28 serotypes (Wolinsky &
 Schaefer, 1973). As revealed by molecular biological studies, the detected insertion sequence IS 901 (Kunze et al.,

1992) is possessed not only by the isolates of the above-named serotypes, but also by isolates, virulent for birds, that
 could not be typed because agglutination occurred (Pavlik *et al.,* 2000). In epidemiological studies, a standardised
 IS901 RFLP methods replaced serotyping (Dvorska *et al.,* 2003).

236 2. Immunological methods

Tests used for export depend on the importing requirement of individual countries. In the main, the tuberculin test or the haemagglutination (stained antigen) test are most frequently used for export testing of poultry.

239 2.1. Tuberculin test

240 The tuberculin test is the most widely used test in-for domestic fowl and the only test for which an international standard for the reagent exists. Tuberculin is the standard avian purified protein derivative (PPD). Birds are tested by 241 intradermal inoculation in the wattle with 0.05 ml or 0.1 ml of tuberculin (containing approximately 2000 International 242 Units [IU]), using a very-fine needle of approximately 26 gauge, 10 mm long × 0.5 mm. The test is read after 48 hours 243 and. A positive reaction is any swelling at the site, from a small firm nodule approximately 5 mm in diameter to gross 244 oedema extending into the other wattle and down the neck. With practice, Even very small wattles on immature birds 245 246 can be inoculated successfully. However, in immature birds the comb may be used in immature birds, although the results are not so-as reliable. Tuberculin testing of the wattle in turkeys is much less reliable consistent than in the 247 248 domestic fewl-chickens. Inoculation in the wing web has been recommended as being-more efficient, but this is still not as good as for domestic fowl-in chickens. Other birds may also be tested in the wing web, but results are not 249 250 generally satisfactory. The bare ornamental skin areas on Muscovy ducks and some species of pheasant species 251 can be used, but reliability dependability is doubtful, and interpretation is difficult. Testing in the foot web of waterfowl has also been described; the test is not very sensitive and is often complicated by infections of the inoculation site. 252

In <u>the common</u> pheasant, the tuberculin test can be performed in either of two ways. In the first, 0.05 ml or 0.1 ml of tuberculin is injected into the skin of the lower eyelid. A positive result is indicated by marked swelling at the injection site after 48 hours. Alternatively, 0.25 ml of tuberculin is injected into the thoracic muscles, and the birds are observed for 6–10 hours. Infected birds will show signs of depression and keep aside from the flock, and there may be cases of sudden death. No clinical signs will be provoked in uninfected birds.

258 2.2. Stained antigen test

267

The stained-antigen agglutination test has been used with good results, especially in domestic and ornamental 259 260 waterfowl. A drop (0.05-0.1 ml) of the antigen is mixed with the same volume of fresh whole blood, obtained by 261 venipuncture, on a white porcelain or enamel tile. The mixture is rocked for 2 minutes and examined for agglutination. 262 The agglutination may be coarse, in which case it is obvious, or quite fine, in which case it may be most clearly seen as an accumulation of the malachite-green-stained antigen around the edge of the drop, leaving the centre a normal 263 blood-red colour. This test is especially useful for screening large flocks for immediate culling and therefore has 264 265 advantages over the tuberculin test for controlling the disease, even in domestic fowl. It has also been claimed that it is more reliable in domestic poultry than the tuberculin test. 266

2.2.1. Preparation of the antigen

268 An antigen stained with 1% malachite green is used for the rapid whole blood plate agglutination test 269 (Rozanska, 1965). The strain used to prepare the stained antigen must be smooth and not auto-270 agglutinate in saline suspension. It must conform to the characteristics of *M. a. avium*, preferably obtained from a culture collection, to guarantee its authenticity. 271 272 A strain that will detect infection with any serotype is recommended instead of the specific serotype most likely to be encountered (in Europe, serotype 2 for domestic fowl, serotype 1 for waterfowl, and 273 274 birds and swine in the USA). Using a highly specific strain for the serotype is recommended. The specificity of strains can be determined only by testing them as antigens, although, in general, a 275 serotype 2 antigen will always detect serotype 3 infection and vice versa. Serotype 1 strains detect a 276 wide spectrum of infections and frequently detect infections with mycobactin-dependent 277 mycobacteria or M. a. silvaticum. There is no reason not to use a culture containing more than one 278 279 strain of *M. a. avium* if it shows the desired properties of sensitivity and specificity. Consistency of results between batches will be easier using pure cultures. 280 The organism should be grown in a suitable liquid medium, such as Middlebrook 7H9 containing 1% 281 sodium pyruvate. Good growth should be obtained in approximately 7 days. The liquid culture is used 282 as a seed for bulk antigen preparation. 283 284 Antigen for acclutination tests is best obtained on a solid medium, such as Löwenstein-Jensen or 285 7H11, containing 1% sodium pyruvate instead of glycerol, using Roux flasks or large bottles. Using

- 286 a solid medium maximizes the chance of detecting contamination, and antigens grown in some liquid 287 media are not agglutinated by specific antibodies. Liquid seed culture should be diluted (based on 288 experience) to give discrete colonies on the solid medium. This will usually give the best yield 289 increasing the chance of detecting contamination. About 10 ml of inoculum will usually allow it to wash over the whole surface and provide sufficient moisture to keep the air in the bottle near 100% 290 291 humidity.
 - The bottles are incubated at 37°C, and good growth should be obtained in 14-21 days with most strains. The antigen is harvested by adding sterile glass beads and twice the volume of sterile normal saline (containing 0.3% formalin) as was used to inoculate the bottle. The bottle is then shaken gently to wash off all the growth, and the washing is collected into a sterile bottle and re-incubated at 37°C for 7 days. The killed bacilli are washed twice in sterile normal saline with 0.2% formalin by centrifugation and re-suspension. This sequence is safer than the original method in which the washing was carried out before the incubation that kills the organisms. Finally, bacilli are again centrifuged and re-suspended in sterile normal saline containing 0.2% formalin and 0.4% sodium citrate to a concentration of about 10¹⁰ bacteria per ml. This corresponds to a concentration ten times that which matches tube No. 4 on McFarland's scale.
- 302 Cultures for antigen should be inspected for contamination daily for the first 5 days of incubation. The suspension made from the culture washings is also re-examined microscopically (for likely 303 304 contaminants such as yeasts) and rechecked by culture to ensure that the formalin has killed the 305 mycobacteria.

306 2.2.2. Validation of the antigen

Cultures should be checked by Gram staining for contamination by organisms other than mycobacteria.

One or more batches for agglutinating antigen must be tested for efficacy in using serum from 309 naturally or artificially infected tuberculous birds by comparison with a standard preparation of known 310 potency. When using animals for research or reagent testing, approval of the procedures and the use 311 312 of animals by the institution's ethics committee should be sought before any testing occurs. The potency relative to that of the standard preparation must not differ significantly from that declared on the label. Each bottle of antigen must be tested with normal chicken serum (to detect 314 autoagglutination) and M. a. avium positive chicken serum of low and high antibody content. This 315 should be done, where possible, alongside a previous batch of stained antigens. Those bottles that give satisfactory agglutination reactions with the antisera can now be pooled and the antigen stained. This is done by adding 3 ml of 1% malachite green solution per 100 ml of suspension. The stained antigen should be checked using whole blood, just as the unstained antigen was tested with serum. 320 The agglutinating antigen should stay in the refrigerator for at least 6 months at 4°C and much longer 321 if frozen at -20°C or below. If a batch has not been used for several weeks, it should be rechecked, especially for autoagglutination. 322

323 It is critical to perform a safety test of the unwashed antigen by culture and incubation to ensure that all the bacilli are dead. 324

Note on limitation of use 325

Neither the tuberculin test with avian tuberculin nor the stained-antigen agglutination test is likely to be of any value in 326 327 cases of *M. tuberculosis* infection in caged pet birds.

328

292

293

294

295

296

297

298

299

300 301

307

308

313

316

317 318

319

C. REQUIREMENTS FOR DIAGNOSTIC BIOLOGICALS

- Background 329 1.
- 330 No vaccines are available.

331 Avian tuberculin is a preparation of purified protein derivatives (PPD-A) made from the heat-treated products of growth of M. a. avium. It is used by intradermal injection to reveal delayed hypersensitivity as a means of identifying to identify birds 332 infected with or sensitised to the same species of tubercle bacillus Mycobacterium. Importantly it is also used as an to aid 333

to differential diagnosis in the comparative intradermal tuberculin test for bovine tuberculosis (see Chapter 3.1.13). An 334 335 international standard preparation of PPD-A is being developed by WOAH to replace the former WHO Standard³⁷.

336 The general principles as given-in Chapter 1.1.8 Principles of veterinary vaccine production, should be followed for injectable diagnostic biologicals such as tuberculin. The standards set out here and in chapter 1.1.8 are intended to be 337 338 general-in nature and may be supplemented by national and regional requirements.

Outline of production and minimum requirements for tuberculin production 339 2.

340 2.1. Characteristics of the seed

341

344

345

346

347

348 349

351

352

353 354

355

356

357

358

359 360

361

362

363 364

365 366

367 368

369

370 371

372

373

374

375

376

377

2.1.1. Biological characteristics of the master seed 342 343

Strains of M. a. avium used to prepare seed cultures should be purchased from a culture collection and identified as to-species by appropriate tests. Several strains are recommended by-for this purpose in different countries. For example, in the European Union (EU), for example, are, D4ER and TB56. Reference may also be made to are recommended. The relevant national recommendations should be followed. Globally there are commercial sources for PPD-A.

Quality criteria (sterility, purity, freedom from extraneous agents) 2.1.2.

Seed cultures should be shown to be free from contaminating organisms and to be capable of producing tuberculin with of sufficient potency. The necessary tests are described below.

350 22 Method of manufacture

2.2.1. Procedure

The seed material is kept as a stock of freeze-dried cultures. If the cultures have been grown on solid media, it will be necessary to adapt the organism to grow as a floating culture. This is most easily accomplished by incorporating a piece of potato in the flasks of liquid medium (e.g. Watson Reid's medium). When the culture has been adapted to a liquid medium, it can be maintained by a passage at 2-4-week intervals (Angus, 1978; Haagsma & Angus, 1995).

The organism is cultivated in modified Dorset-Henley's synthetic medium, then killed by heating in flowing steam and filtered to remove cells. The protein in the filtrate is precipitated chemically (ammonium sulphate or trichloroacetic acid-[TCA] are used), washed, and resuspended. An antimicrobial preservative that does not give rise to false-positive reactions, such as phenol (not more than 0.5% [w/v]), may be added. Mercurial derivatives should not be used. Glycerol (not more than 10% [w/v]) or glucose (2.2% [w/v]) may be added as a stabiliser. The product is dispensed aseptically into sterile neutral glass containers, which are then sealed to prevent contamination. The product may be freeze-dried.

2.2.2. **Requirements for ingredients**

The production culture substrate must be shown to be capable of producing produce a product that conforms to the standards of the European Pharmacopoeia (2000-2024 38) standards or other international standards such the WHO (WHO, 1987). It must be free from ingredients known to cause toxic or allergic reactions.

2.2.3. In-process controls

The production flasks, inoculated from suitable seed cultures, are incubated for the appropriate time period. Any flasks showing contamination or grossly abnormal growth should be discarded after autoclaving. As incubation proceeds, the surface growth of many cultures becomes moist and may sink into the medium or to the bottom of the flask. In PPD-A tuberculin, the pH of the dissolved precipitate (the so-called concentrated tuberculin) should be pH 6.6-6.7. The Kieldahl method determines the protein level (total organic nitrogen) of the PPD-A concentrate is determined by the Kieldahl method. Total nitrogen and trichloroacetic acid precipitable nitrogen are usually compared.

378 2.2.4. Final product batch tests

³⁷ PPD of M. avium tuberculin, WHO (1955) Technical Report Series, no.96, 11.

³⁸ https://www.edgm.eu/en/d/234640?p I back url=%2Fen%2Fsearch%3Fg%3Dpurified%2Bprotein%2Bderivative

6	using a procedure similar to that described in Section C.2.2.4.iv. In guinea pigs sensitised with
,	<i>M. bovis</i> , The potency of the preparation of avian tuberculin must be shown to be not more than 10%

ii)

i)

Sterility

Identity

animals for this purpose should be reviewed and approved by your institution's ethical committee.

biological materials intended for veterinary use).

iii) Safety

Tuberculin PPD<u>-A</u> can be examined for freedom from living mycobacteria using the culture method described previously. This culture method, which does not require <u>the</u> use of animals, is used in many laboratories, and its use is encouraged over the use of animals for this purpose. The following is the previously described method, using experimental animals to evaluate <u>the</u> safety of PPD. <u>The use of animals for this purpose should be reviewed and approved by the institution's ethics committee.</u> Two guinea-pigs, each weighing not less than 250 g and <u>that have</u>-not <u>been treated</u> previously <u>treated</u> with any material that will interfere with the test, are injected subcutaneously with 0.5 ml of the tuberculin under test. No abnormal effects should occur within 7 days.

Sterility testing is generally performed according to the European Pharmacopoeia (2000-2024) or

other guidelines (see-also Chapter 1.1.9 Tests for sterility and freedom from contamination of

One or more batches of tuberculin may be tested for specificity together with a standard preparation

of bovine tuberculin by comparing the reactions produced in guinea-pigs sensitised with M. bovis

of the potency of the standard preparation of bovine tuberculin used in the potency test. The use of

Tests on tuberculin for living mycobacteria may be performed either on the tuberculin immediately before it is dispensed into final containers or on samples taken from the final containers themselves. A sample of at least 10 ml must be taken and this must be injected intraperitoneally or subcutaneously into at least two guinea-pigs, dividing the volume to be tested equally between the guinea-pigs. It is desirable to take a larger sample, 50 ml, and to concentrate any residual mycobacteria by centrifugation or membrane filtration. The guinea-pigs are observed for at least 42 days and are examined macroscopically at post-mortem. Any lesions found are examined microscopically and by culture. Each filled container must be inspected before it is labelled, and any showing abnormalities must be discarded.

A test for the absence of toxic or irritant properties must be carried out <u>conducted</u> according to the <u>specifications of the</u> European Pharmacopoeia (2000<u>2024</u>) <u>specifications or the equivalent</u> <u>regulatory documents for each country or region</u>.

To test for lack of sensitising effect, three guinea-pigs that have not previously been treated with any material that could interfere with the test are each injected intradermally on each of three occasions with the equivalent of 500 IU-<u>International units</u> – one <u>IU</u> is equal to the biological activity 0.02 µg of <u>PPD</u> – of the preparation under test in a 0.1 ml volume. In the USA and Canada, the potency of the tuberculin is expressed as tuberculin unit (TU) rather than IU. One TU is also defined as 0.02 µgs of <u>PPD</u>. Each guinea-pig, together with each of the three control guinea-pigs that have not been injected previously, is injected intradermally 15–21 days after the third injection with the same dose of the same-tuberculin. The reactions of the two groups of guinea-pigs should not be significantly different when measured 24–28 hours later.

iv) Batch potency

The potency of avian tuberculin is determined in guinea-pigs sensitised with *M. a. avium, by* comparison compared with a standard preparation calibrated in IU or TU.

Use no fewer than nine albino guinea-pigs, each weighing 400–600 g. Sensitise the guinea-pigs by administering to each, by deep intramuscular injection, a suitable dose of inactivated or live *M. a. avium* to each by deep intramuscular injection. The test is performed between 4 and 6 weeks later as follows: Shave. Briefly, have the guinea-pigs' flanks so as to provide space for three-to-four injections on each side. Prepare at least three dilutions of the tuberculin under test and at least three dilutions of the standard preparation in <u>an</u> isotonic buffer solution containing 0.0005% (w/v) polysorbate 80 (Tween 80). Choose the dilutions so that the reactions produced have diameters of not less than 8 mm and not more than 25 mm. Allocate the dilutions to the injection sites randomly according to using a Latin square design. The dilutions correspond to 0.001, 0.0002, and 0.00004 mg of protein in a final dose of 0.2 ml, injected intradermally.

440 It is recommended that avian tuberculin should contain the equivalent of at least 25,000 IU/ml or 441 approximately 0.5 mg protein per ml, giving a dose for practical use of 2500 IU/0.1 ml.

442 **3.** Requirements for authorisation/registration/licensing

443 **3.1. Manufacturing process**

The manufacturing process should follow the requirements of European Pharmacopoeia (<u>2000_2024</u>) or other international standards.

446 **3.2. Safety requirements**

3.2.1. Target and non-target animal safety

448 Antimicrobial preservatives or other substances that may be added to a tuberculin must have been shown not 449 to impair the safety and effectiveness of the product. The maximum permitted concentrations for phenol is 450 0.5% (w/v), and for glycerol, it is 10% (v/v). The pH should be between 6.5 and 7.5.

451 3.2.2. Precautions (hazards)

452 Experience-both in humans and animals led to the observation that appropriately diluted tuberculin injected 453 intradermally results in a localised reaction at the injection site without generalised manifestations. Even in 454 very sensitive persons, severe, generalised reactions are extremely rare and limited.

455 **3.3. Stability**

456 During storage, liquid avian tuberculin should be protected from the light and held at a temperature of $5^{\circ}C$ ($\pm 3^{\circ}C$). 457 Freeze-dried preparations may be stored at higher temperatures (but-not exceeding $25^{\circ}C$) and protected from the 458 light. During use, periods of exposure to higher temperatures or to direct sunlight should be kept at a minimum.

459 Provided the tuberculins are Following accepted practice, tuberculin should be stored at a temperature of between 460 2°C and 8°C and protected from light; they may be used up to the end of the following periods subsequent to after 461 the last satisfactory potency test: Liquid PPD tuberculins: 2 years; lyophilised PPD-A tuberculins: 8 years; HCSM 462 (heat-concentrated synthetic-medium) tuberculins diluted: 2 years. Recent research on the temperature stability of 463 human, bovine, and avian tuberculin solutions has shown that they are stable for a year at 37°C. This should be 464 further explored as these products are used in the field in remote areas of the world where maintaining temperature 465 control is very difficult (Maes et al., 2011).

466

447

REFERENCES

467 ANGUS R.D. (1978). Production of Reference PPD tuberculins for Veterinary use in the United States. *J. Biol. Stand.*, **6**, 221.

ARAHAL D.R., BULL C.T., BUSSE H.J., CHRISTENSEN H., CHUVOCHINA M., DEDYSH S.N., FOURNIER P.E., KONSTANTINIDIS K.T.,
 PARKER C.T., ROSSELLO-MORA R., VENTOSA A. & GOKER M. (2023). Guidelines for interpreting the International Code of
 Nomenclature of Prokaryotes and for preparing a Request for an Opinion. *Int. J. Syst. Evol. Microbiol.*, 73.

BUCKWALTER S.P., OLSON S.L., CONNELLY B.J., LUCAS B.C., RODNING A.A., WALCHAK R.C., DEML S.M., WOHLFIEL S.L. &
 WENGENACK N.L. (2016). Evaluation of Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry for
 Identification of Mycobacterium species, Nocardia species, and Other Aerobic Actinomycetes. J. Clin. Microbiol., 54, 376–
 384.

476 COUSINS D., FRANCIS B. & DAWSON D. (1996). Multiplex PCR provides a low-cost alternative to DNA probe methods for rapid 477 identification of *Mycobacterium avium* and *Mycobacterium intracellulare. J. Clin. Microbiol.*, **34**, 2331–2333.

- 478 <u>COWMAN S., VAN INGEN J., GRIFFITH D.E. & LOEBINGER M.R. (2019). Non-tuberculous mycobacterial pulmonary disease. *Eur* 479 <u>.Respir. J., 54(1).</u></u>
- 480 Dvorska L., Bull T.J., Bartos M., Matlova L., Svastova P., Weston R.T., Kintr J., Parmova I., Van Soolincen D. & Pavlik
 481 I. (2003). A standardised restriction fragment length polymorphism (RFLP) method for typing *Mycobacterium avium* isolates
 482 links IS901 with virulence for birds. J. Microbiol. Methods, 55, 11–27.
- DVORSKA L., MATLOVA L., AYELE W. Y., FISCHER O. A., AMEMORI T., WESTON R. T., ALVAREZ J., BERAN V., MORAVKOVA M. &
 PAVLIK I. (2007). Avian tuberculosis in naturally infected captive water birds of the Ardeidae and Threskiornithidae families
 studied by serotyping, IS901 RFLP typing and virulence for poultry. *Vet. Microbiol.*, **119**, 366–374.
- DVORSKA L., MATLOVA L., BARTOS M., PARMOVA I., BARTL J., SVASTOVA P., BULL T. J. & PAVLIK I. (2004). Study of
 Mycobacterium avium complex strains isolated from cattle in the Czech Republic between 1996 and 2000. *Vet. Microbiol.*,
 99, 239–250.
- DVORSKA L., PARMOVA I., LAVICKOVA M., BARTL J., VRBAS V. & PAVLIK I. (1999). Isolation of *Rhodococcus equi* and atypical
 mycobacteria from lymph nodes of pigs and cattle in herds with the occurrence of tuberculoid gross changes in the Czech
 Republic over the period of 1996-1998. Veterinarni Medicina, 44, 321–330.
- EUROPEAN PHARMACOPOEIA (2000-2024). Purified protein derivative (avian). *In:* European Pharmacopoeia, Eleventh Edition.
 Editions of the Council of Europe, Strasbourg, France, https://www.edgm.eu/en/d/234640?p.l.back_url=%2Fen%2Fsearch%3Fg%3Dpurified%2Bprotein%2Bderivative
- 495 GUERRERO C., BERNASCONI C., BURKI D., BODMER T. & TELENTI A. (1995). A novel insertion element from *Mycobacterium* 496 *avium*, IS1245, is a specific target for analysis of strain relatedness. *J. Clin. Microbiol.*, **33**, 304–307.
- HAAGSMA J. & ANGUS R.D. (1995). Tuberculin production. *In: Mycobacterium bovis* Infections in Humans and Animals,
 Steele J.H. & Thoen C.O., eds. Iowa State University Press, Ames, USA, 73–84.
- HALL L., DOERR K.A., WOHLFIEL S.L. & ROBERTS G.D. (2003). Evaluation of the MicroSeq system for identification of mycobacteria by 16S ribosomal DNA sequencing and its integration into a routine clinical mycobacteriology laboratory. J. *Clin. Microbiol.*, **41**, 1447–1453.
- 502 <u>HOOP R. (2002). Mycobacterium tuberculosis infection in a canary (Serinus canaria L.) and a blue-fronted Amazon parrot</u> 503 <u>(Amazona amazona aestiva). Avian Dis., **46**, 502–504.</u>
- INDERLIED C.B., KEMPER C.A. & BERMUDEZ L.E.M. (1993). The *Mycobacterium avium* complex. *Clin. Microbiol. Rev.*, 6, 266–310.
- 506 KAEVSKA M., SLANA I., KRALIK P. & PAVLIK I. (2010). Examination *of Mycobacterium avium* subsp. *avium* distribution in 507 naturally infected hens by culture and triplex quantitative real time PCR. *Veterinarni Medicina*, **55**, 325–330.
- 508 KAZDA J., PAVLIK I., FALKINHAM J. & HRUSKA K. (2009). The Ecology of *Mycobacteria*: Impact on Animal's and Human's 509 Health, First Edition, Springer Science+Business Media BV, 520 pp. ISBN 978-1-4020-9412-5.
- 510 KUNZE Z.M., PORTAELS F. & MCFADDEN J.J. (1992). <u>Biologically distinct subtypes of *Mycobacterium avium* differ in 511 possession of insertion sequence IS901. J. Clin. Microbiol., **30**, 2366–2372.</u>
- KIRSCHNER P., MEIER P.A. & BOTTGER E.C. (1993). Genotypic identification and detection of mycobacteria. *In:* Diagnostic
 Molecular Microbiology, Persing D.H., Smith T.F., Tenover F.C. & White T.C., eds. American Society for Microbiology,
 Washington DC, USA, 173–190.
- 515 <u>LANTERI G., MARINO F., REALE S., VITALE F., MACRI F. & MAZZULLO G. (2011). Mycobacterium tuberculosis in a red-crowned</u>
 516 parakeet (*Cyanoramphus novaezelandiae*). J. Avian Med. Surg., 25, 40–43.
- 517 <u>MAES M., GIMÉNEZ J.F., D'ALESSANDRO A. & DE WAARD J.H. (2011). The stability of human, bovine and avian tuberculin</u> 518 <u>purified protein derivative (PPD). J. Infect. Dev. Ctries</u>, **5**, 781–785.

519 MIJS W., DE HAAS P., ROSSAU R., VAN DER LAAN T., RIGOUTS L., PORTAELS F. & VAN SOOLINGEN D. (2002). Molecular evidence 520 to support a proposal to reserve the designation *Mycobacterium avium* subsp. *avium* to bird-type isolates and *M. avium* 521 subsp. *hominissuis* for the human/porcine type of *M. avium*. *Int. J. Syst. Evol. Microbiol.*, **52**, 1505–1518.

MORAVKOVA M., HLOZEK P., BERAN V., PAVLIK I., PREZIUSO S., CUTERI V. & BARTOS M. (2008). Strategy for the detection and differentiation of *Mycobacterium avium* species in isolates and heavily infected tissues. *Res. Vet. Sci.*, **85**, 257–264.

- 524 <u>NARSANA N., ALEJANDRA PEREZ M. & SUBRAMANIAN A. (2023). Mycobacteria in Organ Transplant Recipients. Infect. Dis. Clin.</u>
 525 <u>North Am., S0891-5520(23)00040-5. doi: 10.1016/j.idc.2023.04.004.</u>
- PAVLIK I., MATLOVA L., DVORSKA L., BARTL J., OKTABCOVA L., DOCEKAL J. & PARMOVA I. (2003). Tuberculous lesions in pigs in the
 Czech Republic during 1990-1999: occurrence, causal factors and economic loses. Veterinarni Medicina, 48, 113–125.
- PAVLIK I., MATLOVA L., DVORSKA L., SHITAYE J. E. & PARMOVA I. (2005). Mycobacterial infections in cattle and pigs caused by
 Mycobacterium avium complex members and atypical mycobacteria in the Czech Republic during 2000–2004. Veterinami
 Medicina. 50, 281–290.
- PAVLIK I., SVASTOVA P., BARTL J., DVORSKA L. & RYCHLIK I. (2000). Relationship between IS901 in the *Mycobacterium avium* complex strains isolated from birds, animals, humans, and the environment and virulence for poultry. *Clin. Diagn. Lab. Immunol.*, 7, 212–217.
- 534 <u>PETERS M., PRODINGER W.M., GUMMER H., HOTZEL H., MOBIUS P. & MOSER I. (2007). Mycobacterium tuberculosis infection in</u>
 535 <u>a blue-fronted amazon parrot (*Amazona aestiva aestiva). Vet. Microbiol.*, **122**, 381–383.
 </u>
- 536 <u>POCKNELL A.M., MILLER B.J., NEUFELD J.L.</u> & <u>GRAHN B.H.</u> (1996). Conjunctival mycobacteriosis in two emus (*Dromaius* 537 *novaehollandiae*). <u>Vet. Pathol.</u>, **33**, 346–348.
- 538 REALINI L., DE RIDDER K., HIRSCHEL B. & PORTAELS F. (1999). Blood and charcoal added to acidified agar media promote the
 539 growth of *Mycobacterium genavense*. *Diagn. Microbiol. Infect. Dis.*, **34**, 45–50.
- RIOJAS M.A., FRANK A.M., GREENFIELD S.R., KING S.P., MEEHAN C.J., STRONG M., WATTAM A.R. & HAZBÓN M.H. (2021).
 Identification and Characterization of Mycobacterial Species Using Whole-Genome Sequences. *Methods Mol. Biol.*, 2314, 399–457.
- 543 RITACCO V., KREMER K., VAN DER LAAN T., PUNENBURG J.E.M., DE HAAS P.E.W. & VAN SOOLINGEN D. (1998). Use of IS901
 544 and IS1245 in RFLP typing of *Mycobacterium avium* complex: relatedness among serovar reference strains, human and
 545 animal isolates. *Int. J. Tuberculosis Lung Dis.*, 2, 242–251.
- ROZANSKA M. (1965). Preparation of antigen for whole blood rapid agglutination test and its specificity for diagnosis of avian
 tuberculosis. *Bull. Vet. Inst. Pulawy*, 9, 20–25.
- 548 SAITO H., TOMIOKA H., SATO K., TASAKA H. & DAWSON D.J. (1990). Identification of various serovar strains of *Mycobacterium* 549 *avium* complex by using DNA probes specific for *Mycobacterium avium* and *Mycobacterium intracellulare. J. Clin.* 550 *Microbiol.*, **28**, 1694–1697.
- 551 <u>SALAMATIAN I., GHANIEI A., MOSAVARI N., NOURANI H., KESHAVARZ R. & ESLAMPANAH M. (2020). Outbreak of avian</u> 552 <u>mycobacteriosis in a commercial turkey breeder flock. *Avian Pathol.*, **49**, 296–304.</u>
- 553 <u>SATTAR A., ZAKARIA Z., ABU J., AZIZ S.A. & ROJAS-PONCE G. (2021). Isolation of Mycobacterium avium and other</u> 554 <u>nontuberculous mycobacteria in chickens and captive birds in peninsular Malaysia. *BMC Vet. Res.*, **17**, 13.</u>
- 555 <u>SCHMIDT V., KOHLER H., HEENEMANN K. & MOBIUS P. (2022). Mycobacteriosis in Various Pet and Wild Birds from Germany:</u>
 556 <u>Pathological Findings, Coinfections, and Characterization of Causative Mycobacteria. *Microbiol. Spectr.*, 10(4): e0045222.
 </u>
- 557 <u>SCHMIDT V., SCHNEIDER S., SCHLOMER J., KRAUTWALD-JUNGHANNS M.E. & RICHTER E. (2008). Transmission of tuberculosis</u> 558 <u>between men and pet birds: a case report. *Avian Pathol.*, **37**, 589–592.</u>
- 559 <u>SCHMITZ A., KORBEL R., THIEL S., WORLE B., GOHL C. & RINDER M. (2018a). High prevalence of Mycobacterium genavense</u> 560 <u>within flocks of pet birds. *Vet. Microbiol.*, **218**, 40–44.</u>
- 561 <u>SCHMITZ A., RINDER M., THIEL S., PESCHEL A., MOSER K., REESE S. & KORBEL R. (2018b). Retrospective Evaluation of Clinical</u> 562 <u>Signs and Gross Pathologic Findings in Birds Infected With *Mycobacterium genavense. J. Avian Med. Surg.*, **32**, 194–204.</u>
- 563 SHITAYE J.E., GRYMOVA V., GRYM M., HALOUZKA R., HORVATHOVA A., MORAVKOVA M., BERAN V., SVOBODOVA J., DVORSKA-564 BARTOSOVA L. & PAVLIK I. (2009). *Mycobacterium avium* subsp. *hominissuis* infection in a pet parrot. *Emerg. Inf. Dis.*, **15**, 565 617–619.
- 566 SHITAYE J.E., HALOUZKA R., SVOBODOVA J., GRYMOVA V., GRYM M., SKORIC M., FICTUM P., BERAN V., SLANY M. & PAVLIK I. 567 (2010). First isolation of *Mycobacterium genavense* in blue headed parrot (*Pionus menstruus*) imported from Surinam 568 (South America) to the Czech Bepublic: a case report. *Veterinarni Medicina* **55**, 320, 347
- 568 (South America) to the Czech Republic: a case report. *Veterinarni Medicina*, **55**, 339–347.

SHITAYE J.E., MATLOVA L., HORVATHOVA A., MORAVKOVA M., DVORSKA-BARTOSOVA L., TREML F., LAMKA J. & PAVLIK I. (2008).
 Mycobacterium avium subsp. *avium* distribution studied in a naturally infected hen flock and in the environment by culture,
 serotyping and IS901 RFLP methods. *Vet. Microbiol.*, **127**, 155–164.

572 SHITAYE J.E., PARMOVA I., MATLOVA L., DVORSKA L., HORVATHOVA A., VRBAS V. & PAVLIK, I. (2006). Mycobacterial and 573 *Rhodococcus equi* infections in pigs in the Czech Republic between the years 1996 and 2004: the causal factors and 574 distribution of infections in the tissues. *Veterinarni Medicina*, **51**, 497–511.http://www.vri.cz/docs/vetmed/51-11-497.pdf

- 575 SLANA I., KAEVSKA M., KRALIK P., HORVATHOVA A. & PAVLIK, I. (2010). Distribution of *Mycobacterium avium* subsp. *avium* and 576 *M. a. hominissuis* in artificially infected pigs studied by culture and IS*901* and IS*1245* quantitative real time PCR. *Vet.* 577 *Microbiol.*, **144**, 437–443.
- 578 SOINI H., EEROLA E. & VILJANEN M.K. (1996). Genetic diversity among *Mycobacterium avium* complex Accu-Probe-positive 579 isolates. *J. Clin. Microbiol.*, **34**, 55–57.
- 580 <u>TELL L.A., FOLEY J., NEEDHAM M.L. & WALKER R.L. (2003a). Comparison of four rapid DNA extraction techniques for</u> 581 <u>conventional polymerase chain reaction testing of three *Mycobacterium* spp. that affect birds. *Avian Dis.*, **47**, 1486–1490.</u>

TELL L.A., LEUTENEGGER C.M., LARSEN R.S., AGNEW D.W., KEENER L., NEEDHAM M.L. & RIDEOUT B.A. (2003b). Real-time
 polymerase chain reaction testing for the detection of *Mycobacterium genavense* and *Mycobacterium avium* complex
 species in avian samples. *Avian. Dis.*, 47, 1406–1415.

- 585 TELL L.A., WOODS L. & CROMIE R.L. (2001). Mycobacteriosis in birds. Rev. Sci. Tech., 20, 180–203.
- THOREL M.F., HUCHZERMEYER H. & MICHEL A.L. (2001). *Mycobacterium avium* and *M. intracellulare* infection in mammals. *Rev. sci. tech. Off. int. Epiz.*, **20**, 204–218.
- 588 THOREL M.F., HUCHZERMEYER H., WEISS R. & FONTAINE J.J. (1997). *Mycobacterium avium* infections in animals. Literature 589 review. *Vet. Res.*, **28**, 439–447.

THOREL M.F., KRICHEVSKY M. & LEVY-FREBAULT V.V. (1990). Numerical taxonomy of mycobactin-dependent mycobacteria,
 emended description of *Mycobacterium avium*, and description of *Mycobacterium avium* subsp. *avium* subsp. nov.,
 Mycobacterium avium subsp. *paratuberculosis* subsp. nov., and *Mycobacterium avium* subsp. *silvaticum* subsp. nov. *Int. J. Syst. Bacteriol.*, **40**, 254–260.

- TORTOLI E., MEEHAN C.J., GROTTOLA A., FREGNI SERPINI G., FABIO A., TROVATO A., PECORARI M. & CIRILLO D.M. (2019).
 Genome-based taxonomic revision detects a number of synonymous taxa in the genus *Mycobacterium*. *Infect. Genet. Evol.*, **75**, 103983.
- VAN INGEN J., AL HAJOJ SAM., BOERE M., AL RABIAH F., ENAIMI M., DE ZWAAN R., TORTOLI E., DEKHUIJZEN R. & VAN SOOLINGEN
 D. (2009). *Mycobacterium riyadhense* sp. nov.; a non-tuberculous species identified as *Mycobacterium tuberculosis* by a
 commercial line-probe assay. *Int. J. Syst. Evol. Microbiol.*, **59**, 1049–1053.
- VAN SOOLINGEN D., BAUER J., RITACCO V., CARDOSO LEAO S., PAVLI I., VINCENT V., RASTOGI N., GORI A., BODMER T., GARZELLI
 C. & GARCIA M.J. (1998). IS1245 restriction fragment length polymorphism typing of *Mycobacterium avium* isolates:
 proposal for standardization. J. Clin. Microbiol., 36, 3051–3054.
- MITTE C., FOWLER J.H., PFEIFFER W., HUNGERFORD L.L., BRAUN J., BURCHELL J., PAPENDICK R. & RIDEOUT B.A. (2021). Social
 network analysis and whole-genome sequencing to evaluate disease transmission in a large, dynamic population: A study
 of avian mycobacteriosis in zoo birds. *PLoS One*, **16**(6): e0252152.
- WOLINSKY E. & SCHAEFER W.B. (1973). Proposed numbering scheme for mycobacterial serotypes by agglutination. *Int. J. Syst. Bacteriol.*, **23**, 182–183.
- WORLD HEALTH ORGANIZATION (WHO) (1987). Requirements for Biological Substances No. 16, Annex 1: Requirement for
 Tuberculins. Technical Report Series No. 745, WHO, Geneva, Switzerland, 31–59.

610 611	* *
612 613 614	NB: There is currently (2024) no WOAH Reference Laboratory for avian tuberculosis (please consult the WOAH Web site for the current list: <u>https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3http://www.oie.int/</u>).
615	NB: FIRST ADOPTED IN 1989 AS TUBERCULOSIS IN BIRDS. MOST RECENT UPDATES ADOPTED IN 2014.

Annex 10. Item 5.1. – Chapter 3.4.1. Bovine anaplasmosis MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION Paris, 4–8 September 2023

1

2

3

4

5

6

7

8

9

SECTION 3.4.

BOVINAE

CHAPTER 3.4.1.

BOVINE ANAPLASMOSIS

SUMMARY

10Definition of the disease: Bovine anaplasmosis results from infection with Anaplasma marginale. A second11species, A. centrale, has long been recognised and usually causes benign infections. Anaplasma marginale12is responsible for almost all outbreaks of clinical disease. Anaplasma phagocytophilum and A. bovis, which13infect cattle, have been recently are also included within the genus-but they are not reported to. Anaplasma14phagocytophilum can cause clinical self-limiting disease in cattle. There are no reports of disease associated15with A. bovis infection. The organism is classified in the genus Anaplasma belonging to the family16Anaplasmataceae of the order Rickettsiales.

17Description of the disease: Anaemia, jaundice in acute, severe cases and sudden-unexpected death are18characteristic signs of bovine anaplasmosis. Other signs include rapid loss of milk production and weight,19but the clinical disease can only be confirmed by identifying the organism. Once infected, cattle may remain20carriers for life, and identification of these animals depends on the detection of specific antibodies using21serological tests, or of rickettsial DNA using molecular amplification techniques. The disease is typically22transmitted by tick vectors, but mechanical transmission by biting insects or by needle can occur.

23 Detection Identification of the agent: Microscopic examination of blood or organ smears stained with 24 Giemsa stain is the most common method of identifying Anaplasma in clinically affected animals. In these 25 smears, A. marginale organisms appear as dense, rounded, intraerythrocytic bodies approximately 0.3-26 1.0 µm in diameter situated on or near the margin of the erythrocyte. Anaplasma centrale is similar in 27 appearance, but most of the organisms are situated toward the centre of the erythrocyte. It can be difficult 28 to differentiate A. marginale from A. centrale in a stained smear, particularly with low levels of rickettsaemia. 29 Commercial stains that give very rapid staining of Anaplasma spp. are available in some countries. 30 Anaplasma phagocytophilum can only be observed in infected granulocytes, mainly neutrophils and A. bovis 31 can only be observed in infected monocytes_infecting granulocytes, mainly neutrophils.

It is important that smears be well prepared and free from foreign matter. Smears from live cattle should
 preferably be prepared from blood drawn from the jugular vein or another large vessel. For post-mortem
 diagnosis, smears should be prepared from internal organs (including liver, kidney, heart and lungs) and

from blood retained in peripheral vessels. The latter are particularly desirable <u>useful</u> if post-mortem decomposition is advanced.

37 Serological tests: A competitive enzyme-linked immunosorbent assay (C-ELISA) has been demonstrated 38 to have good sensitivity in detecting carrier animals. Card agglutination is the next most frequently used 39 assay. The complement fixation test (CFT) is no longer considered a reliable test for disease certification of 40 individual animals due to variable sensitivity. Cross reactivity between Anaplasma spp. can complicate 41 interpretation of serological tests. In general, the C-ELISA has the best specificity, with cross-reactivity 42 described between A. marginale, A. centrale, A. phagocytophilum and Ehrlichia spp. Alternatively, an 43 indirect ELISA using the CFT with modifications (I-ELISA) is a reliable test used in many laboratories and can be prepared in-house for routine diagnosis of anaplasmosis. Finally, a displacement double-antigen 44 sandwich ELISA has been developed to differentiate between A. marginale and A. centrale antibodies. 45

- 46 Nucleic-acid-based tests have been used are often used in diagnostic laboratories and experimentally, 47 and are capable of detecting the presence of low-level infection in carrier cattle and tick vectors. A nested 48 conventional polymerase chain reaction (PCR) reaction is necessary has been used to identify low-level 49 carriers using conventional polymerase chain reaction (PCR), and although nonspecific amplification can 50 occur. Recently, Real-time PCR assays with have analytical sensitivity equivalent to nested conventional 51 PCR have been described and are preferable in a diagnostic setting to reduce the risk of amplicon 52 contamination.
- 53 **Requirements for vaccines:** Live vaccines are used in several countries to protect cattle against 54 A. marginale infection bovine anaplasmosis. A vaccine consisting of live A. centrale is most widely used and 55 gives partial protection against challenge with virulent A. marginale. <u>Vaccination with A. centrale leads to</u> 56 infection and long-term persistence in many cattle. Vaccinated cattle are typically protected from disease 57 caused by A. marginale, but not infection.
- 58 Anaplasma centrale vaccine is provided in chilled or frozen forms. Quality control is very important as other 59 blood-borne agents that may be present in donor cattle can contaminate vaccines and be disseminated 60 broadly. For this reason, frozen vaccine is recommended as it allows thorough post-production quality control, 61 which limits the risk of contamination with other pathogens.
- Anaplasma centrale vaccine is not entirely safe. A practical recommendation is to restrict its use, as far as possible, to calves, as nonspecific immunity will minimise the risk of some vaccine reactions that may require treatment with tetracycline or imidocarb. Partial immunity develops in 6–8 weeks and lasts for several years after a single vaccination. In countries where A. centrale is exotic, it cannot be used as a vaccine against A. marginale.

67

A. INTRODUCTION

68 Outbreaks of bovine anaplasmosis are due to infection with Anaplasma marginale. Anaplasma centrale is capable of 69 producing can produce a moderate degree of anaemia, but clinical outbreaks in the field are extremely rare. New species 70 of Anaplasma, Other members of the family Anaplasmataceae that infect cattle include A. phagocytophilum and A. bovis 71 (Dumler et al., 2001), with a primary reservoir. Anaplasma phagocytophilum has a broad host range and causes the 72 diseases human granulocytic anaplasmosis (HGE), equine granulocytic anaplasmosis (EGA), and canine granulocytic 73 anaplasmosis (CGA), in humans, horses, and dogs, respectively (Matei et al., 2019). In northern Europe-in rodents, A. 74 phagocytophilum causes tick-borne fever, primarily affecting lambs. In cattle, A. phagocytophilum infections have been 75 reported to infect cattle, but do not cause from many geographical regions, however the association with disease is less 76 commonly reported. Naturally occurring clinical disease as reported in Germany was characterised by fever (39.5-41.7° 77 C), sudden reduction in milk production, lower limb oedema, and stiffness with leukopenia, erythropenia, neutropenia, lymphocytopenia and monocytopenia. The affected animals recovered without antibiotic treatment (Dreher et al., 2005; 78 Hofmann-Lehmann et al., 2004 Silaghi et al., 2018). 79

The most marked clinical signs of <u>bovine</u> anaplasmosis are anaemia and jaundice, the latter occurring <u>in acute severe</u>, <u>cases or</u> late in the disease. Haemoglobinaemia and haemoglobinuria are not present, and this may assist in the differential diagnosis of <u>bovine</u> anaplasmosis from babesiosis, which is often endemic in the same regions. The disease can only be confirmed, however, by identification of the organism <u>in erythrocytes from the affected animal</u>. <u>Caution must be exercised</u> if using nucleic acid techniques alone to diagnose *A. marginale* in anaemic cattle. Persistent, low-level infection can be detected by these techniques and may lead to a misdiagnosis of bovine anaplasmosis. Visualisation of *A. marginale* bodies
 in erythrocytes is therefore required for confirmation.

Anaplasma marginale occurs in most tropical and subtropical countries and <u>is widely distributed</u> in some more temperate regions. Anaplasma centrale was first described from South Africa. The organism has since been imported by other countries – including Australia and some countries in South America, South-East Asia and the Middle East – for use as a vaccine against *A. marginale.*

91 Anaplasma species-were, though originally regarded described as protozoan parasites, but further research showed they had no significant attributes to justify this description. Since the last major accepted revision of the are obligate intracellular 92 Gram-negative bacteria. Based on taxonomy established in 2001 (Dumler et al., 2001), the Family Anaplasmataceae 93 94 (Order Rickettsiales) is now-composed of four-five genera, Anaplasma, Ehrlichia, Neorickettsia, and-Wolbachia. The genus and Aegyptianella is retained within the Family Anaplasmataceae as genus incertae sedis. The revised genus. The genus 95 Anaplasma new contains Anaplasma marginale as the type species, A. phagocytophilum the agent of human granulocytic 96 97 ehrlichiosis (formerly Ehrlichia phagocytophila and E. equi), A. platys, and A. bovis (formerly E. bovis). Haemobartonella and Eperythrozoon are now considered most closely related to the mycoplasmas. 98

Anaplasma species are transmitted either mechanically or biologically by arthropod vectors. Reviews based on careful 99 100 study Detection of reported transmission experiments list up pathogen DNA within a tick is insufficient to 19 different ticks as capable of determine the ability of a particular tick species to transmit a pathogen. Studies demonstrating transmission 101 102 of the pathogen are critical in determining the potential role of a particular tick species in pathogen transmission transmitting A. marginale (Kocan et al., 2004). These are: Argas persicus, Ornithodoros lahorensis,. Many studies have demonstrated 103 D. variabilis. 104 the transmission ability of Dermcentor albipictus, D. andersoni, D. hunteri, D. occidentalis, 105 Hyalomma excavatum, H. rufipes, Ixodes ricinus, I. scapularis, and D. albipictus. Additionally, transmission by multiple Rhipicephalus species is well recognised including R. annulatus (formerly Boophilus annulatus), R. bursa, R. calcaratus, 106 R. decoloratus, R. evertsi, R. microplus, R. sanguineus and R. simus. However, the classification of several ticks in these 107 108 reports has been questioned. and R. sanguineous. Other species of Rhipicephalus also likely serve as biological vectors of A. marginale. Anaplasma marginale DNA has been widely reported in Hyalomma species, and transmission has been 109 110 demonstrated with H. excavatum. It is likely that multiple Hyalomma species also serve as vectors of A. marginale (Shkap et al., 2009). 111

Intrastadial or transstadial transmission is the usual mode-<u>can occur</u>, even in the one-host, *Rhipicephalus* species. Male ticks may be particularly important as vectors, <u>as</u> they can become persistently infected and serve as a reservoir are most likely to move between cattle searching for infection female ticks. Experimental demonstration of vector competence does not necessarily imply a role in transmission in the field. However, *Rhipicephalus* species are clearly important vectors of anaplasmosis in countries such as Australia and countries in, many regions of Africa, and Latin America, and some species of <u>Africa</u>, and Latin America, and some species of <u>Africa</u>.

118 Various other biting arthropods have been implicated as mechanical vectors, particularly in the USA. Experimental 119 transmission has been demonstrated with a number of species of *Tabanus* (horseflies), and with mosquitoes of the genus 120 *Psorophora* (Kocan *et al.*, 2004). The importance of biting insects in the natural transmission of anaplasmosis appears to 121 vary greatly from region to region. *Anaplasma marginale* also can be readily transmitted during vaccination against other 122 diseases unless a fresh or sterilised needle is used for injecting each animal. Similar transmission by means of unsterilised 123 surgical instruments has been described (Reinbold *et al.*, 2010a).

124 The main-only known biological vectors of *A. centrale* appear to be multihost ticks is *R. simus*, endemic in Africa, including 125 *R. simus*. The, Though multiple transmission studies have been done, there is no evidence that the common cattle tick 126 (*R. microplus*) has not been shown to be can serve as a vector for *A. centrale*. This is of relevance relevant where 127 *A. centrale* is used as a vaccine in *R. microplus*-infested regions.

Anaplasma marginale infection has not been reported in humans. Thus, There is no-minimal risk of field or laboratory transmission to workers and from laboratories working with *A. marginale*-may operate at the lowest biosafety level, equivalent to BSL1. Nevertheless the agent should be handled with appropriate biosafety and containment procedures as determined by biorisk analysis (see Chapter 1.1.4 Biosafety and biosecurity: Standard for managing biological risk in the veterinary laboratory and animal facilities).

133

134

Table 1. Test methods available for the diagnosis of bovine anaplasmosis and their purpose

B. DIAGNOSTIC TECHNIQUES

			Pur	pose			
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	<u>Contribute to</u> <u>eradication</u> <u>policies</u>	<u>Confirmation</u> of clinical <u>cases</u>	<u>Prevalence</u> of infection – <u>surveillance</u>	Immune status in individual animals or populations (post- vaccination)	
Microscopic examination	-	* =	-	+++	-	_	
	Detection of the agent ^(a)						
PCR	-	++ +	-	+++	-	_	
		Detectio	on of immune r	esponse			
CAT ^{<u>(b)</u>}	_	_	_	_	+	+	
<u>C-</u> ELISA ^(b)	+++	+ <u>+++</u>	+++	-	+++	+++	
IFAT <u>ట</u>	+	_	_	_	++	++	
GFT	-	-	-	-	+	-	
<u>ddasELISA</u>	=	=	=	=	E	<u>++</u>	

135 136 137

138 139

140

141

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; – = not appropriate for this purpose.

Agent id. = agent identification; CAT = card agglutination test; CFT = complement fixation test; C-ELISA = competitive enzyme-linked immunosorbent assay; ddasELISA = displacement double-antigen, sandwich ELISA;

IFAT = indirect fluorescent antibody test; PCR = polymerase chain reaction.

^(a)A combination of agent identification methods applied on the same clinical sample is recommended.

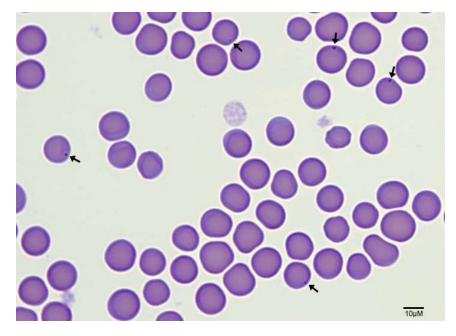
(b)These tests do not distinguish infected from vaccinated animals.

142 **1. Detection of the agent**

143 **1.1. Microscopic examination**

Samples from live cattle should include thin blood smears and blood collected into an anticoagulant. Air-dried thin blood smears can be kept satisfactorily at room temperature for at least 1 week. The blood sample in anticoagulant should be held and transferred at 4°C, unless it can reach the laboratory within a few hours. This sample is useful for preparing fresh smears if those submitted are not satisfactory. In addition, a low packed cell volume and/or erythrocyte count can help to substantiate the involvement of *A. marginale* when only small numbers of the parasites are detected in smears, for example <u>particularly</u> during the recovery stage of the disease.

150 In contrast to Babesia bovis, A. marginale-does-infected erythrocytes do not accumulate in capillaries, so blood drawn from the jugular or other large vessel is satisfactory. Anaplasma marginale replicate in the erythrocytes to form small 151 membrane-bound colonies, also termed inclusion bodies or initial bodies. Because of the rather indistinctive 152 morphology of Anaplasma-These initial bodies can be visualised on a blood smear, but are small and easily confused 153 with debris or stain precipitate (see Figure 1). Thus it is essential that smears are well prepared and . including 154 ensuring slides are free from foreign matter, as specks of debris can confuse diagnosis and stain is recently filtered 155 (Watman #1 filter paper). Thick blood films as are used sometimes for the diagnosis of babesiosis are not appropriate 156 for the diagnosis of anaplasmosis, as Anaplasma A. marginale are difficult to identify once they become dissociated 157 158 from erythrocytes.



160 Figure. 1. Anaplasma marginale initial bodies. A Diff-Quick stained blood smear from a bovine experimentally infected 161 with A. marginale. Arrows point to the A. marginale initial bodies. Photo from S. Noh.

159

- Samples from dead animals should include air-dried thin smears from the liver, kidney, heart and lungs and from a peripheral blood vessel. The latter is particularly recommended should there be a significant delay before postmortem examination because, under these circumstances, bacterial contamination of organ smears often makes identification of *Anaplasma <u>A.</u> marginale* equivocal. Brain smears, which are useful for the diagnosis of some forms of babesiosis, are of no direct value for diagnosing anaplasmosis, but should be included for differential diagnosis where appropriate.
- 168 Blood from organs, rather than organ tissues *per se*, is required for smear preparation, as the aim is to be able to 169 examine microscopically intact erythrocytes for the presence of *Anaplasma-A. marginale colonies*. Organ-derived 170 blood smears can be stored satisfactorily at room temperature for several days.
- 171 Both blood and organ smears can be stained in 10% Giemsa stain for approximately 30 minutes after fixation in 172 absolute methanol for 1 minute. After staining, the smears are rinsed three or four times with tap water to remove 173 excess stain and are then air-dried. Conditions for Giemsa staining vary from laboratory to laboratory, but distilled 174 water is not recommended for dilution of Giemsa stock. Water should be pH 7.2–7.4 to attain best resolution with 175 Giemsa stain. Commercial stains that give very rapid staining of *Anaplasma* <u>A</u> <u>marginale</u> are available in some 176 countries. Smears are <u>must be</u> examined under oil immersion at a magnification of ×700–1000.
- Anaplasma marginale appear as dense, initial bodies are rounded and deeply stained intraerythrocytic bodies, and
 approximately 0.3–1.0 μm in diameter. Most of these bodies are located on or near the margin of the erythrocyte.
 This feature distinguishes *A. marginale* from *A. centrale*, as in the latter most of the organisms have a more central
 location in the erythrocyte. However, particularly at low levels of rickettsaemia, differentiation of these two species in
 smears can be difficult. Appendages associated with the *Anaplasma* body initial body have been described in some
 isolates of *A. marginale* (Kreier & Ristic, 1963; Stich *et al.*, 2004).
- 183 The percentage of infected erythrocytes varies with the stage and severity of the disease. Maximum rickettsaemias 184 in excess of 50% may occur with *A. marginale*. Multiple infections of individual erythrocytes are common during 185 periods of high rickettsaemias.
- The infection becomes visible microscopically 2–6 weeks following transmission. During the course of clinical disease, the rickettsaemia approximately doubles each day for up to about 10 days, and then decreases at a similar rate. Severe anaemia may persist for some weeks after the parasites have become virtually undetectable in blood smears. Following recovery from initial infection, cattle remain latently infected for life.

190 **1.2.** Polymerase chain reaction

191 Nucleic-acid-based tests to detect A. marginale-infection in carrier-infected cattle have been developed although not 192 vet fully validated. The analytical sensitivity of polymerase chain reaction (PCR)-based methods has been estimated at 0.0001% infected erythrocytes, but at this level, only a proportion of carrier cattle would be detected. A nested PCR 193 has been used to identify A. marginale carrier cattle with a capability of identifying as few as 30 infected erythrocytes 194 per ml of blood, well below the lowest levels in carriers. However, nested PCR is time consuming as it requires two 195 196 full PCR reactions, and poses significant quality control and specificity problems for routine use (Torioni De Echaide 197 et al., 1998). Real-time PCR assays are reported to achieve a level of analytical sensitivity equivalent to nested PCR has also been described for identification of A. marginale and should be considered instead of the nested PCR (Carelli 198 et al., 2007; Decaro et al., 2008; Reinbold et al., 2010b). Two-Advantages of this technique the real-time PCR, which 199 uses a single closed tube for amplification and analysis, are reduced opportunity for amplicon contamination and a 200 semi-quantitative assay result. Equipment and reagents needed for real-time PCR is are expensive, requires 201 preventive maintenance, and may be beyond the capabilities of some laboratories. Real-time PCR assays may target 202 one of several genes (Carelli et al., 2007; Decaro et al., 2008), or 16S rRNA (Reinbold et al., 2010b), and are reported 203 to achieve a level of analytical sensitivity equivalent to nested conventional PCR (Carelli et al., 2007; Decaro et al., 204 2008: Reinbold et al., 2010b). 205

- 206The most widely cited assays for the detection A. marginale in individual animals use a probe for increased specificity207and are designed to detect msp1b (Carelli et al., 2007) or msp5 (Futse et al., 2003) in genomic DNA extracted from208whole blood. The assay based on detection of msp1b has been partially validated to detect the pathogen in individual209animals and was used to define samples for the validation of a C-ELISA (Carelli et al., 2007; Chung et al., 2014). The210analytical test performance of this assay is robust, and exclusivity testing confirmed other bacterial and protozoal tick-211borne pathogens of cattle were not detected. The assay, evaluated using 51 blood samples from 18 cattle herds in212three regions of southern Italy, had 100% concordance with nested PCR.
- Msp1b is a multigene family. Based on the annotation of the St. Maries strain of A. marginale, the designed primers
 and probe will amplify multiple members of this gene family, including msp1b-1, msp1b-2, and msp1-pg3). This may
 help increase diagnostic sensitivity, but may pose challenges if quantification of the pathogen is desired. Additionally,
 some A. marginale strains have single nucleotide polymorphisms in msp1b within the primer and probe binding
 regions. Thus, if msp1b is used as a diagnostic target, primer and probe design should consider local A. marginale
 strains. Msp1b has the advantage as a target in that orthologs of this gene family are absent in the related A.
 phagocytophilum and Ehrlichia spp., including E. ruminantium, thus helping ensure specificity of the test.
- 220Msp5 has also been used as a target to detect A. marginale in cattle in field samples and more frequently in221experimental samples (Futse et al., 2003). Msp5 is highly conserved among A. marginale strains and is a single copy222gene, thus providing some advantages as a target for ensuring detection of widely variant strains of A. marginale.223However, the related Anaplasma spp. and Ehrlichia spp. all have msp5 orthologs with 50% identity to an E.224ruminantium gene (NCBI accession: L07385.1), thus specificity must be determined in laboratory and field samples.225Additionally, little work has been done to validate an msp5-based real-time PCR test for diagnostic purposes.
- 226A third primer—probe set is designed to detect A. marginale using real-time, reverse transcriptase PCR. The primers227amplify a 16sRNA gene segment from A. marginale and A. phagocytophilum, while the probe differentiates between228the two species (Reinbold et al., 2010b). The analytical performance of this assay is robust. However, the diagnostic229sensitivity, specificity, and of particular importance with 16sRNA sequence-based tests, exclusivity for other tick-230borne pathogens of cattle have not been evaluated. Additionally, this assay is designed for use following RNA231extraction and reverse transcription, which is more laborious and expensive than DNA extraction. Bacterial RNA is232rapidly degraded, and this may ultimately reduce diagnostic sensitivity of this assay.

233 In regions that use A. centrale as a vaccine, it may be useful to differentiate between A. marginale and A. centrale infected/vaccinated animals. PCR is best suited for this task. The real-time PCR assay developed by Carelli et al. 234 can also be used in a duplex reaction to detect and differentiate between A. centrale and A. marginale (Decaro et al., 235 236 2008). Primers and probe have been designed to specifically amplify a region of A. centrale groEL, but not A. marginale groEL, despite 97% sequence identity between the two genes. The A. marginale-specific primers and 237 probes perform similarly in the single and duplex PCR (Carelli et al., 2007). Using the same 51 field samples from 238 239 cattle in Italy, the A. centrale assay had less analytical sensitivity compared with nested PCR and discordance in 4 240 of 51 samples between an A. centrale reverse line blot test and the duplex PCR assay.

		-			
<u>Assay</u>	<u>Reference</u>	<u>Oligonucleotides^(a)</u>	Sequence 5'–3' ^(b)	<u>Amplicon</u> <u>size (bp)</u>	<u>NCBI</u> <u>accession</u> <u>number</u>
		<u>Am_msp1b_F</u>	TTG-GCA-AGG-CAG-CAG-CTT		
Real-time PCR	<u>Carelli <i>et al.,</i></u>	<u>Am_msp1b_R</u>	TTC-CGC-GAG-CAT-GTG-CAT	95	<u>M59845</u>
	<u>2007</u>	<u>Am_msp1b_PB</u>	TCG-GTC-TAA-CAT-CTC-CAG-GCT-TTC- AT	<u> 20</u>	
	<u>Futse <i>et al.,</i></u> 2003	<u>Am_msp5_F</u>	GCC-AAG-TGA-TGG-TGA-TAT-CGA	<u>151</u>	<u>M93392</u>
Real-time PCR		<u>Am_msp5_R</u>	AGA-ATT-AAG-CAT-GTG-ACC-GCT-G		
		<u>Am_msp5_PB</u>	AAC-GTT-CAT-GTA-CCT-CAT-CAA		
Reverse-	<u>Reinbold et</u> <u>al., 2010</u>	<u>16S rRNA_F^(c)</u>	CTC-AGA-ACG-AAC-GCT-GG		
transcription		<u>16S rRNA_R(c)</u>	CAT-TTC-TAG-TGG-CTA-TCC-C	<u>142</u>	<u>M60313</u>
<u>real-time PCR</u>		Am_16S rRNA_PB ^(d)	CGC-AGC-TTG-CTG-CGT-GTA-TGG-T		
		<u>Ac_groEL_F^(e, f)</u>	CTA-TAC-ACG-CTT-GCA-TCT-C	<u>77</u>	<u>CP001759.1</u>
Real-time PCR ^(d)	<u>Decaro et</u> <u>al., 2008</u>	<u>Ac_groEL_R^(e, f)</u>	CGC-TTT-ATG-ATG-TTG-ATG-C		
		Ac_groEL_PB ^(e, f)	TCA-TCA-TTC-TTC-CCC-TTT-ACC-TCG-T		



Table 1. Oligonucleotides used in PCR assays to detect A. marginale and A. centrale

242

243

244

245 246 247 (b)Fluorophores and quenchers not included in probe sequences.
(c)Amplifies A. phagocytophilum and A. marginale 16S rRNA gene.

^(d)Probe is specific for *A. marginale* 16S rRNA gene.

^(e)Can be used as a duplex PCR with *msp1b* primers and probe based on Carelli *et al.*, 2007.

^(a)Am denotes A. marginale, Ac denotes A. centrale, Pb denotes probe sequence.

^(f)Primers and probe amplify A. centrale groEL.

248 2. Serological tests

In general, unless animals have been treated or are at a very early stage of infection (<14 days), serology using the competitive enzyme-linked immunosorbent assay (C-ELISA), indirect ELISA (I-ELISA) or card agglutination test (CAT) (see below) may be the preferred methods of identifying infected animals in most laboratories. *Anaplasma <u>marginale</u>* infections usually persist for the life of the animal. However, except for occasional small recrudescences, *Anaplasma <u>Marginale</u> A. marginale initial bodies* cannot readily be detected in blood smears after acute rickettsaemia and, even-end-point PCR may not detect the presence of *Anaplasma* the pathogen in blood samples from asymptomatic carriers. Thus, a number of serological tests have been developed with the aim of detecting persistently infected animals.

A feature of the serological diagnosis of anaplasmosis is the highly variable results with regard to both sensitivity and 256 257 specificity reported for many of the tests from different laboratories. This is due at least in part to inadequate evaluation validation of the tests using significant numbers of known positive and negative animals. Importantly, the capacity of several 258 assays to detect known infections of long standing duration has been inadequately addressed. An exception is a C-ELISA 259 (see below), which has been was initially validated using true positive and negative animals defined by nested PCR (Torioni 260 De Echaide et al., 1998), and the card agglutination assay, for which relative sensitivity and specificity in comparison with 261 the C-ELISA has been evaluated (Molloy et al., 1999). And updated in 2014 (Chung et al., 2014). Therefore, while most of 262 the tests described in this section are useful for obtaining broad-based epidemiological data, caution is advised on their 263 use for disease certification. The C-ELISA, I-ELISA and CAT are described in detail below. 264

It should be noted that there is a high degree of cross-reactivity between *A. marginale* and *A. centrale*, as well as cross-reactivity with both *A. phagocytophilum* and *Ehrlichia* spp. in serological tests (Al-Adhami *et al.*, 2011; Dreher *et al.*, 2005).
While the infecting species can sometimes be identified using antigens from homologous and heterologous species, equivocal results are obtained on many occasions. Efforts have been made to develop tests that differentiate between naturally acquired immunity to *A. marginale* and vaccine acquired immunity due to immunisation with *A. centrale* (Bellezze *et al.*, 2023; Sarli *et al.*, 2020).

271 **2.1.** Competitive enzyme-linked immunosorbent assay

272A C-ELISA using a recombinant antigen termed Major surface protein 5 (MSP5) is an immunodominant protein273expressed by A. marginale, A. ovis, and A. centrale. In A. marginale the gene is highly conserved making it a useful274target across broad geographical regions with high A. marginale strain diversity (Knowles et al., 1996; Torioni De

Echaide et al., 1998). Thus, a C-ELISA based on recombinantly expressed (rMSP5-and MSP5-) in combination with 275 276 an MSP5-specific monoclonal antibody (mAb) has proven very sensitive and specific for detection of Anaplasmainfected animals (Hofmann Lehmann et al., 2004-Molloy et al., 1999; Reinbold et al., 2010b; Strik et al., 2007). All 277 278 A. marginale strains tested, along with Additionally, A. ovis and A. centrale, express the MSP5 antigen and induce infected animals produce antibodies against the immunodominant epitope recognised by the MSP5-specific mAb. A 279 recent report mAb used in the C-ELISA. This C-ELISA was updated in 2014 to improve performance by using 280 glutathione S-transferase (GST) instead of maltose binding protein (MBP) as the tag on the rMSP5 (Chung et al., 281 2014). This assay no longer requires adsorption to remove the antibodies directed against MBP, thus it is faster and 282 283 easier than the previous version of the C-ELISA. The diagnostic sensitivity is 100% and the diagnostic specificity is 284 99.7% using a cut-off of 30% inhibition as determined by receiver operating characteristic (ROC) plot (Chung et al., 2014). For this validation, 385 sera defined as negative were from dairy cattle maintained in tick-free facilities from 285 farms with no clinical history of bovine anaplasmosis. The 135 positive sera were from cattle positive for A. marginale 286 using nested PCR and serology. 287

288 One study suggested that antibodies from cattle experimentally infected with A. phagocytophilum will test positive in the C-ELISA (Dreher et al., 2005). However, in another study no cross-reactivity could be demonstrated, and the mAb 289 used in the assay did not react with A. phagocytophilum MSP5 in direct binding assays (Strik et al., 2007). Cross 290 291 reactivity has been demonstrated between A. marginale and Ehrlichia spp, in naturally and experimentally infected cattle (Al Adhami et al, 2011). Earlier studies had shown that the C ELISA was 100% specific using 261 known 292 negative sera from a non-endemic region, detecting acutely infected cattle as early as 16 days after experimental tick 293 294 or blood inoculation, and was demonstrated to detect cattle that have been experimentally infected as long as 6 years previously (Knowles et al., 1996). In detecting persistently infected cattle from an anaplasmosis-endemic region that 295 were defined as true positive or negative using a nested PCR procedure, the rMSP5 C-ELISA had a sensitivity of 296 96% and a specificity of 95% (Torioni De Echaide et al., 1998) A. marginale and Ehrlichia sp. BOV2010 isolated in 297 298 Canada, in naturally and experimentally infected cattle (Al-Adhami et al, 2011).

Test results using the rMSP5 C-ELISA are available in less than 2.5-hours. A test kit <u>is</u> available commercially <u>that</u> contains specific instructions. <u>Users should follow the manufacturer's instructions</u>. In general, however, it is conducted as follows.

302	<u>2.1.1.</u>	Kit reagents
303		A 96-well microtitre plate coated with rMSP5 antigen,
304		A 96-well coated adsorption/transfer plate for serum adsorption to reduce background binding,
305		100×Mab_peroxidase conjugate,
306		10× wash solution and ready-to-use conjugate-diluting buffer,
307		Ready to use substrate and stop solutions,
308		Positive and negative controls
309	2.1.2.	Test procedure
310 311		i) Add 70 µl of undiluted serum sample to the coated adsorption/transfer plate and incubate at room temperature for 30 minutes.
312 313		ii) Transfer 50 µl per well of the adsorbed <u>undiluted</u> serum to the rMSP5-coated plate and incubate at room temperature for 60 minutes.
314		iii) Discard the serum and wash the plate twice using diluted wash solution.
315 316		iv) Add 50 µl per well of the 1× diluted MAb_peroxidase conjugate to the rMSP5-coated plate wells, and incubate at room temperature for 20 minutes.
317 318		 Discard the 1×diluted MAb_peroxidase conjugate and wash the plate four times using diluted wash solution.
319 320		vi) Add 50 µl per well of the substrate solution, cover the plate with foil, and incubate for 20 minutes at room temperature.
321 322		vii) Add 50 μl per well of stop solution to the substrate solution already in the wells and gently tap the sides of the plate to mix the wells.
323		viii) Immediately read the plate in the plate reader at 620 <u>, 630 or 650</u> nm.

324	2.1.3. Test validation					
325	The mean average optical density (OD) of the negative control must range from 0.40 to 2.10. The average per					
326	cent inhibition of the positive control must be ≥30%.					
020						
327	2.1.4. Interpretation of the results					
328	The % inhibition is calculated as follows:					
	Sample OD × 100					
	100 - Mean negative control OD = Per cent inhibition					
	Weah negative control OD					
329	<u>% inhibition = 100[1 – (Sample OD ÷ Negative Control OD)]</u>					
330	Samples with <30% inhibition are negative. Samples with ≥30% inhibition are positive.					
331	Specificity of the MSP5 C-ELISA may be increased by using a higher percentage inhibition cut-off value					
332	(Bradway et al., 2001); however the effect of this change on sensitivity has not been thoroughly evaluated.					
333	Recently, an improved MSP5 C ELISA was developed by replacing rMBP MSP5 with rGST MSP5 in addition to an improvement in the antigen-coating method by using a specific catcher system. The new rMSP5-GST					
334 335	C-ELISA was faster, simpler, had a higher specificity and an improved resolution compared with the rMSP5-					
336	MBP C-ELISA with MBP adsorption (Chung et al., 2014).					
337	2.2. Indirect enzyme-linked immunosorbent assay					
338	An I-ELISA was first developed using the CAT antigen <u>, which is a crude <i>A. marginale</i> lysate</u> (see below). and it The					
339	test can be implemented where the commercial C-ELISA is not available. Unlike the C-ELISA, most reagents, such					
340	as buffers and ready-to dissolve substrates, are available commercially in many countries. Any laboratory can prepare					
341	the antigen using local strains of A. marginale, though standardised methods have not been developed. I-ELISA uses					
342	small amounts of serum and antigen <u>that</u> and the sensitivity and specificity of the test standardised with true positive					
343 344	and negative sera is as good as for the C-ELISA. As it can be prepared in each laboratory <u>.</u> Only the general procedure is described here (Barry <i>et al.</i> , 1986). For commercial kits, the manufacturer's instructions should be followed. In the					
344 345	case of in-house I-ELISA-The sensitivity and specificity of the test was 87.3% and 98.4–99.6% respectively, though					
346	this varied by laboratory (Nielsen <i>et al.</i> , 1996). For general methods, refer to Barry <i>et al.</i> (1986). Initial bodies and					
347	membranes are obtained as for the complement fixation test (Rogers et al., 1964). This antigen is treated with 0.1%					
348	sodium dodecyl sulphate for 30 minutes prior to fixing the antigen to the microtitre plate. For each laboratory, the					
349	specific amount of antigen has to <u>must</u> be adjusted <u>optimised</u> to obtain the best reading and the least expenditure.					
250	Alternatively, rMSP5 can be used as the antigen in this test. This eliminates the need for preparation and					
350 351	standardisation of antigen derived from splenectomised, A. marginale infected animals (Silva et al., 2006). In a					
352	comparison between I-ELISA using the CAT antigen and rMSP with a histidine tag (rMSP5-HIS), these two I-ELISAs					
353	performed identically. In this comparison, IFAT was used as the gold standard test (Silva et al., 2006).					
354	Test results using the I-ELISA are available in about 4 to 5 hours. It is generally conducted as follows:					
355	2.2.1. Test reagents					
356	A 96-well microtitre plate coated with crude A. marginale antigen,					
357	PBS/Tween buffer, (PBS 0.1 M, pH 7.2, Tween 20 0.05%),					
358	Blocking reagent (e.g. commercial dried skim milk)					
250	Trip buffer 0.1 M. MgCL, 0.1 M. NoCL 005 M, pH 0.8					

- 359 Tris buffer 0.1 M, MgCl₂, 0.1 M, NaCl, 005 M, pH 9.8
- 360 Substrate *p*-Nitrophenyl phosphate disodium hexahydrate
- 361 Positive and negative controls.

362	2.2.2.	est procedure (this test is run in triplicate)	
363		Plates can be prepared ahead of time and kept under airtight conditions at -20° C.	
364 365		Carefully remove the plastic packaging before using plates, being careful not to touch the bottom of them as this can distort the optical density reading.	I
366 367		Remove the lid and deposit 200 μ l PBST20 solution in each well and incubate for 5 minutes a room temperature (RT).	t
368) For one plate, dissolve 1.1 g of skim milk (blocking agent) in 22 ml of PBST20.	
369 370		Remove the plate contents and deposit in each well 200 μ l of blocking solution, put the lid or and incubate for 60 minutes at 37°C.	۱
371) Wash the plate three times for 5 minutes with PBST20.	
372) Dilute all serum samples including controls 1/100 in PBST20 solution.	
373 374		 Remove the contents of the plate and deposit 200 µl of diluted serum in each of the three wells for each dilution, starting with the positive and negative and blank controls. 	3
375) Incubate plate at 37°C covered for 60 minutes.	
376		Wash three times as described in subsection vi.	
377 378		Dilute 1/1000 anti-IgG bovine alkaline phosphatase conjugate in PBST20 solution. Add 200 µ of the diluted conjugate per well. Incubate the covered plate at 37°C for 60 minutes.	I
379 380) Remove the lid and <u>wash three times as described in point vi above</u> make three washes with PBST20.	ł
381 382		 Remove the contents of the plate and deposit 195 μl of 0.075% <i>p</i>-Nitrophenyl phosphate disodium hexahydrate in Tris buffer <u>in each well</u> and incubate for 60 minutes at 37°C. 	9
383 384		v) The reaction is quantified by a microplate reader spectrophotometer, adjusted to 405 nm wavelength. The data are expressed in optical density (OD).	ı
385	2.2.3.	ata analysis	
386		nalysis of results should take into account the following parameters.	
387		The mean value of the blank wells.	
388		The mean value of the positive wells with their respective standard deviations.	
389		The mean value of negative wells with their respective standard deviations.	
390 391) The mean value of the blank wells is subtracted from the mean of all the other samples if no automatically subtracted by the ELISA reader.	t
392 393		Control sera are titrated to give optical density values ranging from 0.90 to 1.50 for the positive and, 0.15 to 0.30 for the negative control.	;
394 395		ositive values are those above the cut-off calculated value which is the sum of the average of the egative and two times the standard deviation.	3
396 397 398		or purposes of assessing the consistency of the test operator, the error "E" must alsoo be estimated is is calculated by determining the percentage represented by the standard deviation of any agains eir mean serum.	
399 400		<u>s with all diagnostic tests, it is important to measure reproducibility. For more details see Chapter</u> 2.4 Measurement uncertainty.	<u>r</u>
401 402		<u>ment double-antigen sandwich ELISA to differentiate between <i>A. marginal</i>e and <u>le antibodies</u></u>	<u>1</u>
403 404 405	A. centrale-va	<u>vaccination with <i>A. centrale</i> is used to control bovine anaplasmosis, differentiation betweer</u> ated and <i>A. marginale</i> -infected animals may be useful. Because there is often high amino acio A. marginale and <i>A. centrale</i> surface proteins, identifying unique targets for serological assays for	b

405 identity between A. marginale and A. centrale surface proteins, identifying unique targets for serological assays for
 406 this purpose is difficult. Epitopes from MSP5 (aa28-210, without the transmembrane region) that are not shared
 407 between A. marginale and A. centrale were used to develop a displacement double-antigen sandwich ELISA

- 408 (ddasELISA) (Bellezze et al., 2023; Sarli et al., 2020). The recombinant MSP5 epitopes from A. marginale or A. 409 centrale are expressed in E. coli with a histidine tag and purified. The ELISA plates are then coated with either the 410 recombinant A. marginale MSP5 epitope, or the A. centrale MSP5 epitope and blocked. Serum is added to the wells and allowed to incubate. Following washing, a combination of biotinylated and non-biotinylated recombinant proteins 411 412 are added to improve specificity of the reaction (see below for specifics). The protein-biotin binding to the serum 413 antibody is detected with a peroxidase-streptavidin based detection system. The optical density for the A. marginale MSP5-coated well (ODAm) and the OD for the A. centrale MSP5 (ODAc) coated well for each animal is measured. If 414 415 the OD for either target is <0.2, the sample is excluded from the analysis. For the remaining samples, the ratio 416 between the OD values (ODAm/ODAc) is calculated. If the ratio is >0.38 the sample is considered positive for anti-417 A. marginale antibodies, and a ratio ≤ 0.38 is classified as vaccinated with A. centrale.
- 418For the detection of A. marginale the test has a diagnostic specificity of 98% and a diagnostic sensitivity of 98.9%.419For 702 field samples evaluated, 131 (19%) had an OD <0.2 in the ddasELISA and thus were excluded from the</td>420analysis. Of those animals, 52% were nested PCR positive for A. marginale, 23% were nested PCR positive for A.421centrale, 4.6% were nested PCR positive for A. marginale and A. centrale, 20% were nested PCR negative for both,422suggesting the ddasELISA may lack sensitivity.
- 423Of the 571 ddasELISA positive field samples, the agreement between the ddasELISA and nested PCR was 84% and424the kappa coefficient was 0.70 (95% CI: 0.635–0.754), indicating substantial agreement between tests. There was425agreement between the ddasELISA and nested PCR for 93% of the *A. marginale* ddasELISA positive samples and42686% of the *A. centrale* ddasELISA positive samples. Additionally, 36 nested PCR negative samples tested positive427for antibodies against *A. marginale* (n=28) or *A. centrale* (n=8) by ddasELISA. This test could not identify animals428with co-infections, meaning animals vaccinated with *A. centrale* that are then infected with *A. marginale*, which is not429uncommon.
- 430 <u>Test results using the ddasELISA are available in 5–6 hours. It is conducted as outlined below, see Bellezze *et al.*,
 431 <u>2023 for more details.</u>
 </u>
- 2.3.1. **Test reagents** 432 A 96-well microtitre plate coated with either A. marginale or A. centrale recombinant protein 433 i) 434 ii) PBS/Tween buffer (PBS (50mM sodium phosphate, 150 mM NaCL, pH 7.2) with 0.05% Tween-435 20) iii) Blocking reagent (PBS with 10% commercial dried skim milk) 436 437 iv) Purified recombinant A. marginale MSP5 epitopes and A. centrale epitopes 438 V) Biotinylated recombinant A. marginale MSP5 epitopes and A. centrale epitopes 439 vi) Streptavidin-horse radish peroxidase (HRP) detection system 440 Chromogenic substrate (1 mM 2,2'-Azinobis [3-ethylbenzothiazoline-6-sulfonic acid]vii) diammonium salt in0.05 M sodium citrate, pH 4.5, 0.0025% V/V H2O2 (100 µl/well). 441 442 viii) ELISA plate reader (405 nm reading) Positive and negative control sera for A. marginale and A. centrale 443 ix) 2.3.2. Test procedure 444 Plates are coated overnight. 445 i) 446 ii) Block with blocking buffer for 1 hour at room temperature and wash three times with PBS/Tween buffer. 447 448 iii) Add undiluted serum 100 ul/well and incubate for 1 hour at 25°C at 100 rpm. Wash three times with PBS/Tween buffer. 449 iv) Add 100 µl of A. marginale MSP5-biotin (1 µg/ml) plus A. centrale MSP5 (10 µg/ml) to 450 V) A. marginale test wells. Add A. centrale MSP5-biotin (1 µg/ml) plus A. marginale MSP5 451 (10 µg/ml) in PBS/Tween buffer + 10% fat-free dried milk to A. centrale test wells. 452 Incubate 1 hour at 25°C, 100 rpm and wash the plate five times with PBS/Tween buffer. 453 vi) 454 vii) To detect the bound protein-biotin complex, add streptavidin-HRP diluted in 1/500 in PBS/Tween buffer with 10% dried milk for 1 hour at 25°C, 100 rpm. 455 viii) Wash five times with PBS/Tween buffer. 456

457 ix) Add chromogenic substrate based on manufacturer's instructions. The reaction is measured by microplate reader spectrophotometer at 405 nm wavelength. The 458 X) 459 data are expressed in optical density (OD). OD_{405nm} <0.2 is considered negative. 460 xi) 461 xii) Results are expressed as the ratio between antibodies specific for A. marginale MSP5 and for A. centrale MSP5 (ODAm/ODAc). If the ratio is >0.38 the sample is considered positive for anti-462 463 A. marginale antibodies, and a ratio ≤ 0.38 is classified as vaccinated with A. centrale.

464 **2.4. Card agglutination test**

The advantages of the CAT are that it is sensitive-The sensitivity of the CAT is from 84% to 98% (Gonzalez et al., 465 1978; Molloy et al., 1999) and the specificity is 98.6% (Molloy et al., 1999). Though sometimes giving variable results, 466 the CAT can be useful under certain circumstances, as it may be undertaken either in the laboratory or in the field, 467 and it gives a result within a few minutes. Nonspecific reactions may be a problem, and subjectivity in interpreting 468 469 assay reactions can result in variability in test interpretation. In addition, the CAT antigen, which is a suspension lysate of A. marginale particles isolated from erythrocytes, can be difficult to prepare and can vary from batch to batch 470 471 and laboratory to laboratory. To obtain the antigen, splenectomised calves are infected by intravenous inoculation 472 with blood containing Anaplasma A. marginale-infected erythrocytes. When the rickettsaemia exceeds 50%, the animal is exsanguinated, the infected erythrocytes are washed, lysed, and the erythrocyte ghosts and Anaplasma 473 particles <u>A. marginale</u> are pelleted. The pellets are sonicated, washed, and then resuspended in a stain solution to 474 produce the antigen suspension. 475

476 A test procedure that has been slightly modified from that originally described (Amerault & Roby, 1968; Amerault *et* 477 *al.*, 1972) is as follows, and is based on controlled conditions in a laboratory setting:

	,	
478	2.4.1.	Test procedure
479 480		 Ensure all test components are at a temperature of 25–26°C before use (this constant temperature is critical for the test).
481 482 483 484		 ii) On each circle of the test card (a clear perspex/plastic or glass plate marked with circles that are 18 mm in diameter), place next to each other, but not touching, 10 μl of bovine serum factor (BSF), 10 μl of test serum, and 5 μl of CAT antigen³⁹. Negative and low positive control sera must be tested on each card.
485 486 487 488		iii) BSF is serum from a selected animal with high known conglutinin level. If the conglutinin level is unknown, fresh serum from a healthy animal known to be free from <i>Anaplasma</i> can be used. The BSF must be stored at -70°C in small aliquots, a fresh aliquot being used each time the tests are performed. The inclusion of BSF improves the sensitivity of the test.
489 490		iv) Mix well with a glass stirrer. After mixing each test, wipe the stirrer with clean tissue to prevent cross-contamination.
491		v) Place the test card in a humid chamber and rock at 100–110 rpm for 7 minutes.
492 493 494		vi) Read immediately against a backlight. Characteristic clumping of the antigen (graded from +1 to +3) is considered to be a positive result. The test is considered to give a negative result when there is no characteristic clumping.
495 496 497 498		<u>A latex card agglutination test, a relatively simple and rapid test platform, has been partially validated.</u> <u>This test uses rMSP5-HIS rather than <i>A. marginale</i> lysate and does not require BSF. The performance of this test was compared with that of the I-ELISA using rMSP5-HIS as the antigen. The relative sensitivity was 95.2% and relative specificity was 91.86% (Ramos <i>et al.</i>, 2014).</u>

³⁹ The test as conducted in the USA and Mexico uses larger volumes of reagents: antigen (15 µl), serum (30 µl), and bovine serum factor (30 µl), and a 4-minute reaction time (see step iv).

499 2.4. Complement fixation test

500 The complement fixation (CF) test has been used extensively for many years; however, it shows variable sensitivity 501 (ranging from 20 to 60%), possibly reflecting differences in techniques for antigen production, and poor reproducibility. 502 In addition, it has been demonstrated that the CF assay fails to detect a significant proportion of carrier cattle (Bradway 503 *et al.*, 2001). It is also uncertain as to whether or not the CF test can identify antibodies in acutely infected animals 504 prior to other assays (Coetzee *et al.*, 2007; Molloy *et al.*, 1999). Therefore, the CF test is no longer recommended as 505 a reliable assay for detecting infected animals.

2.5. Indirect fluorescent antibody test

Because of the limitations on the number of indirect fluorescent antibody (IFA) tests that can be performed daily by 507 508 one operator, other serological tests are generally preferred to the IFA test. The IFA test is performed as described for bovine babesiosis in chapter 3.4.2, except that A. marginale infected blood is used for the preparation of antigen 509 smears. A serious problem encountered with the test is nonspecific fluorescence. The reported sensitivity is 97.6% 510 and specificity 89.6% (Gonzalez et al., 1978). Antigen made from blood collected as soon as adequate rickettsaemia 511 (5-10%) occurs is most likely to be suitable. Nonspecific fluorescence due to antibodies adhering to infected 512 erythrocytes can be reduced by washing the erythrocytes in an acidic glycine buffer before antigen smears are 513 prepared. Infected erythrocytes are washed twice in 0.1 M glycine buffer (pH 3.0, centrifuged at 1000 g for 15 minutes 514 at 4°C) and then once in PBS, pH 7.4. Recently published data show that the IFA, like the C-ELISA, can cross react 515 with other members of the Anaplasmataceae family, and specifically an Ehrlichia spp. identified as BOV2010 (Al-516 517 Adhami et al., 2011).

518 **2.6. Complement fixation test**

519The complement fixation test (CFT) was used extensively for many years; however, it has variable sensitivity (ranging520from 20 to 60%), possibly reflecting differences in techniques for antigen production, and poor reproducibility. In521addition, the CF assay fails to detect a significant proportion of carrier cattle (Bradway *et al.*, 2001). It is also uncertain522as to whether or not the CF test can identify antibodies in acutely infected animals prior to other assays (Coetzee *et al.*, 2007; Molloy *et al.*, 1999). Therefore, the CF test is no longer recommended as a reliable assay for detecting524infected animals.

525

C. REQUIREMENTS FOR VACCINES

526 1. Background

527 Several immunisation methods have been used to protect cattle against anaplasmosis in countries where the disease is 528 endemic, but none is ideal to date (McHardy, 1984). A review of *A. marginale* vaccines and antigens has been published 529 (Kocan *et al.*, 2003-2010; Noh *et al.*, 2012). Use of the less pathogenic *A. centrale*, which gives partial cross-protection 530 against *A. marginale*, is the most widely accepted method, although not used in many countries where the disease is 531 exotic, including north America.

In this section, the production of live *A. centrale* vaccine is described. It involves infection of a susceptible, splenectomised
calf and the use of its blood as a vaccine. Detailed accounts of the production procedure are available and reference should
be made to these publications for details of the procedures outlined here (Bock *et al.*, 2004; de Vos & Jorgensen, 1992;
Pipano, 1995).

Guidelines for the production of veterinary vaccines are given in Chapter 1.1.8 *Principles of veterinary vaccine production*.
 The guidelines given here and in Chapter 1.1.8 are intended to be general in nature and may be supplemented by national and regional requirements.

539 *Anaplasma centrale* vaccine can be provided in-either frozen or chilled-form depending on demand, transport networks, 540 and the availability of liquid nitrogen or dry ice supplies. Frozen vaccine is recommended in most instances, as it allows 541 for thorough post-production quality control of each batch. It is, however, more costly to produce and more difficult to 542 transport than chilled vaccine. The risk of contamination makes post-production control essential, but may be prohibitively 543 expensive.

544 2. Outline of production and minimum requirements for conventional vaccines

545 2.1. Characteristics of the seed

2.1.1. Biological characteristics

Anaplasma centrale was isolated in 1911 in South Africa and has been used as a vaccine in South America, Australia, Africa, the Middle East, and South-East Asia. It affords only partial, but adequate, protection in regions where the challenging circulating strains are of moderate virulence (e.g. Australia) (Bock & de Vos, 2001). In the humid tropics where *A. marginale* appears to may be a very more virulent-rickettsia, the protection afforded by *A. centrale* may be inadequate to prevent disease in some animals.

553Anaplasma centrale usually causes benign infections, especially if used in calves under 9 months of554age. Severe reactions following vaccination have been reported when adult cattle are inoculated. The555suitability of an isolate of *A. centrale* as a vaccine can be determined by inoculating susceptible cattle,556monitoring the subsequent reactions, and then challenging the animals and susceptible controls with557a virulent local strain of *A. marginale*. Both safety and efficacy can be judged by monitoring558rickettsaemias in stained blood films and the depression of packed cell volumes of inoculated cattle559during the vaccination and challenge reaction periods.

Infective material for preparing the vaccine is readily stored as frozen stabilates of infected blood in liquid nitrogen or dry ice. Dimethyl sulphoxide (DMSO) and or polyvinylpyrrolidone M.W. 40,000 (Bock *et al.*, 2004) are the recommended cryopreservatives, as they allow for intravenous administration after thawing of the stabilate. A detailed account of the freezing technique using DMSO is reported elsewhere (Mellors *et al.*, 1982), but briefly involves the following: infected blood is collected, chilled to 4°C, and cold cryoprotectant (4 M DMSO in PBS) is added slowly with stirring to a final blood:protectant ratio of 1:1, to give a final concentration of 2 M DMSO. The entire dilution procedure is carried out in an ice bath and the diluted blood is dispensed into suitable containers (e.g. 5 ml cryovials), and frozen, as soon as possible, in the vapour phase of a liquid nitrogen container.

2.1.2. Quality criteria

Evidence of purity of the *A. centrale* isolate can be determined by serological testing of paired sera from the cattle used in the safety test for possible <u>contaminants pathogens</u> that may be present (Bock *et al.*, 2004; Pipano, 1997). Donor calves used to expand the seed for vaccine production should be examined for all blood-borne infections prevalent in the vaccine-producing country, including *Babesia, Anaplasma, Ehrlichia, Theileria* and *Trypanosoma*. This can be done by routine examination of stained blood films after splenectomy, <u>PCR</u>, and preferably also by serology. Any calves showing evidence of natural infections of any of these agents should be rejected. The absence of other infective agents should also be confirmed. These may include the agents of enzootic bovine leukosis, mucosal disease, infectious bovine rhinotracheitis, ephemeral fever, Akabane disease, bluetongue, <u>and</u> foot and mouth disease, and rinderpest. The testing procedures will depend on the diseases prevalent in the country and the availability of tests but should involve serology of paired sera at the very least and, in some cases, virus isolation, antigen, or DNA/RNA detection (Bock *et al.*, 2004; Pipano, 1981; 1997).

584 2.2. Method of manufacture

2.2.1. Procedure

i) Production of frozen vaccine

Quantities of the frozen stabilate (5–10 ml) are thawed by immersing the vials in water preheated to 40°C. The thawed material is kept on ice and used as soon as possible (within 30 minutes) to infect a susceptible, splenectomised calf by intravenous inoculation.

The rickettsaemia of the this donor calf is monitored daily by examining stained films of jugular blood, and the blood is collected for vaccine production when suitable rickettsaemias are reached. A rickettsaemia of 1×10^8 /ml (approximately 2% rickettsaemia in jugular blood) is the minimum required for production of vaccine as this is the dose to vaccinate a bovine. If a suitable rickettsaemia is not obtained, passage of the strain by subinoculation of 100–200 ml of blood to a second splenectomised calf may be necessary.

596	Blood from the donor is collected by aseptic jugular or carotid cannulation using heparin as an
	anticoagulant (5 International Units [IU] heparin/ml blood). The use of blood collection units for human
597	
598	use are also suitable and guarantee sterility and obviate the need to prepare glass flasks that make
599	the procedure more cumbersome.
600	In the laboratory, the infective blood is mixed in equal volumes with 3 M glycerol in PBS supplemented
601	with 5 mM glucose at 37°C (final concentration of glycerol 1.5 M). The mixture is then equilibrated at
602	37°C for 30 minutes and dispensed into suitable containers (e.g. 5 ml cryovials). The vials are cooled
603	at approximately 10°C/minute in the vapour phase of liquid nitrogen and, when frozen, stored in the
604	liquid phase (Bock <i>et al.,</i> 2004).
004	
005	DMOO and he could be a compared at the start in the science of starting in the course of startin
605	DMSO can be used as a cryoprotectant in the place of glycerol. This is done in the same way as
606	outlined for the preparation of seed stabilate (Mellors <i>et al.,</i> 1982; Pipano, 1981).
607	If glycerolised vaccine is to be diluted, the diluent should consist of PBS with 1.5 M glycerol and 5 mM
608	glucose (Jorgensen et al., 1989). Vaccine cryopreserved with DMSO should be diluted with diluent
609	containing the same concentration of DMSO as in the original cryopreserved blood (Pipano et al.,
610	1986).
611	ii) Production of chilled vaccine
612	Infective material for chilled vaccine is prepared in the same way as for frozen vaccine, but it must
613	be issued and used as soon as possible after collection. The infective blood can be diluted to provide
614	1×10^7 parasites per dose of vaccine. A suitable diluent is 10% sterile bovine serum in a
615	glucose/balanced salt solution containing the following quantities per litre: NaCI (7.00 g), MgCI ₂ .6H ₂ O
616	(0.34 g), glucose (1.00 g), Na₂HPO₄(2.52 g), KH₂PO₄(0.90 g), and NaHCO₃(0.52 g).
617	If diluent is not available, acid citrate dextrose (20% [v/v]) or citrate phosphate dextrose (20% [v/v])
618	should be used as anticoagulant to provide the glucose necessary for survival of the organisms.
010	
010	
619	iii) Use of vaccine
620	In the case of frozen vaccine, vials should be thawed by immersion in water, preheated to 37°C to
621	40°C, and the contents mixed with suitable diluent to the required dilution. If glycerolised vaccine is
622	prepared, it should be kept cool and used within 8 hours (Bock <i>et al.</i> , 2004). If DMSO is used as a
623	cryoprotectant, the prepared vaccine should be kept on ice and used within 15–30 minutes (Pipano,
624	1981). The vaccine is most commonly administered subcutaneously.
024	1901). The vaccine is most commonly auministered subcutaneously.
625	iv) Chilled vaccine should be kept refrigerated and used within 4–7 days of preparation.
025	V of line vaccine should be kept reingerated and used within $4-7$ days of preparation.
626	The strain of A. centrale used in the vaccine is of reduced virulence, but is not entirely safe. A practical
627	recommendation is, therefore, to limit the use of vaccine to calves, where nonspecific immunity will
628	minimise the risk of vaccine reactions. When older animals have to be vaccinated, there is a risk of
629	severe reactions. These reactions occur infrequently, but valuable breeding stock or pregnant
630	animals obviously warrant close attention, -and should be observed daily for 3 weeks post-
631	vaccination. Clinically sick animals should be treated with oxytetracycline or imidocarb at dosages
632	recommended by the manufacturers. Protective immunity develops in 6–8 weeks and usually lasts
633	for several years.
004	
634	Anaplasmosis and babesiosis vaccines are often used concurrently, but it is not advisable to use any
635	other vaccines at the same time (Bock <i>et al.,</i> 2004).
636	2.2.2. Requirements for substrates and media
637	Anaplasma centrale cannot can be cultured in vitro Rhipicephalus appendiculatus and Dermacentor variabilis
638	cells lines, though antigen expression and immunogenicity of the cultured A. centrale need to be tested (Bell-
639	<u>Sakyi et al., 2015</u>). No substrates or media other than buffers and diluents are used in vaccine production.
640	DMSO or glycerol should be purchased from reputable companies.
0-10	Emere el giyoriol enouid de puloridoca nom reputable companies.
641	2.2.3. In-process controls
641	2.2.3. In-process controls
642	i) Source and maintenance of vaccine donors
640	A course of column from natural infactions of Anonlasma A marginals and other tick haves
643	A source of calves free from natural infections of <i>Anaplasma <u>A</u>. marginale</i> and other tick-borne
644	diseases should be identified. If a suitable source is not available, it may be necessary to breed the
645	calves under tick-free conditions specifically for the purpose of vaccine production.

646 647 648 649 650	The calves should be maintained under conditions that will prevent exposure to infectious diseases and to ticks and biting insects. In the absence of suitable facilities, the risk of contamination with the agents of infectious diseases present in the country involved should be estimated, and the benefits of local production of vaccine weighed against the possible adverse consequences of spreading disease (Bock <i>et al.</i> , 2004).
651	ii) Surgery
652 653	Donor calves should be splenectomised to allow maximum yield of organisms for production of vaccine. This is best carried out in young calves and under general anaesthesia.
654	iii) Screening of vaccine donors before inoculation
655 656 657 658 659 660 661 662 663 664	As for preparation of seed stabilate, donor calves for vaccine production should be examined for all blood-borne infections prevalent in the vaccine-producing country, including <i>Babesia</i> , <i>Anaplasma</i> , <i>Ehrlichia</i> , <i>Theileria</i> and <i>Trypanosoma</i> . This can be done by routine examination of stained blood films after splenectomy, and preferably also by serology. Any calves showing evidence of natural infections of any of these agents should be rejected. The absence of other infective agents should also be confirmed. These may include the agents of enzootic bovine leukosis, bovine viral diarrhoea, infectious bovine rhinotracheitis, ephemeral fever, Akabane disease, bluetongue, and foot and mouth disease. The testing procedures will depend on the diseases prevalent in the country and the availability of tests, but should involve serology of paired sera at the very least and, in some cases, virus isolation, antigen, or DNA/RNA detection (Bock <i>et al.</i> , 2004; Pipano, 1981; 1997).
665	iv) Monitoring of rickettsaemias following inoculation
666 667 668	It is necessary to determine the concentration of rickettsia in blood being collected for vaccine. The rickettsial concentration can be estimated from the erythrocyte count and the rickettsaemia (percentage of infected erythrocytes).
669	v) Collection of blood for vaccine
670 671 672	All equipment should be sterilised before use (e.g. by autoclaving). Once the required rickettsaemia is reached, the blood is collected in heparin using strict aseptic techniques. This is best done if the calf is sedated and with the use of a closed-circuit collection system.
673 674 675	Up to 3 litres of heavily infected blood can be collected from a 6-month-old calf. If the calf is to live, the transfusion of a similar amount of blood from a suitable donor is indicated. Alternatively, the calf should be killed immediately after collection of the blood.
676	vi) Dispensing of vaccine
677 678 679 680	All procedures are performed in a suitable environment, such as a laminar flow cabinet, using standard sterile techniques. Use of a mechanical or magnetic stirrer will ensure thorough mixing of blood and diluent throughout the dispensing process. Penicillin (500,000 IU/litre) and streptomycin (370,000 μ g/litre) are added to the vaccine at the time of dispensing.
681	2.2.4. Final product batch tests
682 683 684	The potency, safety and sterility of vaccine batches cannot be determined in the case of chilled vaccine, and specifications for frozen vaccine depend on the country involved. The following are the specifications for frozen vaccine produced in Australia.
685	i) Sterility and purity
686 687	Standard tests for sterility are employed for each batch of vaccine and diluent (see Chapter 1.1.9 <i>Tests for sterility and freedom from contamination of biological materials intended for veterinary use</i>).
688 689 690 691 692 693 694 695 696	The absence of contaminants is determined by doing appropriate serological testing of donor cattle, by inoculating donor lymphocytes into sheep and then monitoring them for evidence of viral infection, and by inoculating cattle and monitoring them serologically for infectious agents that could potentially contaminate the vaccine. Cattle inoculated during the test for potency (see Section C.2.2.4.iii) are suitable for the purpose. Depending on the country of origin of the vaccine, these agents include the causative organisms of enzootic bovine leukosis, infectious bovine rhinotracheitis, bovine viral diarrhoea, ephemeral fever, Akabane disease, Aino virus, bluetongue, parainfluenza, foot and mouth disease, lumpy skin disease, rabies, Rift Valley fever, contagious bovine pleuropneumonia, Jembrana disease, heartwater, pathogenic <i>Theileria</i> and <i>Trypanosoma spp.</i> , <i>Brucella abortus</i> ,

700		there are many publications describing primers, and assay conditions for any particular disease.	
701		ii) Safety	
702		Vaccine reactions of the cattle inoculated in the test for potency (see Chapter 1.1.8 Principles of	
703		veterinary vaccine production) are monitored by measuring rickettsaemia and depression of packed	
704		cell volume. Only batches with pathogenicity levels equal to or lower than a predetermined standard	
705		are released for use.	
706		iii) Potency	
707		Vaccine is thawed and diluted 1/5 with a suitable diluent (Bock et al., 2004). The diluted vaccine is	
708		then incubated for 8 hours at 4°C, and five cattle are inoculated subcutaneously with 2 ml doses. The	
709		inoculated cattle are monitored for the presence of infections by examination of stained blood smears.	
710		All should become infected for a batch to be accepted. A batch proving to be infective is	
711		recommended for use at a dilution of 1/5 with isotonic diluent.	
712	2.3. Requir	rements for authorisation	
713	2.3.1.	Safety	
		-	
714		The strain of A. centrale used in vaccine is of reduced virulence but is not entirely safe. A practical	
715		recommendation is, therefore, to limit the use of vaccine to calves, where nonspecific immunity will	
716		minimise the risk of vaccine reactions. When older animals have to be vaccinated, there is a risk of	
717		severe reactions. These reactions occur infrequently, but valuable breeding stock or pregnant	
718		animals obviously warrant close attention, and should be observed daily for 3 weeks post-	
719		vaccination. Clinically sick animals should be treated with oxytetracycline or imidocarb at dosages	
720		recommended by the manufacturers.	
120			
721		Anaplasma centrale is not infective to other species, and the vaccine is not considered to have other	
722		adverse environmental effects. The vaccine is not infective for humans. When the product is stored	
723		in liquid nitrogen, the usual precautions pertaining to the storage, transportation and handling of deep-	
724		frozen material applies.	
124			
725	2.3.2.	Efficacy requirements	
726		Partial but long-lasting immunity results from one inoculation. There is no evidence that repeated	
727		vaccination will have a boosting effect. Immunisation with live A. centrale results in long-term infection	
728		of the vaccinee, thus repeated vaccination is unnecessary. Infection with A. centrale does not prevent	
729		subsequent infection with A. marginale, but does at least result in protection from disease (Shkap et	
730		<u>al., 2009).</u> The vaccine is used for control of clinical anaplasmosis in endemic areas. It will not provide	
731		sterile immunity, and should not be used for eradication of <i>A. marginale</i> .	
732	2.3.3.	Stability	
733		The vaccine can be kept for 5 years when stored in liquid nitrogen. Once thawed, it rapidly loses its	
734		potency. Thawed vaccine cannot be refrozen.	
735	3. Vaccines ba	ased on biotechnology	
736	There are no vaccin	es based on biotechnology available for anaplasmosis.	
737		REFERENCES	
738	AL-ADHAMI B., SCANDRETT W.B., LOVANOV V.A. & GAJADHAR A.A. (2011). Serological cross reactivity between Anaplasma		
739	marginale and Ehrlichia species in naturally and experimentally infected cattle. J. Vet. Diagn. Invest., 23, 1181–1188.		
740 741	AMERAULT T.E. & ROBY T.O. (1968). A rapid card agglutination test for bovine anaplasmosis. <i>J. Am. Vet. Med. Assoc.</i> , 153 , 1828–1834.		

Coxiella, and Leptospira (Bock et al., 2004; Pipano, 1981; 1997). Other pathogens to consider include

the causal agents of bovine tuberculosis and brucellosis as they may spread through contaminated

blood used for vaccine production. Most of these agents can be tested by means of specific PCR and

697

698

699

- AMERAULT T.E., ROSE J.E. & ROBY T.O. (1972). Modified card agglutination test for bovine anaplasmosis: evaluation with serum and plasma from experimental and natural cases of anaplasmosis. *Proc. U.S. Anim. Health Assoc.*, **76**, 736–744.
- BARRY D.N., PARKER R.J., DE VOS A.J., DUNSTER P. & RODWELL B.J. (1986). A microplate enzyme-linked immunosorbent assay for measuring antibody to *Anaplasma marginale* in cattle serum. *Aust. Vet. J.*, **63**, 76–79.
- BELL-SAKYI L., PALOMAR A.M., BRADFORD E.L. & SHKAP V. (2015). Propagation of the Israeli vaccine strain of Anaplasma
 centrale in tick cell lines. Vet. Microbiol., **179**, 270–276.
- BELLEZZE J., THOMPSON C.S., BOSIO A.S., TORIONI S.M. & PRIMO M.E. (2023). Development and field evaluation of an ELISA
 to differentiate Anaplasma marginale-infected from A. centrale-vaccinated cattle. J. Vet. Diagn. Invest., 35, 204–208.
- 750 BOCK R., JACKSON L., DE VOS A. & JORGENSEN W. (2004). Babesiosis of cattle. Parasitology, 129, Suppl, S247–269.
- BOCK R.E. & DE VOS A.J. (2001). Immunity following use of Australian tick fever vaccine: a review of the evidence. *Aust. Vet. J.*, **79**, 832–839.
- BRADWAY D.S., TORIONI DE ECHAIDE S., KNOWLES D.P., HENNAGER S.G. & MCELWAIN T.F. (2001). Sensitivity and specificity of
 the complement fixation test for detection of cattle persistently infected with *Anaplasma marginale*. J. Vet. Diagn. Invest.,
 13, 79–81.
- CARELLI G., DECARO N., LORUSSO A., ELIA G., LORUSSO E., MARI V., CECI L. & BUONAVOGLIA C. (2007). Detection and quantification of *Anaplasma marginale* DNA in blood samples of cattle by real-time PCR. *Vet. Microbiol.*, **124**, 107–114.
- COETZEE J.F., SCHMIDT P.L., APLEY M.D., REINBOLD J.B. & KOCAN K.M. (2007). Comparison of the complement fixation test
 and competitive ELISA for serodiagnosis of *Anaplasma marginale* infection in experimentally infected steers. *Am. J. Vet. Res.*, 68, 872–878.

CHUNG C., WILSON C., BANDARANAYAKA-MUDIYANSELAGE C.-B., KANG E., ADAMS D.S., KAPPMEYER L.S., KNOWLES D.P.,
 MCELWAIN T.F., EVERMANN J.F., UETI M.W., SCOLES G.A., LEE S.S. & MCGUIRE T.C. (2014). Improved diagnostic performance
 of a commercial *Anaplasma* antibody competitive enzyme-linked immunosorbent assay using recombinant major surface
 protein 5-glutathione S-transferase fusion protein as antigen. *J. Vet. Diagn. Invest.*, 26, 61–71.

DECARO N., CARELLI G., LORUSSO E., LUCENTE M.S., GRECO G., LORUSSO A., RADOGNA A., CECI L. & BUONAVOGLIA C. (2008).
 Duplex real-time polymerase chain reaction for simultaneous detection and quantification of *Anaplasma marginale* and
 Anaplasma centrale. J. Vet. Diagn. Invest., **20**, 606–611.

DE VOS A.J. & JORGENSEN W.K. (1992). Protection of cattle against babesiosis in tropical and subtropical countries with a
 live, frozen vaccine. *In*:Tick Vector Biology, Medical and Veterinary Aspects, Fivaz B.H., PetneyT.N. & Horak I.G., eds.
 Springer Verlag, Berlin, Germany, 159–174.

DREHER U.M., DE LA FUENTE J., HOFMANN-LEHMANN R., MELI M.K., PUSTERIA N., KOCAN K.M., WOLDEHIWET A., REGULA G. &
 STAERK K.D.C. (2005). Serologic cross reactivity between *Anaplasma marginale* and *Anaplasma phagocytophilum*. *Clin. Vaccine. Immunol.*, **12**, 1177–1183.

DUMLER J.S., BARBET A.F., BEKKER C.P., DASCH G.A., PALMER G.H., RAY S.C., RIKIHISA Y. & RURANGIRWA F.R. (2001).
 Reorganization of genera in the Families Rickettsiaceae and Anaplasmataceae in the order *Rickettsiales:* unification of
 some species of *Ehrlichia* with *Anaplasma*, *Cowdria* with *Ehrlichia*, and *Ehrlichia* with *Neorickettsia*, descriptions of five
 new species combinations and designation of *Ehrlichia equi* and 'HGE agent' as subjective synonyms of *Ehrlichia phagocytophila. Int. J. Syst. Evol. Microbiol.*, **51**, 2145–2165.

GONZALEZ, E. F., LONG R. F. & TODOROVIC R. A. (1978). Comparisons of the complement-fixation, indirect fluorescent antibody, and card agglutination tests for the diagnosis of bovine anaplasmosis. *Am. J. Vet. Res.*, **39**, 1538-1541.

HOFMANN-LEHMANN R., MELI M.L., DREHER U.M., GÖNCZI E., DEPLAZES P., BRAUN U., ENGELS M., SCHÜPBACH J., JÖRGER K.,
 THOMA R., GRIOT C., STÄRKK.D.C., WILLI B., SCHMIDT J., KOCAN K.M. & LUTZ H. (2004). Concurrent infections with vector-

FUTSE J.E., UETI M.W., KNOWLES D.P. JR. & PALMER G.H. (2003). Transmission of Anaplasma marginale by Boophilus
 microplus: retention of vector competence in the absence of vector-pathogen interaction. J. Clin. Microbiol.. 41, 3829– 3834.

- borne pathogens associated with fatal haemolytic anemia in a cattle herd in Switzerland. J. Clin. Microbiol., **42**, 3775– 3780.
- JORGENSEN W.K., DE VOS A.J. & DALGLIESH R.J. (1989). Infectivity of cryopreserved *Babesia bovis, Babesia bigemina* and
 Anaplasma centrale for cattle after thawing, dilution and incubation at 30°C. *Vet. Parasitol.*, **31**, 243–251.
- KOCAN K.M., DE LA FUENTE J., BLOUIN E.F. & GARCIA GARCIA J.C. (2004). Anaplasma marginale (Rickettsiales:
 Anaplasmataceae): recent advances in defining host-pathogen adaptations of a tick-borne rickettsia. Parasitology, 129, 5285–5300.
- 793 KOCAN K.M., DE LA FUENTE J., GUGLIELMONE A.A. & MELENDÉZ R.D. (2003). Antigens and alternatives for control of 794 *Anaplasma marginale* infection in cattle. *Clin. Microbiol. Rev.*, **16**, 698–712.
- KOCAN K. M., J. DE LA FUENTE, BLOUIN E. F., COETZEE J. F. & EWING S. A. (2010). The natural history of Anaplasma marginale.
 Vet. Parasitol., **167**, 95–107.
- KNOWLES D., TORIONI DE ECHAIDE S., PALMER G., MCGUIRE T., STILLER D. & MCELWAIN T. (1996). Antibody against an
 Anaplasma marginale MSP5 epitope common to tick and erythrocyte stages identifies persistently infected cattle. *J. Clin. Microbiol.*, **34**, 2225–2230.
- KREIER J.P. & RISTIC M. (1963). Anaplasmosis. X Morphological characteristics of the parasites present in the blood of
 calves infected with the Oregon strain of *Anaplasma marginale*. *Am. J Vet. Res.*, 24, 676–687.
- 802 MCHARDY N. (1984). Immunization against anaplasmosis: a review. Prev. Vet. Med., 2, 135–146.
- MATELI.A., ESTRADA-PENA A., CUTLER S.J., VAYSSIER-TAUSSAT M., VARELA-CASTRO L., POTKONJAK A., ZELLER H. & MIHALCA
 A.D. (2019). A review on the eco-epidemiology and clinical management of human granulocytic anaplasmosis and its agent
 in Europe. Parasit. Vectors, 12, 599.
- MELLORS L.T., DALGLIESH R.J., TIMMS P., RODWELL B.J. & CALLOW L.L. (1982). Preparation and laboratory testing of a frozen vaccine containing *Babesia bovis*, *Babesi abigemina* and *Anaplasma centrale*. *Res. Vet. Sci.*, **32**, 194–197.
- MOLLOY J.B., BOWLES P.M., KNOWLES D.P., MCELWAIN T.F., BOCK R.E., KINGSTON T.G., BLIGHT G.W. & DALGLIESH R.J. (1999).
 Comparison of a competitive inhibition ELISA and the card agglutination test for detection of antibodies to *Anaplasma marginale* and *Anaplasma centrale* in cattle. *Aust. Vet. J.*, **77**, 245–249.
- NIELSEN K., SMITH P., GALL D., DE ESHAIDE S. T, WAGNER G. & DAJER A. (1996). Development and validation of an indirect
 enzyme immunoassay for detection of antibody to *Anaplasma marginale* in bovine sera. *Vet. Parasitol.*, 67, 133–142.
- NOH S. M. & BROWN W.C. (2012). Adaptive immune responses to infection and opportunities for vaccine development (Anaplasmataceae). Intracellular Pathogens II: Rickettsiales. G. H. Palmer. Washington, DC, USA, ASM Press. II: 330– 365.
- PIPANO E. (1981). Frozen vaccine against tick fevers of cattle. *In:* XI International Congress on Diseases of Cattle, Haifa,
 Israel. Mayer E., ed. Bregman Press, Haifa, Israel, 678–681.
- 818 PIPANO E. (1995). Live vaccines against hemoparasitic diseases in livestock. Vet. Parasitol., 57, 213–231.
- PIPANO E. (1997). Vaccines against hemoparasitic diseases in Israel with special reference to quality assurance. *Trop. Anim. Health Prod.*, **29** (Suppl. 4), 86S–90S.
- 821 PIPANO E., KRIGEL Y., FRANK M., MARKOVICS A. & MAYER E. (1986). Frozen *Anaplasma centrale* vaccine against 822 anaplasmosis in cattle. *Br. Vet. J.*, **142**, 553–556.
- RAMOS C.A., ARAUJO F.R., SANTOS R.C., MELO E.S., SOUSA L.C., VIDAL C.E., GUERRA N.R. & RAMOS R.A. (2014).
 Development and assessment of a latex agglutination test based on recombinant MSP5 to detect antibodies against
 Anaplasma marginale in cattle. *Braz. J. Microbiol.*, **45**, 199–204.

REINBOLD J.B., COETZEE J.F., HOLLIS L.C., NICKELL J.S., RIEGEL C.M., CHRISTOPHER J.A. & GANTA R.R. (2010a). Comparison of iatrogenic transmission of *Anaplasma marginale* in Holstein steers via needle and needle-free injection techniques. *Am. J. Vet. Res.*, **71**, 1178–1188.

REINBOLD J.B., COETZEE J.F., SIRIGIREDDY K.R. & GANTA R.R. (2010b). Detection of *Anaplasma marginale* and *A. phagocytophilum* in bovine peripheral blood samples by duplex real-time reverse transcriptase PCR assay. *J. Clin. Microbiol.*, **48**, 2424–2432.

- ROGERS T.E., HIDALCO R.-J. & DIMOPOULLOS G.T. (1964). Immunology and serology of Anaplasma marginale. I.
 Fractionation of the complement fixing antigen. J. Bacteriol., 88, 81–86.
- SARLI M., THOMPSON C.S., NOVOA M., VALENTINI B.S., MASTROPAOLO M., ECHAIDE I.E., DE ECHAIDE S. T. & PRIMO M.E. (2020).
 Development and evaluation of a double-antigen sandwich ELISA to identify *Anaplasma marginale*-infected and
 A. centrale-vaccinated cattle. J. Vet. Diagn. Invest., 32, 70–76.

<u>SHKAP V., KOCAN K., MOLAD T., MAZUZ M., LEIBOVICH B., KRIGEL Y., MICHOYTCHENKO A., BLOUIN E., DE LA FUENTE J., SAMISH
 <u>M., MTSHALI M., ZWEYGARTH E., FLEIDEROVICH E. L. & FISH L. (2009). Experimental transmission of field Anaplasma marginale</u>
 and the A. centrale vaccine strain by Hyalomma excavatum, Rhipicephalus sanguineus and Rhipicephalus (Boophilus)
 annulatus ticks. Vet. Microbiol., 134, 254–260.
</u>

841 <u>SILAGHI C., NIEDER M., SAUTER-LOUIS C., KNUBBEN-SCHWEIZER G., PFISTER K. & PFEFFER M. (2018). Epidemiology, genetic</u>
 842 <u>variants and clinical course of natural infections with *Anaplasma phagocytophilum* in a dairy cattle herd. *Parasit. Vectors*,
 843 <u>11, 20.</u>
</u>

 STICH R.W., OLAH G.A., BRAYTON K.A., BROWN W.C., FECHHEIMER M., GREEN CHURCH K., JITTAPALAPONC S., KOCAN K.M.,
 MCGUIRE T.C., RURANCIRWA F.R. & PALMER G.H. (2004).Identification of a novel Anaplasma marginale appendageassociated protein that localizes with actin filaments during intraerythrocytic infection. Infect Immun., 72, 7257–7264.

SILVA V.M., ARAUJO F.R., MADRUGA C.R., SOARES C.O., KESSLER R.H., ALMEIDA M.A., FRAGOSO S.P., SANTOS L.R., RAMOS
 C.A., BACANELLI G. & TORRES R.A. (2006). Comparison between indirect enzyme-linked immunosorbent assays for
 Anaplasma marginale antibodies with recombinant Major Surface Protein 5 and initial body antigens. Mem. Inst. Oswaldo
 Cruz., 101, 511–516.

STRIK N.I., ALLEMAN A.R., BARBET A.F., SORENSON H.L., WANSLEY WAMSLEY H.L., GASCHEN F.P., LUCKSCHANDER N., WONG S.,
 CHU F., FOLEY J.E., BJOERSDORFF A., STUEN S. & KNOWLES D.P. (2007). Characterization of *Anaplasma phagocytophilum* major surface protein 5 and the extent of its cross-reactivity with *A. marginale. Clin. Vaccine Immunol.*, 14, 262–268.

TORIONI DE ECHAIDE S., KNOWLES D.P., MCGUIRE T.C., PALMER G.H., SUAREZ C.E. & MCELWAIN T.F. (1998). Detection of cattle naturally infected with *Anaplasma marginale* in a region of endemicity by nested PCR and a competitive enzymelinked immunosorbent assay using recombinant major surface protein 5. *J. Clin. Microbiol.*, **36**, 777–782.

857	*
858	* *
859	NB: There is a WOAH Reference Laboratory for anaplasmosis (please consult the WOAH Web site:
860	https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3)
861	Please contact the WOAH Reference Laboratory for any further information on
862	diagnostic tests, reagents and vaccines for bovine anaplasmosis
863	NB: First adopted in 1991. Most recent updates adopted in 2015.

1	Annex 11. Item 5.1. – Chapter 3.4.7. Bovine viral diarrhoea
2	MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION
3	Paris, 4–8 September 2023
4	
5	CHAPTER 3.4.7.
6	BOVINE VIRAL DIARRHOEA

SUMMARY

7

8 9

14

16

21

Cattle of all ages are susceptible to infection with bovine viral diarrhoea viruses (BVDV), including BVDV type 1 (Pestivirus bovis), type 2 (Pestivirus tauri), and Hobi-like pestiviruses (type 3 [Pestivirus brazilense]). 10 Distribution is world-wide although some countries have recently eradicated the virus. BVDV infection results 11 in a wide variety of clinical manifestations, including enteric and respiratory disease in any class of cattle, or 12 reproductive and fetal disease following infection of a susceptible breeding female. Infection may be 13 subclinical or extend to severe fatal disease. Animals that survive in-utero infection in the first trimester of gestation are almost always persistently infected (PI). PI animals provide the main reservoir of the virus in a 15 population and excrete large amounts of virus in urine, faeces, discharges, milk and semen. Identification of such PI cattle is a key element in controlling the infection. It is important to avoid the trade of such animals. 17 They may appear clinically healthy, or weak and unthrifty. Many PI animals die before reaching maturity. 18 They may infrequently develop mucosal disease with anorexia, gastrointestinal erosions, and profuse diarrhoea, invariably leading to death. Mucosal disease can arise only in PI animals. Latent infections 19 generally do not occur following recovery from acute infection. However bulls may rarely have a persistent 20 testicular infection and excrete virus in semen for prolonged periods.

22 Detection of the agent: BVDV is a pestivirus in the family Flaviviridae and is closely related to classical 23 swine fever <u>virus (Pestivirus suis)</u> and ovine border disease viruses (<u>Pestivirus ovis)</u>. <u>BVD viruses are</u> classified into the distinct species Pestivirus bovis (commonly known as BVDV type 1), Pestivirus tauri 24 (BVDV type 2) and Pestivirus brazilense (BVDV type 3 or Hobi-like pestivirus). The two genotypes (types 1 25 and 2) are classified as separate species in the genus Pestivirus. A third putative genotype, BVDV type 3, 26 27 has also recently been proposed. Although both cytopathic and non-cytopathic biotypes of BVDV type 1 and 28 type 2 exist, non-cytopathic strains are usually encountered in field infections and are the main focus of diagnostic virus isolation in cell cultures. PI animals can be readily identified by a variety of methods aimed 29 to detect viral antigens or viral RNA directly in blood and tissues. Virus can also be isolated by inoculation 30 of specimens onto susceptible cell cultures followed by immune-labelling methods to detect the replication 31 32 of the virus in the cultures. Persistence of virus infection should be confirmed by resampling after an interval 33 of at least 3 weeks, when virus will again be detected. PI animals are usually seronegative. Viraemia in acute 34 cases is transient and usually difficult to detect. Virus isolation in semen from bulls that are acutely or persistently infected requires special attention to specimen transport and testing. RNA detection assays are 35 particularly useful because they are rapid, have very high sensitivity and do not depend on the use of cell 36 37 cultures.

38 Serological tests: Acute infection with BVDV is best confirmed by demonstrating seroconversion using sequential paired samples, ideally from several animals in the group. The testing of paired (acute and 39 convalescent samples) should be done a minimum of 21 days apart and samples should be tested 40

41 concurrently in the same assay. Enzyme-linked immunosorbent assays and the virus neutralisation test are 42 the most widely used.

43 Requirements for vaccines: There is no standard vaccine for BVD, but a number of commercial 44 preparations are available. An ideal vaccine should be able to prevent transplacental infection in pregnant cows. Modified live virus vaccine should not be administered to pregnant cattle (or to their sucking calves) 45 due to the risk of transplacental infection. Live vaccines that contain cytopathic strains of BVDV present a 46 risk of inducing mucosal disease in PI animals. Inactivated viral vaccines are safe and can be given to any 47 48 class of animal but generally require booster vaccinations. BVDV is a particularly important hazard to the 49 manufacture of vaccines and biological products for other diseases due to the high frequency of contamination of batches of fetal calf serum used as a culture medium supplement. 50

51

A. INTRODUCTION

52 **1. Impact of the disease**

53 Cattle of all ages are susceptible to infection with bovine viral diarrhoea viruses (BVDV). Distribution of the virus is worldwide although some countries have recently eradicated the virus. BVDV infection results in a wide variety of clinical 54 manifestations, including enteric and respiratory disease in any class of cattle or reproductive and fetal disease following 55 infection of a susceptible breeding female. Infection may be subclinical or extend to severe fatal disease. Clinical 56 57 presentations and severity of disease may vary with different strains of virus. BVDV viruses also cause immune suppression, which can render infected animals more susceptible to infection with other viruses and bacteria. The clinical 58 impact may be more apparent in intensively managed livestock. Animals that survive in-utero infection in the first trimester 59 of gestation are almost always persistently infected (PI). PI animals provide the main reservoir of the virus in a population 60 61 and excrete large amounts of virus in urine, faeces, discharges, milk and semen. The virus spreads mainly by close contact between PI animals and other cattle. Virus shedding by acutely infected animals is usually less important. This virus may 62 also persist in the environment for short periods or be transmitted with-via contaminated reproductive materials. Vertical 63 64 transmission plays an important role in its epidemiology and pathogenesis.

65 Infections of the breeding female may result in conception failure or embryonic and fetal infection which results in abortions, stillbirths, teratogenic abnormalities or the birth of PI calves. Persistently viraemic animals may be born as weak, unthrifty 66 calves or may appear as normal healthy calves and be unrecognised clinically for a long time. However, PI animals have 67 68 a markedly reduced life expectancy, with a high proportion dying before reaching maturity. Infrequently, some of these 69 animals may later develop mucosal disease with anorexia, gastrointestinal erosions, and profuse diarrhoea, invariably leading to death. Mucosal disease can arise only in PI animals. It is important to avoid the trade of viraemic animals. It is 70 71 generally considered that serologically positive, non-viraemic cattle are 'safe', providing that they are not pregnant. 72 However, a small proportion of persistently viraemic animals may produce antibodies to some of the viral proteins if they 73 are exposed to another strain of BVDV that is antigenically different to the persisting virus. Consequently, seropositivity cannot be completely equated with 'safety'. Detection of PI animals must be specifically directed at detection of the virus 74 or its components (RNA or antigens). Latent infections generally do not occur following recovery from acute infection. 75 However, semen collected from bulls during an acute infection is likely to contain virus during the viraemic period and often 76 for a short time afterwards. Although extremely rare, some recovered bulls may have a persistent testicular infection and 77 78 excrete virus in semen, perhaps indefinitely.

While BVDV strains are predominantly pathogens of cattle, interspecies transmission can occur following close contact with sheep, goats or pigs. Infection of pregnant small ruminants or pigs with BVDV can result in reproductive loss and the birth of PI animals. BVDV infections have been reported in both New World and Old World camelids. Additionally, strains of border disease virus (BDV) have infected cattle, resulting in clinical presentations indistinguishable from BVDV infection. The birth of cattle PI with BDV and the subsequent development of mucosal disease have also been described. Whilst BVDV and BDV have been reported as natural infections in pigs, the related virus of classical swine fever does not naturally infect ruminants.

Although ubiquitous, control of BVDV can be achieved at the herd level, and even at the national level, as evidenced by the progress towards eradication made in many European countries (Moennig *et al.*, 2005<u>; Schweizer *et al.*, 2021</u>).

88 2. The causal agent

Bovine viral diarrhoea virus (BVDV) is a single linear positive-stranded RNA virus in the genus *Pestivirus* of the family *Flaviviridae*. The genus contains a number of species including the two genotypes of bovine viral diarrhoea virus (BVDV)
(types 1 [*Pestivirus bovis*], and 2 [*Pestivirus tauri*] and 3 [*Pestivirus brazilense*]) and the closely related classical swine
fever (*Pestivirus suis*) and ovine border disease viruses (*Pestivirus ovis*). Viruses in these genotypes pestivirus species
show considerable antigenic difference from each other and, within the type 1 and type 2 species *Pestivirus bovis* and

94 tauri, BVDV isolates exhibit considerable biological and antigenic diversity. Within the two BVDV genotypes species 95 Pestivirus bovis and tauri, further subdivisions are discernible by genetic analysis (Vilcek et al., 2001). The two genetypes 96 species may be differentiated from each other, and from other pestiviruses, by monoclonal antibodies (MAbs) directed against the major glycoproteins E2 and ERNS or by genetic analysis. Reverse-transcription polymerase chain reaction (RT-97 98 PCR) assays enable virus typing direct from blood samples (Letellier & Kerkhofs, 2003; McGoldrick et al., 1999). Type 1 viruses are generally more common although the prevalence of type 2 strains can be high in North America. BVDV of both 99 genotypes species (Pestivirus bovis and tauri) may occur in non-cytopathic and cytopathic forms (biotypes), classified 100 according to whether or not microscopically apparent cytopathology is induced during infection of cell cultures. Usually, it 101 is the non-cytopathic biotype that circulates freely in cattle populations. Non-cytopathic strains are most frequently 102 responsible for disease in cattle and are associated with enteric and respiratory disease in any class of cattle or 103 reproductive and fetal disease following infection of a susceptible breeding female. Infection may be subclinical or extend 104 to severe fatal disease (Brownlie, 1985). Cytopathic viruses are encountered in cases of mucosal disease, a clinical 105 syndrome that is relatively uncommon and involves the 'super-infection' of an animal that is PI with a non-cytopathic virus 106 by a closely related cytopathic strain. The two virus biotypes found in a mucosal disease case are usually antigenically 107 108 closely related if not identical. Type 2 viruses are usually non-cytopathic and have been associated with outbreaks of severe acute infection and a haemorrhagic syndrome. However some type 2 viruses have also been associated with a 109 110 disease indistinguishable from that seen with the more frequently isolated type 1 viruses. Further, some type 1 isolates 111 have been associated with particularly severe and fatal disease outbreaks in adult cattle. Clinically mild and inapparent infections are common following infection of non-pregnant animals with either genotype-virus species. 112

113 There is an increasing awareness of an "atypical" or "HoBi like" pestivirus a putative BVDV type 3 Pestivirus H strains are also associated with clinical disease in cattle, but they appear mainly restricted to South American and Asian cattle 114 populations, in cattle, also associated with clinical disease (Bauermann et al., 2013; Chen et al., 2021), but its distribution 115 is presently unclear. These viruses are readily detected by proven pan-reactive assays such as real-time RT-PCR. Some 116 commercial antigen ELISAs (enzyme-linked immunosorbent assays) have been shown to detect these strains (Bauermann 117 118 et al., 2012); generally virus isolation, etc., follows the same principles as for BVDV 1 (Pestivirus bovis) and 2 (Pestivirus tauri). It should be noted however, that antibody ELISAs vary in their ability to detect antibody to BVDV type 3 (Pestivirus 119 brazilense) and vaccines designed to protect against BVDV 1 and 2 may not confer full protection against infection with 120 these novel pestiviruses (Bauermann et al., 2012; 2013). 121

122 **3.** Pathogenesis

123

3.1. Acute infections

Acute infections with BVDV are encountered more frequently in young animals, and may be clinically inapparent or 124 associated with fever, diarrhoea (Baker 1995), respiratory disease and sometimes sudden death. The severity of disease 125 126 may vary with virus strain and the involvement of other pathogens (Brownlie, 1990). In particular, outbreaks of a severe form of acute disease with haemorrhagic lesions, thrombocytopenia and high mortality have been reported sporadically 127 from some countries (Baker, 1995; Bolin & Ridpath, 1992). Infection with type 2 viruses (Pestivirus tauri) in particular 128 has been demonstrated to cause altered platelet function. During acute infections there is a brief viraemia for 7-10 days 129 and shedding of virus can be detected in nasal and ocular discharges. There may also be a transient leukopenia, 130 thrombocytopenia or temperature response, but these can vary greatly among animals. Affected animals may be 131 predisposed to secondary infections with other viruses and bacteria. Although BVDV may cause a primary respiratory 132 disease on its own, the immunosuppressive effects of the virus exacerbate the impact of this virus. BVDV is one of the 133 134 major pathogens of the bovine respiratory disease complex in feedlot cattle and in other intensive management systems 135 such as calf raising units.

Infection of breeding females immediately prior to ovulation and in the first few days after insemination can result in
 conception failure and early embryonic loss (McGowan & Kirkland, 1995). Cows may also suffer from infertility,
 associated with changes in ovarian function and secretions of gonadotropin and progesterone (Fray *et al.*, 2002).
 Bulls may excrete virus in semen for a short period during and immediately after infection and may suffer a temporary
 reduction of fertility. Although the virus level in this semen is generally low it can result in reduced conception rates
 and be a potential source of introduction of virus into a naive herd (McGowan & Kirkland, 1995).

142 **3.2.** *In-utero* infections

Infection of a breeding female can result in a range of different outcomes, depending on the stage of gestation at 143 which infection occurred. Before about 25 days of gestation, infection of the developing conceptus will usually result 144 145 in embryo-fetal death, although abortion may be delayed for a considerable time (McGowan & Kirkland, 1995). Surviving fetuses are normal and uninfected. However, infection of the female between about 30-90 days will 146 invariably result in fetal infection, with all surviving progeny PI and seronegative. Infection at later stages and up to 147 148 about day 150 can result in a range of congenital defects including hydranencephaly, cerebellar hypoplasia, optic defects, skeletal defects such as arthrogryposis and hypotrichosis. Growth retardation may also occur, perhaps as a 149 result of pituitary dysfunction. Fetal infection can result in abortion, stillbirth or the delivery of weak calves that may 150 die soon after birth (Baker, 1995; Brownlie, 1990; Duffell & Harkness, 1985; Moennig & Liess, 1995). Some PI calves 151

may appear to be normal at birth but fail to grow normally thrive. They remain PI for life and are usually seronegative.
 <u>exceptions may be young calves that ingested colostrum containing antibodies</u>. The onset of the fetal immune
 response and production of antibodies occurs between approximately day 90–120, with an increasing proportion of
 infected calves having detectable antibodies while the proportion in which virus may be detected declines rapidly.
 Infection of the bovine fetus after day 180 usually results in the birth of a normal seropositive calf.

157 **3.3. Persistent infections**

Persistently viraemic animals are a continual source of infective virus to other cattle and are the main reservoir of BVDV in a population. In a population without a rigorous BVDV control programme, approximately 1–2% of cattle are PI. During outbreaks in a naive herd or breeding group, if exposure has occurred in the first trimester of pregnancy, a very high proportion of surviving calves will be PI. If a PI animal dies, there are no pathognomonic lesions due to BVDV and the pathology is often complicated by secondary infections with other agents. Some PI animals will survive to sexual maturity and may breed successfully but their progeny <u>of female PI animals</u> will also always be PI. Animals being traded or used for artificial breeding should first be screened to ensure that they are not PI.

165 **3.4. Mucosal disease**

Persistently viraemic animals may later succumb to mucosal disease (Brownlie, 1985). However, cases are rare. This syndrome has been shown to be the outcome of the infection of a PI animal with an antigenically similar cytopathic virus, which can arise either through superinfection, recombination between non-cytopathic biotypes, or mutation of the persistent biotype (Brownlie, 1990). There is usually little need to specifically confirm that a PI animal has succumbed to mucosal disease as this is largely a scientific curiosity and of little practical significance, other than that the animal is PI with BVDV. However, cases of mucosal disease may be the first indication in a herd that BVDV infection is present, and should lead to more in depth investigation and intervention.

173 **3.5. Semen and embryos**

Bulls that are PI usually have poor quality, highly infective semen and reduced fertility (McGowan & Kirkland, 1995). 174 All bulls used for natural or artificial insemination should be screened for both acute and persistent BVDV infection. 175 A rare event, possibly brought about by acute infection during pubescence, can result in persistent infection of the 176 177 testes and thus strongly seropositive bulls that intermittently excrete virus in semen (Voges et al., 1998). This phenomenon has also been observed following vaccination with an attenuated virus (Givens et al., 2007). Embryo 178 donor cows that are PI with BVDV also represent a potential source of infection, particularly as there are extremely 179 high concentrations of BVDV in uterine and vaginal fluids. While oocvsts without an intact zona pellucida have been 180 181 shown to be susceptible to infection in vitro, the majority of oocysts remain uninfected with BVDV. Normal uninfected progeny have also been 'rescued' from PI animals by the use of extensive washing of embryos and in-vitro fertilisation. 182 Female cattle used as embryo recipients should always be screened to confirm that they are not PI, and ideally, are 183 seropositive or were vaccinated at least 4 weeks before first use. 184

Biological materials used for *in-vitro* fertilisation techniques (bovine serum, bovine cell cultures) have a high risk of contamination and should be screened for BVDV. Incidents of apparent introduction of virus via such techniques have highlighted this risk. It is considered essential that serum supplements used in media should be free of contaminants as detailed in Chapter 1.1.9 *Tests for sterility and freedom from contamination of biological materials intended for veterinary use*, using techniques described in Section B.3-1.1 of this chapter.

4. Approaches to diagnosis and sample collection

191 The diagnosis of BVDV infection can sometimes be complex because of the delay between infection and clinical 192 expression. While detection of PI animals should be readily accomplished using current diagnostic methods, the recognition 193 of acute infections and detection of BVDV in reproductive materials can be more difficult.

194 **4.1. Acute infections**

195 Unlike PI animals, acutely infected animals excrete relatively low levels of virus and for a short period of time (usually about 7-10 days) but the clinical signs may occur during the later stages of viraemia, reducing the time to detect the 196 197 virus even further. In cases of respiratory or enteric disease, samples should be collected from a number of affected animals, preferentially selecting the most recently affected. Swabs should be collected from the nares and conjunctiva 198 of animals with respiratory disease or from rectum and faeces if there are enteric signs. Lung and spleen are preferred 199 from dead animals. Viral RNA may be detected by real-time RT-PCR assays and have the advantages of high 200 sensitivity and being able to detect genome from non-infectious virus. As the virus levels are very low, it is not usually 201 practical to undertake virus isolation unless there is a need to characterise the strain of BVDV involved. Serology 202 203 undertaken on paired acute and convalescent sera (collected at least 21 days after the acute sample and from 8-10 animals) is worthwhile and gives a high probability of incriminating or excluding BVDV infection. 204

205 Confirmation that an abortion, stillbirth or perinatal death is caused by BVDV is often difficult to establish because 206 there can be a long delay between initial infection and death or expulsion of the fetus. Sampling should take into consideration the need to detect either viral components or antibodies. Spleen and lung are preferred samples for 207 208 virus detection while pericardial or pleural fluids are ideal samples for serology. The stomach of newborn calves should be checked to confirm that sucking has not occurred. While virus may be isolated from fetal tissue in some 209 cases, emphasis should be placed on the detection of viral antigen by ELISA or RNA by real-time RT-PCR. For 210 serology, both ELISAs and virus neutralisation test (VNT) are suitable though sample quality and bacterial 211 contamination may compromise the ability to detect antibodies by VNT. Maternal serology, especially on a group of 212 animals, can be of value, with the aim of determining whether there has been recent infection in the group. A high 213 antibody titre (>1/1000) to BVDV in maternal serum is suggestive of fetal infection and is probably due to the fetus 214 215 providing the dam with an extended exposure to virus.

216 4.2. Persistent infections

In the past, identification of PI animals relied heavily on the use of virus isolation in cell cultures. However, antigen 217 detection ELISAs and real-time RT-PCR assays, each with relatively high sensitivity, are widely used for the detection 218 of viral antigens or RNA in both live and dead animals. Virus isolation aimed at the detection of non-cytopathic BVDV 219 220 in blood is also used, while in some countries, the virus has been identified by immunohistochemistry (IHC). Skin samples have been collected from live animals while a wide range of tissues from dead animals are suitable. Both 221 virus isolation and IHC are labour intensive and costly and can be technically demanding. Virus isolation from blood 222 can be confounded by the presence of maternal antibody to BVDV in calves less than 4-5 months of age (diagnostic 223 224 gap). Also for antigen detection ELISAs and flow cytometry from blood or blood leukocytes, there are restrictions that 225 limit when animals that ingested colostrum that contains antibodies against BVDV can be reliably tested. In older 226 animals with persistent viraemia infection, low levels of antibody may be present due to their ability to seroconvert to strains of BVDV (including vaccines) antigenically different to the persisting virus (Brownlie, 1990). Bulk (tank) or 227 individual milk samples have been used to monitor dairy herds for the presence of a PI animal. Antigen ELISA, real-228 time PCR and virus isolation have all been used. To confirm a diagnosis of persistent infection, animals should be 229 retested after an interval of at least 3 weeks by testing of blood samples for the presence of the virus and for evidence 230 absence of seroconversion. Care should be taken with retesting of skin samples as it has been shown that, in some 231 acute cases, viral antigen may persist for many weeks in skin (Cornish et al., 2005). 232

233 4.3. Mucosal disease

Although not undertaken for routine diagnostic purposes, for laboratory confirmation of a diagnosis of mucosal disease it is necessary to isolate the cytopathic virus. This biotype may sometimes be isolated from blood, but it can be recovered more consistently from a variety of other tissues, in particular spleen, intestine and Peyer's patch tissue. Virus isolation is readily accomplished from spleen which is easy to collect and is seldom toxic for cell culture.

238 4.4. Reproductive materials

239 Semen donor bulls should be sampled for testing for freedom from BVDV infection prior to collection of semen, in accordance with the Terrestrial Animal Health Code. It is necessary to confirm that these bulls are not PI, are not 240 undergoing an acute infection and to establish their serological status. This initial testing should be carried out on 241 whole blood or serum samples. To establish that a seropositive bull does not have a persistent testicular infection 242 243 (PTI), samples of semen should be collected on at least three separate occasions at intervals of not less than 7 days due to the possibility of intermittent low level virus excretion, especially during the early stages of infection. There is 244 also a need to submit a number of straws from each collection, or an appropriate volume of raw semen. Particular 245 care should be taken to ensure that sample transport recommendations are adhered to and that the laboratory 246 247 documents the condition of the samples on arrival at the laboratory. Further details of collection, transport and test requirements are provided in sections that follow. 248

B. DIAGNOSTIC TECHNIQUES

Purpose						
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	<u>Contribute to</u> <u>eradication</u> <u>policies</u>	Confirmation of clinical cases	<u>Prevalence of</u> <u>infection –</u> <u>surveillance</u>	Immune status in individual animals or populations (post- vaccination)
		Detec	ction of the ag	jent ^(a)		
Virus isolation	+	++ ±	++	++ +	_	-
Antigen detection by ELISA	++ <u>+</u>	+++	+++	+++	+++	_
Antigen detection by IHC	-	_	_	++	_	_
NA detection by real-time RT-PCR	+++	+++	+++	+++	+++	-
Detection of immune response						
ELISA	+++	++	+++	- <u>+</u> ^(a)	+++	+++
VN	+	++_+	++	-	+	+++

Table 1. Test methods available for diagnosis of bovine viral diarrhoea and their purpose

251 252 253

249

250

52

253

254 255 + = suitable in very limited circumstances; - = not appropriate for this purpose. ELISA = enzyme-linked immunosorbent assay; IHC = immunohistochemistry method; NA = nucleic acid; RT-PCR = reverse-

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

transcription polymerase chain reaction; ISH = *in-situ* hybridisation; VN = virus neutralisation. ^(a)A combination of agent detection methods applied on the same clinical sample is recommended.

1. Detection of the agent

To prevent the shipment of either animals or animal derivatives (especially semen and embryos) that are infected with 257 258 BVDV, it is necessary to test for the presence of the infectious virus (virus isolation), viral antigens (antigen detection ELISA) or RNA (real-time RT-PCR) in the blood of the animal being shipped, or the donor of the germplasm (semen or 259 embrvos). The exception is for seropositive bulls where semen must be tested rather than the donor bull. Serology only 260 261 plays a role for establishing that seronegative animals are not undergoing an acute infection or, to establish the serological status of donor bulls. Due to their variable sensitivity without prior virus amplification, procedures such as IHC or in-situ 262 hybridisation (ISH) directly on tissues are not considered to be suitable for certification for freedom from BVDV for 263 international trade purposes. In contrast, immune-staining is an essential component of virus isolation in cell culture to 264 detect the presence of non-cytopathic strains of BVDV which predominate in field infections. 265

All test methods must be extensively validated by testing on known uninfected and infected populations of cattle, including animals with low- and high-titre viraemias. Methods based on polyclonal or MAb-binding assays (ELISA or IHC), immune labelling (VI) or on nucleic acid recognition (PCR) must be shown to detect the full range of antigenic and genetic diversity found among BVD viruses. There are three-designated WOAH Reference Laboratories for BVDV that can assist with relevant information; the reference laboratories for classical swine fever could also be approached to offer some advice.

271 **1.1. Virus isolation**

When performed to a high standard, BVDV isolation is very reliable. However, it does have very exacting requirements to ensure that the cell cultures and medium components give a system that is very sensitive and are not compromised by the presence of either low levels of BVDV specific antibody or virus. Virus isolation only has the capacity to detect infectious virus which imposes certain limits on sample quality. Further, to detect low levels of virus that may be present in some samples, particularly semen, it may be necessary to examine larger volumes of specimen than is usual. Some of these limitations can be overcome by the use of antigen detection ELISAs with proven high analytical sensitivity, or the use of real-time RT-PCR. 279 The virus may be isolated in a number of bovine monolayer cell cultures (e.g. kidney, lung, testis or turbinate). In 280 some instances, ovine cells are also suitable. Primary or secondary cultures can be frozen as cell suspensions in liquid nitrogen. These can then be tested over a series of passages, or seeded to other susceptible cells and checked 281 282 for freedom from contaminants and to evaluate their sensitivity compared to an approved batch of cells before routine use. Such problems may be reduced by the use of continuous cell lines, which can be obtained BVD-free, however, 283 284 their BVDV-free status and susceptibility must be monitored regularly. Continuous cells should be used under a 'seed lot' system where they are only used over a limited passage range, within which they have been shown to have 285 acceptable sensitivity to BVDV infection. Although particular continuous cell lines are considered to be appropriate 286 for use for BVDV isolation, there can be significant variation in batches of cells from different sources due to differing 287 passage histories so their suitability must still be confirmed before routine use. 288

- 289 Non-cytopathic BVDV is a common contaminant of bovine tissues and cell cultures must be checked for freedom from adventitious virus by regular testing. Cells must be grown in proven cell culture medium components and a large 290 area of cells must be examined. It is not appropriate to screen a few wells of a 96 well plate - examining all wells of 291 a 96 well plate will be more convincing evidence of freedom. The fetal bovine serum that is selected for use in cell 292 culture must also be free not only from virus, but also and of equal or perhaps even greater importance, from BVDV 293 294 neutralising antibody. Heat treatment (56°C for 30-45 minutes) is inadequate for the destruction of BVDV in contaminated serum; irradiation with a dose of at least 25 kiloGrays (2.5 Mrad) is more certain. Commercial batches 295 296 of fetal bovine serum mostly test positive by real-time RT-PCR even after the virus has been inactivated by irradiation. 297 Further, most commercially collected batches of fetal bovine serum contain antibodies to pestiviruses, sometimes at levels that are barely detectable but sufficient to inhibit virus isolation. To overcome this, serum can be obtained from 298 BVD virus and antibody free donor animals and used with confidence. Testing of donors for both virus and antibody 299 occurs on an individual animal basis. Although horse serum has been substituted for bovine fetal serum, it is often 300 301 found to have poorer cell-growth-promoting characteristics. Further there has sometimes been cross contamination 302 with fetal bovine serum during processing, negating the objective of obtaining a BVDV-free product.
- 303 Buffy coat cells, whole blood, washed leukocytes or serum are suitable for isolation of the virus from live animals. 304 Maternal antibody may interfere with isolation from serum in young calves. Tissue suspensions from post-mortem 305 cases should be prepared by standard methods. Confirmation that a bull is not PI with BVDV is most readily achieved 306 by testing of a blood sample. However, persistent testicular infections (PTI) have been detected in some bulls that have recovered from acute infection, are no longer viraemic and are now seropositive (Voges et al., 1998). Virus may 307 be detected in most but not all collections of semen from these bulls. Although still considered to be uncommon, to 308 exclude the potential for a PTI it is essential to screen semen from all seropositive bulls. To be confident that a bull 309 310 does not have a PTI, batches of semen collected over several weeks should be screened. Once a series of collections 311 have been screened, further testing of semen from a seropositive bull is not warranted. Raw semen, and occasionally extended semen, is cytotoxic and must be diluted in culture medium. For these reasons, it is important to monitor the 312 313 health of the cells by microscopic examination at intervals during the incubation. These problems are largely overcome by the use of real-time RT-PCR which has several advantages over virus isolation, including higher 314 sensitivity and the potential to be completed within a few hours rather than weeks for virus isolation. 315
- 316 There are many variations of procedure in use for virus isolation. All should be optimised to give maximum sensitivity 317 of detection of a standard virus preparation. All biological components used for cell culture should be screened and 318 shown to be free of both BVDV and antibodies to BVDV. Cell cultures (whether primary or continuous lines) should be regularly checked to confirm that they maintain maximum susceptibility to virus infection. Depending on the 319 specimen type and purpose for testing, virus isolation is likely to require one or more passages in cell cultures. While 320 321 PI animals can be readily identified by screening blood or serum with one passage, semen should be routinely cultured for three passages and biological products such as fetal bovine serum up to five times (original inoculation 322 323 plus four passages). Conventional methods for virus isolation are used, with the addition of a final immune-staining step (immunofluorescence or, more frequently, peroxidase staining) to detect growth of non-cytopathic virus. Thus 324 tube cultures should include flying cover-slips, while microplate cultures can be fixed and labelled directly in the plate. 325 Examples are given below. Alternatively, culture supernatant from the final passage can be screened by real-time 326 RT-PCR (see below). 327

328

329

330

331

332 333

334

335

336

1.1.1. Microplate immunoperoxidase method for mass screening for virus detection in serum samples (Meyling, 1984)

- 10-25 µl of the serum sample is placed into each of four wells of a 96-well tissue-culture grade i) microplate. This is repeated for each sample. Known positive and negative controls are included.
- ii) 100 µl of a cell suspension at the appropriate concentration (usually about 150,000 cells/ml) in medium without fetal calf serum (FCS) is added to all wells. Note: the sample itself acts as the cell-growth supplement. If testing samples other than serum, use medium with 10% FCS that is free of antibodies to ruminant pestiviruses.

337 338	iii)	The plate is incubated at 37° C for 4 days, either in a 5% CO ₂ atmosphere or with the plate sealed.
339 340	iv)	Each well is examined microscopically for evidence of cytopathology (cytopathic effect or CPE), or signs of cytotoxicity.
341 342	v)	The cultures are frozen briefly at approximately -80° C and 50 µl of the culture supernatant is passaged to new cell cultures, repeating steps 3 <u>1</u> .1.1.i to iv above.
343	vi)	The cells are then fixed and stained by one of two methods:
344	•	Paraformaldehyde
345	a)	Add 200 µl of a 1/10 dilution of formaldehyde solution (approximately 3% concentration) to
346		the plate and leave at room temperature for 10 minutes.
347	b)	The contents of the plate are then discarded and the plate is washed.
348 349	c)	Wash plates 5 times with 0.05% Tween 20 in water (an automatic microplate washer can be used with a low pressure and speed setting).
350 351 352	d)	To each well add 50 μ l of an antiviral antibody at the appropriate dilution (prepared in phosphate buffered saline/ PBS containing 1% gelatin) and incubate for 60–90 minutes at 37°C in a humidified chamber.
353	e)	Wash plates five times as in step c).
354 355 356 357	f)	Dilute the appropriate peroxidase conjugated antiserum to the optimum dilution in 1% gelatin/PBS (e.g. peroxidase conjugated rabbit anti-mouse immunoglobulin when the antiviral antibody is a mouse monoclonal). The optimum concentration should be determined for each batch of conjugate by "checkerboard" titration against reference positive and negative controls.
358 359	g)	To each well of the microplate add 50 μ l of the diluted peroxidase conjugate and incubate for 90 minutes at 37°C in a humidified chamber.
360	h)	Wash plates five times as in step c).
361 362	i)	"Develop" the plate by adding 3-amino-9-ethyl carbazole (AEC) substrate (100 μl/well) and allowing to react for 30 minutes at room temperature.
363	j)	Add 100 μI of PBS to each well and add a lid to each plate.
364 365 366	k)	Examine the wells by light microscopy, starting with the negative and positive control wells. There should be no or minimal staining apparent in the cells that were uninfected (negative control). The infected (positive control) cells should show a reddish- brown colour in the cytoplasm.
367	٠	Acetone
368	a)	The plate is emptied by gentle inversion and rinsed in PBS.
369 370 371 372 373	b)	The cells are fixed as follows: the plate is dipped into a bath of 20% acetone in PBS, emptied immediately and then transferred to a fresh bath of 20% acetone in PBS for 10 minutes. The plate is drained thoroughly and as much fluid as possible is removed by tapping and blotting. The plate is dried thoroughly for at least 3 hours at a temperature of 25–30°C (e.g. using radiant heat from a bench lamp). <i>Note:</i> the drying is part of the fixation process.
374	c)	The fixed cells are rinsed by adding PBS to all wells.
375 376 377	d)	The wells are drained and the BVD antibody (50 μ I) is added to all wells at a predetermined dilution in PBS containing 1% Tween 80 (PBST) and 5% horse serum or 1% gelatin. (Horse serum or gelatin may be added to reduce nonspecific staining.)
378	e)	Incubate at 37°C for 15 minutes.
379	f)	Empty the plate and wash three times in PBST.
380 381	g)	Drain and add the appropriate anti-species serum conjugated to peroxidase at a predetermined dilution in PBST (50 μ l per well) for 15 minutes at 37°C.
382	h)	Empty the plate and wash three times in PBST.
383	i)	Rinse the plate in distilled water. <u>Ensure</u> all fluid is tapped out from the plate.
384 385	j)	Add freshly prepared hydrogen peroxide substrate with a suitable chromogen, e.g. 3-amino- 9-ethyl carbazole (AEC).

386 387 388		An alternative substrate can be made, consisting of 9 mg diaminobenzidine tetrahydrochloride and 6 mg sodium perborate tetrahydrate dissolved in 15 ml of PBS. Though the staining is not quite so intense, these chemicals have the advantage that they can be shipped by air.
389 390	k)	The plate is examined microscopically. Virus-positive cells show red-brown cytoplasmic staining.
391 392 393	3.8.3	rnative methods for fixation of the cells may be used and include the use of heat (see Chapter 3 <i>Classical swine fever</i> , Section B.2.2.1.viii). These should be first evaluated to ensure that the acity to detect viral antigen is not compromised.
394	1.1.2. Tub	e method for tissue or buffy coat suspensions
395 396		is method can also be conveniently adapted to 24-well plastic dishes. <i>Note:</i> a minimum of 2 and bly 3 passages (including primary inoculation) is required.
397 398		ssue samples are ground up and a 10% suspension in culture medium is made. This is then ntrifuged to remove the debris.
399 400		est tube cultures with newly confluent or subconfluent monolayers of susceptible bovine cells are oculated with 0.1 ml of the sample. The culture is left to adsorb for 1 hour at 37°C.
401 402		e culture is washed with 1 ml of medium; this is then discarded and 1 ml of culture maintenance edium is added.
403 404		e culture is incubated for 4–5 days at 37°C, and examined microscopically for evidence of CPE signs of cytotoxicity.
405 406 407 408 409 410 411 412 413 414	tw pa for im us in sli pe	the culture should then be frozen and thawed for passage to fresh cultures for one or preferably o more passages (including the culture inoculated for the final immunostaining). At the final issage, after freeze-thaw the tissue culture fluid is harvested and passaged on to microtitre plates r culture and staining by the immunoperoxidase method (see section B.3 <u>1</u> .1.1 above) or by the munofluorescent method. For immunofluorescence, cover-slips are included in the tubes and ed to support cultured cells. At the end of the culture period, the cover slips are removed, fixed 100% acetone and stained with an immunofluorescent conjugate to BVDV. Examine the cover ps under a fluorescent microscope for diffuse, cytoplasmic fluorescence characteristic of estiviruses. Alternatively, culture supernatant from the final passage can be screened by real-time T-PCR (see below).
415	1.1.3. Viru	is isolation from semen
416 417 418 419 420 421 422 423 424 425 426 427	Sem shou term unde 1/10 shou sem mini enco raw	samples used for the test are, typically, extended bovine semen or occasionally raw semen. Then samples should be transported to the laboratory in liquid nitrogen, or on dry ice. The samples all be stored in liquid nitrogen or at lower than -70° C (for long-term storage) or 4°C (for short- in storage of not more than 1–2 days). The receiving laboratory should document the condition er which samples are received. Raw semen is generally cytotoxic and should be prediluted (e.g. 0 in BVDV free bovine serum) before being added to cell cultures. At least 0.1 ml of raw semen all be tested with three passages in cell culture. Toxicity may also be encountered with extended en. For extended semen, an approximation should be made to ensure that the equivalent of a mum of 0.1 ml raw semen is examined (e.g. a minimum of 1.0 ml extended semen). If toxicity is pountered, multiple diluted samples may need to be tested to reach a volume equivalent to 0.1 ml semen (e.g. 5 × 1 ml of a sample of extended semen that has been diluted 1/5 to reduce toxicity). Iggested method is as follows:
428 429 430	i)	Dilute 200 µl fresh semen in 1.8 ml bovine serum containing antibiotics. This can be the same serum as is being used for supplementing the cell cultures, and must be shown to be free from antibodies against BVDV.
431	ii)	Mix vigorously and leave for 30 minutes at room temperature.
432 433	iii)	Inoculate 1 ml of the semen/serum mixture into a monolayer of susceptible cells (see virus isolation from tissue above) in cell culture tubes or a six-well tissue culture plate.
434	iv)	Incubate the cultures for 1 hour at 37°C.
435 436	v)	Remove the mixture, wash the monolayer several times with maintenance medium and then add new maintenance medium to the cultures.

437 vi) Include BVDV negative and positive controls in the test. Special caution must be taken to avoid 438 accidental contamination of test wells by the positive control, for example always handling the positive control last. 439 440 vii) Observe plates microscopically to ensure freedom from contamination and cytotoxicity. No cytopathology is expected as a result of BVDV infection but other viruses such as BHV-1 could 441 be inadvertently isolated. 442 443 viii) After 5–7 days, the cultures are frozen at or below approximately –70°C and thawed, clarified by centrifugation, and the supernatant used to inoculate fresh monolayers. 444 At the end of the second passage, the supernatant from the freeze-thaw preparation should be 445 ix) 446 passaged onto cultures in a suitable system for immunoperoxidase staining or other antigen detection or by real-time RT-PCR after 5 days of culture. This is most readily achieved in 96 447 well microplates. The sample is considered to be negative, if there is no evidence of viral antigen 448 449 or BVDV RNA detected.

450 **1.2. Nucleic acid detection**

451 Conventional gel based RT-PCR has in the past been used for the detection of BVD viral RNA for diagnostic 452 purposes. A multiplex RT-PCR has been used for the simultaneous amplification and typing of virus from cell culture, or direct from blood samples. However, gel based RT-PCR has the disadvantage that it is relatively labour intensive, 453 454 expensive and prone to cross contamination. These problems had been markedly reduced following the introduction of probe-based real-time or quantitative RT-PCR methods. Nevertheless, stringent precautions should still be taken 455 456 to avoid nucleic acid contamination in the test system and general laboratory areas where samples are handled and prepared (see Chapter 1.1.6 Principles and methods of validation of diagnostic assays for infectious diseases and 457 Chapter 2.2.3 Development and optimisation of nucleic acid assays). These assays have even higher sensitivity than 458 gel based RT-PCR and can be completed in a few hours. They are in widespread use for the diagnosis of infectious 459 diseases, allowing the direct detection of viral RNA from a wide range of specimens including serum, whole blood, 460 tissues, milk and semen. The high analytical sensitivity allows the adoption of strategies to screen pools of individual 461 462 samples or testing of bulk tank milk. By using this approach the presence of one or more PI animals can be identified in herds containing several hundred cows. However, it is not appropriate to pool blood samples taken from calves 463 464 between day 7 and 40 of life, when colostrum that contains antibodies against BVDV was ingested. During this time 465 the sensitivity of PCR can be reduced and infected animals escape detection. In contrast, the detection of viral RNA in skin biopsy samples remains unaffected (Fux & Wolf, 2012). Although slightly more expensive than immunostaining 466 467 methods, real-time RT-PCR is a quick and reliable method that can also be used to screen culture supernatant from the final passage of cell cultures. While real-time RT-PCR has very high sensitivity and can be applied to the screening 468 469 of biological materials used for vaccine manufacture, caution is needed in the interpretation of results, as the detection of viral RNA does not imply per se that infective virus is present. Real-time RT-PCR assays based on fluorescent-470 labelled DNA probes can also be used to differentiate pestiviruses (e.g. McGoldrick et al., 1999). 471

472 Primers for the assay should be selected in highly conserved regions of the genome, ideally the 5'-noncoding region, or the NS3 (p80 gene). There are published assays that are broadly reactive across the pestivirus genus, detecting 473 all BVDV types (Pestivirus bovis, tauri and brazilense), CSFV (Pestivirus suis), some strains of BDV (Pestivirus ovis) 474 and most of the several 'atypical' pestiviruses (e.g. Hoffman et al., 2006). A sensitive broadly reactive assay is 475 recommended for diagnostic applications because interspecies transfer of different pestiviruses is occasionally 476 encountered. When further identification of the specific virus is required, pestivirus species-specific assays can be 477 478 applied to further type the virus. It is important to thoroughly optimise all aspects of the real-time RT-PCR assay, including the nucleic acid extraction and purification. Optimal concentrations of Mg²⁺, primers, probe and polymerase, 479 and the cycling parameters need to be determined. However, fully formulated and optimised 'ready to use' 480 'mastermixes' are now available commercially and only require addition of optimised concentrations of primers and 481 probe. Optimised cycling conditions are often recommended for a particular mastermix. 482

483 A variety of commercially available nucleic acid purification systems are available in kit form and several can be semiautomated. Systems based on the capture and purification of RNA using magnetic beads are in widespread use and 484 485 allow rapid processing of large numbers of samples. Specific products should be evaluated to determine the optimal 486 kit for a particular sample type and whether any preliminary sample processing is required. For whole blood samples, the type of anticoagulant and volume of blood in a specimen tube is important. More problems with inhibitors of the 487 PCR reaction are encountered with samples collected into heparin treated blood than EDTA. These differences are 488 also exacerbated if the tube does not contain the recommended volume of blood, thereby increasing the concentration 489 of anticoagulant in the sample. To identify possible false-negative results, it is recommended to spike an exogenous 490 491 ('internal control') RNA template into the specimen prior to RNA extraction (e.g. Hoffman et al., 2006). By the inclusion of PCR primers and probe specific to the exogenous sequence, the efficiency of both the RNA extraction and also 492 493 the presence of any PCR inhibitors can be monitored. While valuable for all sample types, the inclusion of an internal control is particularly desirable when testing semen and whole blood. When using an internal control, extensive testing 494

is necessary to ensure that PCR amplification of the internal control does not compete with the diagnostic PCR and
 thus lower the analytical sensitivity (see also chapter 1.1.6).

When it is suspected that a sample may contain substances that are adversely affecting either the efficiency of RNA extraction or the real-time RT-PCR assay, modest dilution of the sample in saline, cell culture medium or a buffer solution (e.g. <u>phosphate buffered gelatin saline [PBGS]</u>) will usually overcome the problem. Dilution of a semen sample by 1/4 and whole unclotted blood at 1/10 is usually adequate. As the real-time RT-PCR has extremely high analytical sensitivity, dilution of the sample rarely has a significant impact on the capacity of the assay to detect viral RNA when present.

1.2.1. Real-time polymerase chain reaction for BVDV detection in semen

Real-time RT-PCR has been shown to be extremely useful to screen semen samples to demonstrate freedom from BVDV and, apart from speed, often gives superior results to virus isolation in cell culture, especially when low virus levels are present, such as may be found in bulls with a PTI. The real-time RT-PCR described here uses a pair of sequence-specific primers for amplification of target <u>D-R</u>NA and a 5'-nuclease oligoprobe for the detection of amplified products. The oligoprobe is a single, sequence-specific oligonucleotide, labelled with two different fluorophores. The primers and probe are available commercially and several different fluorophores options are available. This panpestivirus real-time RT-PCR assay is designed to detect viral <u>D-R</u>NA of all strains of BVDV <u>types 1</u> (*Pestivirus bovis*)-and BVDV, 2 (*Pestivirus tauri*) and 3 (*Pestivirus brazilense*) as well as BDV, CSFV (*Pestivirus suis*), some strains of BDV (*Pestivirus ovis*) and most atypical pestiviruses. The assay selectively amplifies a 208 base pair sequence of the 5' non-translated region (5' NTR) of the pestivirus genome. Details of the primers and probes are given in the protocol outlined below.

- i) Sample preparation, equipment and reagents
 - a) The samples used for the test are, typically, extended bovine semen or occasionally raw semen. If the samples are only being tested by real-time RT-PCR, it is acceptable for them to be submitted chilled but they must still be cold when they reach the laboratory. Otherwise, if a cold chain cannot be assured or if virus isolation is being undertaken, the semen samples should be transported to the laboratory in liquid nitrogen or on dry ice. At the laboratory, the samples should be stored in liquid nitrogen or at lower than -70°C (for long-term storage) or 4°C (for short-term storage of up to 7 days). Note: samples for virus isolation should not be stored at 4°C for more than 1–2 days.
 - b) Due to the very high analytical sensitivity of real-time RT-PCR, much smaller volumes of semen may be used. However, at least three straws (minimum 250 µl each) from each collection batch of semen should be processed. The semen in the three straws should be pooled and mixed thoroughly before taking a sample for nucleic acid extraction.
 - c) A real-time PCR detection system, and the associated data analysis software, is required to perform the assay. A number of real-time PCR detection systems are available from various manufacturers. Other equipment required for the test includes a micro-centrifuge, a chilling block, a micro vortex, and micropipettes. As real-time RT-PCR assays are able to detect very small amounts of target nucleic acid molecules, appropriate measures are required to avoid contamination. , including dedicated and physically separated 'clean' areas for reagent preparation (where no samples or materials used for PCR are handled), a dedicated sample processing area and an isolated area for the PCR thermocycler and associated equipment. Each area should have dedicated reagents and equipment. Furthermore, a minimum of one negative sample should be processed in parallel to monitor the possibility of low level contamination. Sources of contamination may include product carry-over from positive samples or, more commonly, from cross contamination by PCR products from earlier work.
 - d) The real-time RT-PCR assay involves two separate procedures.
 - Firstly, BVDV RNA is extracted from semen using an appropriate validated nucleic acid extraction method. Systems using magnetic beads for the capture and purification of the nucleic acid are recommended. It is also preferable that the beads are handled by a semi-automated magnetic particle handling system.
 - 2) The second procedure is the RT-PCR analysis of the extracted RNA template in a real-time RT-PCR system.

ii) Extraction of RNA

503

504

505 506

507 508

509

510

511

512 513

514

515

516

517

518

519 520

521

522

523

524

525

526 527

528 529

530

531

532

533 534

535

536

537 538

539

540

541

542

543

544

545

546

547

548

549

Report of the Meeting of the Biological Standards Commission / September 2023

550 551 552 553 554 555 556 557 558 559 560		from extra eval syst the gela µl o extra resu	the same action kit uated to e ems and k semen san tin saline (f the dilut action kits ilts are obta	ucleic acid is extracted from the pooled (three straws collected at the same time animal) semen sample. Use of a commercially available magnetic bead based is recommended. However, the preferred kit should be one that has been ensure optimal extraction of difficult samples (semen and whole blood). Some it protocols are sufficiently refined that it is not necessary to remove cells from nple. Prior to extraction dilute the pooled semen sample 1/4 in phosphate buffered PBGS) or a similar buffered solution. Complete the RNA extraction by taking 50 ed, pooled sample and add it to the sample lysis buffer. Some commercial may require the use of a larger volume. It has also been found that satisfactory ained by adding 25 µl of undiluted pooled sample to sample lysis buffer. Complete by following the kit manufacturer's instructions.
561	iii)	Rea	I-time RT-I	PCR assay procedure
562 563 564 565 566		a)	available the real-ti by variou	mixture: There are a number of commercial real-time PCR amplification kits from various sources and the particular kits selected need to be compatible with me PCR platform selected. The required primers and probes can be synthesised is commercial companies. The WOAH Reference Laboratories for BVDV can information on suitable suppliers.
567 568 569 570 571 572 573		b)	as a 2 × 6 for applic nuclease solutions use or lin	nd storage of reagents: The real-time PCR reaction mixture is normally provided concentration ready for use. The manufacturer's instructions should be followed ation and storage. Working stock solutions for primers and probe are made with -free water at the concentration of 20 μ M and 3 μ M, respectively. The stock are stored at –20°C and the probe solution should be kept in the dark. Single-nited use aliquots can be prepared to limit freeze–thawing of primers and probes nd their shelf life.
574		c)	Primers a	and probe sequences
575 576			Selection below.	of the primers and probe are outlined in Hoffmann <i>et al</i> . (2006) and summarised
577			Forward:	BVD 190-F 5'-GRA-GTC-GTC-ART-GGT-TCG-AC
578			Reverse:	V326 5'-TCA-ACT-CCA-TGT-GCC-ATG-TAC
579			Probe:	TQ-pesti 5'-FAM-TGC-YAY-GTG-GAC-GAG-GGC-ATG-C-TAMRA-3'
580		d)	Preparati	on of reaction mixtures
581 582 583 584 585 586 587 588 589			PCR acti included. <u>and</u> two controls a added af volume o system b	R reaction mixtures are prepared in a separate room that is isolated from other vities and sample handling. For each PCR test, appropriate controls should be As a minimum, a no template control (NTC), appropriate negative control (NC) positive controls (PC1, PC2) should be included. The positive and negative are included in all steps of the assay from extraction onwards while the NTC is ter completion of the extraction. The PCR amplifications are carried out in a f 25 μ l. The protocol described is based on use of a 96 well microplate based ut other options using microtubes are also suitable. Each well of the PCR plate bontain 20 μ l of reaction mix and 5 μ l of sample as follows:
590			12.5 µl	2× RT buffer – from a commercial kit.
591			1 µl	BVD 190-F Forward primer (20 μM)
592			1 µl	V326 Reverse primer (20 μM)
593			1 µl	TQ-pesti Probe (3 μM)
594			2 µl	tRNA (40 ng/µl)
595			1.5 µl	nuclease free water
596			1 µl	25× enzyme mix
597			5 µl	sample (or controls – NTC, NC, PC1, PC2)

598	e)	Selection of controls
599		NTC: usually consists of nuclease free water or tRNA in nuclease free water that is added
600		in place of a sample when the PCR reaction is set up.
601		NC: In practice, many laboratories use PBGS or a similar buffer. Ideally the controls for
602		testing of semen samples should be negative semen, from seronegative bulls. However,
603		as a minimum, the assay in use should have been extensively validated with negative and
604		positive samples to confirm that it gives reliable extraction and amplification with semen.
605		PCs: There are two positive controls (PC1=moderate - [Ct 29-32] and PC2=weak [Ct 32-
606		35] positive). Positive semen from naturally infected bulls is preferable as a positive
607 608		control. However, this is likely to be difficult to obtain. Further, semen from a PI bull is not considered suitable because the virus loads are usually very high and would not give a
609		reliable indication of any moderate reduction in extraction or assay performance. Negative
610		semen spiked with defined quantities of BVDV virus could be used as an alternative. If
611		other samples are used as a routine PC, as a minimum the entire extraction process and
612		PCR assay in use must have been extensively validated using known positive semen from
613		bulls with a PTI or from bulls undergoing an acute infection. If these samples are not
614 615		available and spiked samples are used for validation purposes, a number of samples spiked with very low levels of virus should be included. On a day to day basis, the inclusion
616		of an exogenous control with each test sample will largely compensate for not using
617		positive semen as a control and will give additional benefits by monitoring the efficiency of
618		the assay on each individual sample. Positive control samples should be prepared
619		carefully to avoid cross contamination from high titred virus stocks and should be prepared
620		in advance and frozen at a 'ready to use' concentration and ideally 'single use' volume.
621	f)	Extracted samples are added to the PCR mix in a separate room. The controls should be
622		added last, in a consistent sequence in the following order: NTC, negative and then the
623	,	two positive controls.
624	g)	Real-time polymerase chain reaction
625		The PCR plate or tubes are placed in the real-time PCR detection system in a separate,
626 627		designated PCR room. Some mastermixes have uniform reaction conditions that are suitable for many different assays. As an example, the PCR detection system is
628		programmed for the test as follows:
629		1× 48°C 10 minutes
630		1× 95°C 10 minutes
631		45 × (95°C 15 seconds, 60°C 1 minute)
632	h) A	nalysis of real-time PCR data
633	Т	he software program is usually set to automatically adjust results by compensating for any
634	b	ackground signal and the threshold level is usually set according to the manufacturer's
635	ir	nstructions for the selected analysis software used. In this instance, a threshold is set at 0.05.
636		nterpretation of results
637	а) Test controls – all controls should give the expected results with positive controls (PC1
638 639		and PC2) falling within the designated range and both the negative control (NC) and no template control (NTC) should have no Ct values.
640	b) Test samples
641 642		 Positive result: Any sample that has a cycle threshold (Ct) value less than 40 is regarded as positive.
643		2) Negative result: Any sample that shows no Ct value is regarded as negative. However,
644		before reporting a negative result for a sample, the performance of the exogenous
645		internal control should be checked and shown to give a result within the accepted range
646		for that control (for example, a Ct value no more than 2–3 Ct units higher than the
647		NTC).

648 **1.3. Enzyme-linked immunosorbent assay for antigen detection**

649 Antigen detection by ELISA has become a widely adopted method for the detection of individual PI animals. These 650 assays are not intended for the detection of acutely infected animals (though from to time this may be achieved). Importantly, these assays are not designed for screening of semen or biological materials used in assays or vaccine 651 manufacture. Several methods for the ELISA for antigen detection have been published and a number of commercial 652 kits are available. Most are based on the sandwich ELISA principle, with a capture antibody bound to the solid phase, 653 654 and a detector antibody conjugated to a signal system, such as peroxidase. Amplification steps such as the use of 655 biotin and streptavidin in the detection system are sometimes used to increase assay sensitivity. Both monoclonaland polyclonal-based systems are described. The test measures BVD antigen (NS2-3 or ERNS) in lysates of peripheral 656 blood leukocytes; the new generation of antigen-capture ELISAs (ERNS capture ELISAs) are able to detect BVD 657 658 antigen in blood as well as in plasma or serum samples. The best of the methods gives a sensitivity similar to virus isolation, and may be preferred in those rare cases where persistent infection is combined with seropositivity. Due to 659 transient viraemia, the antigen ELISA is less useful for virus detection in acute BVD infections. 660

661The NS2-3 antigen detectionELISAs may be less effective in young calves that have had colostrum due to the662presence of BVDV maternal antibodies, especially when blood samples or blood leucocytes are tested (Fux & Wolf,6632012). Blood or blood leucocytes should not be tested in the first month (ERNs capture ELISA) or the first 3 months664(NS2-3 ELISA) of life due to the inhibitory effect of maternal antibodies. The real-time RT-PCR is probably the most665sensitive detection method for this circumstance, but the ERNS ELISA has also been shown to be a sensitive and666reliable test, particularly when used with skin biopsy (ear-notch) samples (Cornish *et al.*, 2005).

667 **1.4. Immunohistochemistry**

Enzyme-labelled methods are useful to detect BVDV antigen in tissue sections, particularly where suitable MAbs are
 available. However, these assays are not appropriate to certify animals for international trade and use should be
 limited to diagnostic investigations. It is important that the reagents and procedures used be fully validated, and that
 nonspecific reactivity be eliminated. For PI cattle almost any tissue can be used, but particularly good success has
 been found with lymph nodes, thyroid gland, skin, brain, abomasum and placenta. Skin biopsies, such as ear-notch
 samples, have shown to be useful for *in-vivo* diagnosis of persistent B<u>V</u>DV infection.

674 2. Serological tests

675 Antibody to BVDV can be detected in cattle sera by a standard VNT or by ELISA, using one of several published methods or with commercial kits (e.g. Edwards, 1990). Serology is used to identify levels of herd immunity, for the detection of the 676 presence of PI animals in a herd, to assist with investigation of reproductive disease and possible involvement of BVDV 677 and to establish the serological status of bulls being used for semen collection and to identify whether there has been a 678 679 recent infection. ELISA for antibody in bulk milk samples can give a useful indication of the BVD status of a herd (Niskanen, 1993). High ELISA values (0.8 or more absorbance units) in an unvaccinated herd indicates a high probability of the herd 680 681 having been exposed to BVDV in the recent past, most likely through one or more persistently viraemic animals being 682 present. In contrast, a-very low or negative values (≤ 0.2) -indicates that it is unlikely that persistently viraemic animals are present. However, ELISA values are not always a reliable indicator of the presence of PI animals on farms, due to differing 683 husbandry (Zimmer et al., 2002), recent administration of vaccine and also due to the presence of viral antigen in bulk milk, 684 which may interfere with the antibody assay itself. Determination of the antibody status of a small number of young stock 685 (9-18 months) has also been utilised as an indicator of recent transmission of BVDV in the herd (Houe et al., 1995), but 686 687 this approach is also dependent on the degree of contact between different groups of animals in the herd and the potential 688 for exposure from neighbouring herds. VN tests are more frequently used for regulatory purposes (e.g. testing of semen 689 donors) while ELISAs (usually in the form of commercially prepared kits) are commonly used for diagnostic applications. Whether ELISA or VNT, control positive and negative standard sera must be included in every test. These should give 690 results within predetermined limits for the test to be considered valid. In the VNT, a 'serum control' to monitor sample 691 692 toxicity should also be included for each test sample.

693 **2.1. Virus neutralisation test**

694 Selection of the virus strain to include in a VNT is very important. No single strain is likely to be ideal for all 695 circumstances, but in practice one should be selected that detects the highest proportion of serological reactions in the local cattle population. Low levels of antibody to BVD type 2 virus (Pestivirus tauri) may not be detectable by a 696 neutralisation test that uses type 1 strain of the virus, and vice versa (Fulton et al., 1997). It is important that BVD 697 type 1 and BVD type 2 (Pestivirus bovis and tauri) be used in the test and not just the one that the diagnostician 698 thinks is present, as this can lead to under reporting. Because it makes the test easier to read, most laboratories use 699 highly cytopathic, laboratory-adapted strains of BVDV for VN tests. Two widely used cytopathic strains are 'Oregon 700 C24V' and 'NADL'. However immune-labelling techniques are now available that allow simple detection of the growth 701

or neutralisation of non-cytopathic strains where this is considered desirable, especially to support the inclusion of a
 locally relevant virus strain. An outline protocol for a microtitre VN test is given below (Edwards, 1990):

2.1.1. Test procedure

704 705

706

707 708

709

710

711

712

713

714

715

716

717 718

719

720 721

722

723 724

725 726

727

745

746

747

748

749

750

751 752

753

- i) The test sera are heat-inactivated for 30 minutes at 56°C.
- ii) From a starting dilution of 1/4, serial twofold dilutions of the test sera are made in a cell-culture grade flat-bottomed 96-well microtitre plate, using cell culture medium as diluent. For each sample, three or four wells are used at each dilution depending on the degree of precision required. At each dilution of serum, for each sample one well is left without virus to monitor for evidence of sample toxicity that could mimic viral cytopathology or interfere with virus replication. Control positive and negative sera should also be included in each batch of tests.
- An equal volume (e.g. 50 μl) of a stock of cytopathic strain of BVDV containing 100 TCID₅₀ (50% tissue culture infective dose) is added to each well. A back titration of virus stock is also done in some spare wells to check the potency of the virus (acceptance limits 30–300 TCID₅₀).
 - iv) The plate is incubated for 1 hour at 37°C.
 - v) A flask of suitable cells (e.g. bovine turbinate, bovine testis) is trypsinised and the cell concentration is adjusted to 1.5 × 10⁵/ml. 100 µl of the cell suspension is added to each well of the microtitre plate.
 - vi) The plate is incubated at 37°C for 4–5 days, either in a 5% CO₂ atmosphere or with the plate sealed.
- vii) The wells are examined microscopically for CPE or fixed and stained by immunoperoxidase staining using an appropriate monoclonal antibody. The VN titre for each serum is the dilution at which the virus is neutralised in 50% of the wells. This can be calculated by the Spearman– Kärber or Reed Muench methods. A seronegative animal will show no neutralisation at the lowest dilution (1/4), equivalent to a final dilution of 1/8. For accurate comparison of antibody titres, and particularly to demonstrate significant (more than fourfold) changes in titre, samples should be tested in parallel in the same test.

728 2.2. Enzyme-linked immunosorbent assay

Both indirect and blocking types of test can be used. A number of commercial kits are available. As with the virus neutralisation test, ELISAs configured using antigen from one <u>genotype species</u> of BVD<u>V</u> may not efficiently detect antibody induced by another <u>genotype virus species</u>. Tests should therefore be selected for their ability to detect antibody to the spectrum of types and strains circulating in the country where the test is to be performed.

The chief difficulty in setting up the test lies in the preparation of a viral antigen of sufficient potency. The virus must 733 734 be grown under optimal culture conditions using a highly permissive cell type. Any serum used in the medium must not inhibit growth of BVDV. The optimal time for harvest should be determined experimentally for the individual culture 735 system. The virus can be concentrated and purified by density gradient centrifugation. Alternatively, a potent antigen 736 can be prepared by treatment of infected cell cultures with detergents, such as Nonidet P40, N-decanoyl-N-737 738 methylglucamine (Mega 10), Triton X-100 or 1-octylbeta-D-glucopyranoside (OGP). Some workers have used fixed, infected whole cells as antigen. In the future, Increasing use may be is made of artificial antigens manufactured by 739 expressing specific viral genes in bacterial or eukaryotic systems. Such systems should be validated by testing sera 740 specific to a wide range of different virus strains. In the future, this technology should enable the production of 741 serological tests complementary to subunit or marker vaccines, thus enabling differentiation between vaccinated and 742 naturally infected cattle. An example outline protocol for an indirect ELISA is given below (Edwards, 1990). 743

744 2.2.1. Test procedure

- Roller cultures of secondary calf testis cells with a high multiplicity of infection (about one), are inoculated with BVDV strain Oregon C24V, overlaid with serum-free medium and incubated for 24 hours at 37°C.
 - ii) The cells are scraped off and pelleted. The supernatant medium is discarded. The pellet is treated with two volumes of 2% OGP in PBS for 15 minutes at 4°C, and centrifuged to remove the cell debris. The supernatant antigen is stored in small aliquots at -70°C, or freeze-dried. Non-infected cells are processed in parallel to make a control antigen.
- iii) The antigen is diluted to a predetermined dilution in 0.05 M bicarbonate buffer, pH 9.6. Alternate rows of an ELISA-grade microtitre plate are coated with virus and control antigens overnight at

754 4°C. The plates are then washed in PBS with 0.05% Tween 20 or Tween 80 (PBST) before use in the test. 755 756 iv) Test sera are diluted 1/50 in serum diluent (0.5 M NaCl; 0.01 M phosphate buffer; 0.05% Tween 20; 0.001 M ethylene diamine tetra-acetic acid; 1% polyvinyl pyrrolidone, pH 7.2) and added to 757 virus- and control-coated wells for 1 hour at 37°C. The plates are then washed five times in 758 759 PBST. 760 V) Rabbit anti-bovine IgG peroxidase conjugate is added at a predetermined dilution (in serum diluent) for 1 hour at 37°C, then the plates are again washed five times in PBST. 761 A suitable enzyme substrate is added, such as hydrogen peroxide/tetramethyl benzidine. After 762 vi) colour development, the reaction is stopped with sulphuric acid and the absorbance is read on 763 an ELISA plate reader. The value obtained with control antigen is subtracted from the test 764 reaction to give a net absorbance value for each serum. 765 It is recommended to convert net absorbance values to sample:positive ratio (or percentage 766 vii) positivity) by dividing net absorbance by the net absorbance on that test of a standard positive 767 serum that has a net absorbance of about 1.0. This normalisation procedure leads to more 768 769 consistent and reproducible results.

C. REQUIREMENTS FOR VACCINES

771 **1. Background**

770

772 BVDV vaccines are used primarily for disease control purposes. Although they can convey production advantages especially in intensively managed cattle such as in feedlots. In some countries where BVDV eradication is being 773 774 undertaken, PI animals are removed and remaining cattle are vaccinated to maintain a high level of infection antibody positivity and prevent the generation of further PI animals. Vaccination to control BVDV infections can be challenging due 775 in part to the antigenic variability of the virus and the occurrence of persistent infections that arise as a result of fetal 776 777 infection. Ongoing maintenance of the virus in nature is predominantly sustained by PI animals that are the product of inutero infection. The goal for a vaccine should be to prevent systemic viraemia and the virus crossing the placenta. If this 778 is successfully achieved it is likely that the vaccine will prevent the wide range of other clinical manifestations, including 779 reproductive, respiratory and enteric diseases and immunosuppression with its secondary sequelae. There are many 780 different vaccines available in different countries. Traditionally, BVD vaccines fall into two classes: modified live virus or 781 782 inactivated vaccines. Experimental recombinant subunit vaccines based on BVD viral glycoprotein E2 expressed with 783 baculovirus, or transgenic plants or heterologous viruses and BVDV E2 DNA vaccines have been described but few, if any, are in commercial production. They offer a future prospect of 'marker vaccines' when used in connection with a 784 785 complementary serological test.

786 **1.1. Characteristics of a target product profile**

Traditionally, BVD vaccines fall into two classes: modified live or inactivated virus vaccines. The essential requirement 787 for both types is to afford-provide a high level of fetal infection protection. Many of the live vaccines have been based 788 on a cytopathic strain of the virus which is considered to be unable to cross the placenta. However, it is important to 789 ensure that the vaccine virus does not cause fetal infection. In general, vaccination of breeding animals should be 790 completed well before insemination to ensure optimal protection and avoid any risk of fetal infection. Live virus vaccine 791 may also be immunosuppressive and precipitate other infections. On the other hand, modified live virus vaccines may 792 only require a single dose. Use of a live product containing a cytopathic strain of BVDV may precipitate mucosal 793 disease by superinfection of persistently viraemic animals. Properly formulated inactivated vaccines are very safe to 794 use but, to obtain satisfactory levels of immunity, they usually require booster vaccinations, which may be 795 796 inconvenient. A combined vaccination protocol using inactivated followed by live vaccine may reduce the risk of adverse reaction to the live strain. Whether live or inactivated, because of the propensity for antigenic variability, the 797 798 vaccine should contain strains of BVDV that are closely matched to viruses found in the area in which they are used. 799 For example, in countries where strains of BVDV type 2 (Pestivirus tauri) are found, it is important for the vaccine to contain a suitable type 2 strain. For optimal immunity against type 1 strains (Pestivirus bovis), antigens from the 800 801 dominant subtypes (e.g. 1a and 1b) should be included. Due to the need to customise vaccines for the most commonly 802 encountered strains within a country or region, it is not feasible to produce a vaccine antigen bank that can be drawn 803 upon globally.

604 Guidance for the production of veterinary vaccines is given in Chapter 1.1.8 *Principles of veterinary vaccine production.* The guidelines given here and in chapter 1.1.8 are intended to be general in nature and may be supplemented by national and regional requirements.

807 2. Outline of production and minimum requirements for vaccines

808 **2.1. Characteristics of the seed**

814

834

835

836 837

838

839

840

841

842

843

844

845

846 847

848

849 850

851 852

For optimal efficacy, it is considered that there should be a close antigenic match between viruses included in a vaccine and those circulating in the target population. BVDV type 2 strains (<u>Pestivirus tauri</u>) should be included as appropriate. Due to the regional variations in genotypes species and subtypes of BVDV, many vaccines contain more than one strain of BVDV to give acceptable protection. A good appreciation of the antigenic characteristics of individual strains can be obtained by screening with panels of MAbs (Paton *et al.*, 1995).

2.1.1. Biological characteristics of the master seed

- Isolates of cytopathic virus are often mixed with the noncytopathic biotype. The separation and 815 purification of the two biotypes from an initial mixed culture is important to maintain the expected 816 characteristics of the seen seed and depends on several cycles of a limiting dilution technique for the 817 818 noncytopathic virus, or plaque selection for the cytopathic virus. Purity of the cytopathic virus should be confirmed by at least one additional passage at limiting dilution. When isolates have been cloned, 819 their identity and key antigenic characteristics should be confirmed. The identity of the seed virus 820 821 should be confirmed by sequencing. Where there are multiple isolates included in the vaccine, each has to be prepared separately. 822
- 823 While retaining the desirable antigenic characteristics, the strains selected for the seed should not show any signs of disease when susceptible animals are vaccinated. Live attenuated vaccines should 824 not be transmissible to unvaccinated 'in-contact' animals and should not be able to infect the fetus. 825 826 Ideally seeds prepared for the production of inactivated vaccines should grow to high titre to minimise the need to concentrate the antigens and there should be a minimal amount of protein from the cell 827 cultures incorporated into the final product. Master stocks for either live or inactivated vaccines should 828 be prepared under a seed lot system involving master and working stocks that can be used for 829 production in such a manner that the number of passages can be limited and minimise antigenic drift. 830 While there are no absolute criteria for this purpose, as a general guide, the seed used for production 831 should not be passaged more than 20 times beyond the master seed and the master seed should be 832 833 of the lowest passage from the original isolate as is practical.

2.1.2. Quality criteria (sterility, purity, freedom from extraneous agents)

It is crucial to ensure that all materials used in the preparation of the bulk antigens have been extensively screened to ensure freedom from extraneous agents. This should include master and working seeds, the cell cultures and all medium supplements such as bovine serum. It is particularly important to ensure that any serum used that is of bovine origin is free of both adventitious BVDV of all genotypes and antibodies against BVDV strains because low levels of either virus or antibody can mask the presence of the other. Materials and vaccine seeds should be tested for sterility and freedom from contamination with other agents, especially viruses as described in the chapter 1.1.8 and chapter 1.1.9.

2.1.3. Validation as a vaccine strain

All vaccines should pass standard tests for efficacy. Tests should include as a minimum the demonstration of a neutralising antibody response following vaccination, a reduction in virus shedding after challenge in vaccinated cattle and ideally a prevention of viraemia. Efficacy tests of BVD vaccines by assessing clinical parameters in non-pregnant cattle can be limited by the difficulty of consistently establishing clinical signs but, when employed, clinical parameters such as a reduction in the rectal temperature response and leukopenia should be monitored. Although it can be difficult by using virus isolation in cell culture to consistently demonstrate the low levels of viraemia associated with an acute infection, real-time PCR could be considered as an alternative method to establish the levels of circulating virus.

853 If a vaccine passes basic tests, the efficacy of vaccination should ultimately be measured by the capacity to prevent transplacental transmission. If there is a substantial reduction and ideally 854 complete prevention of fetal infection, a vaccine would be expected to be highly effective in other 855 situations (for example prevention of respiratory disease). A suitable challenge system can be 856 857 established by intranasal inoculation of noncytopathic virus into pregnant cows between 60 and 90 days of gestation (Brownlie et al., 1995). Usually this system will reliably produce persistently viraemic 858 offspring in non-immune cows. In countries where BVDV type 2 viruses (Pestivirus tauri) are 859 860 commonly encountered, efficacy in protecting against BVDV type 2 infections should be measured.

2.2. Method of manufacture

2.2.1. Procedure

Both cytopathic and noncytopathic biotypes will grow in a variety of cell cultures of bovine origin. Standard procedures may be used, with the expectation for harvesting noncytopathic virus on days 4–7 and cytopathic virus on days 2–4. The optimal yield of infectious virus will depend on several factors, including the cell culture, isolate used and the initial seeding rate of virus. These factors should be taken into consideration and virus replication kinetics investigated to establish the optimal conditions for large scale virus production. Whether a live or inactivated vaccine, the essential aim will be to produce a high-titred virus stock. This bulk antigen preparation can subsequently be prepared according to the type of vaccine being considered.

2.2.2. Requirements for ingredients

Most BVDV vaccines are grown in cell cultures of bovine origin that are frequently supplemented with medium components of animal origin. The material of greatest concern is bovine serum due to the potential for contamination with BVD viruses and antibodies to these viruses. These adventitious contaminants not only affect the efficiency of production but also may mask the presence of low levels of infectious BVDV that may have undesirable characteristics. In addition to the virus seeds, all materials should be tested for sterility and freedom from contamination with other agents, especially viruses as described in chapters 1.1.8 and 1.1.9. Further, materials of bovine or ovine origin should originate from a country with negligible risk for transmissible spongiform encephalopathies [TSEs] (see chapter 1.1.9).

2.2.3. In-process controls

In-process controls are part of the manufacturing process. Cultures should be inspected regularly to ensure that they remain free from contamination, and to monitor the health of the cells and the development or absence of CPE, as appropriate. While the basic requirement for efficacy is the capacity to induce an acceptable neutralising antibody response, during production, target concentrations of antigen required to achieve an acceptable response may be monitored indirectly by assessment of the quantity of infectious virus or antigen mass that is produced. Rapid diagnostic assays such as the ELISA are useful to monitor BVDV antigen production. Alternatively, the quality of a batch of antigen may be determined by titration of the quantity of infectious virus present, although this may underestimate the quantity of antigen. For inactivated vaccines, infectivity is evaluated before inactivation. For inactivated vaccines the inactivation kinetics should be established so that a suitable safety margin can be determined and incorporated into the routine production processes. At the end of production, *in-vitro* cell culture assays should be undertaken to confirm that inactivation has been complete. These innocuity tests should include a sufficient number of passages and volume of inoculum to ensure that very low levels of infectious virus would be detected if present.

2.2.4. Final product batch tests

i) Sterility

Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in Chapter 1.1.9.

900 ii) Identity

Identity tests should demonstrate that no other strain of BVDV is present when several strains are propagated in a facility producing multivalent vaccines.

903 iii) Safety

Safety tests shall consist of detecting any abnormal local or systemic adverse reactions to the vaccine by all vaccination route(s). Batch-to-batch safety tests are required unless safety of the product is demonstrated and APPROVED in the registration dossier and production is consistent with that described in chapter 1.1.8.

The safety test is different to the inocuity test (see above).

909	Live vaccines must either be demonstrated to be safe in pregnant cattle (i.e. no transmission to the
910	fetus), or should be licensed with a warning not to use them in pregnant animals. Live vaccines
911	containing cytopathic strains should have an appropriate warning of the risk of inducing mucosal
912	disease in PI cattle.

913 iv) Batch potency 914 BVD vaccines must be demonstrated to produce adequate immune responses, when used in their 915 final formulation according to the manufacturer's published instructions. The minimum quantity of 916 infectious virus and/or antigen required to produce an acceptable immune response should be 917 determined. *In-vitro* assays should be used to monitor individual batches during production.

918 2.3. Requirements for authorisation/registration/licensing

2.3.1. Manufacturing process

For registration of a vaccine, all relevant details concerning manufacture of the vaccine and quality control testing should be submitted to the relevant authorities. Unless otherwise specified by the authorities, information should be provided from three consecutive vaccine batches with a volume not less than 1/3 of the typical industrial batch volume.

There is no standard method for the manufacture of a BVD vaccine, but conventional laboratory techniques with stationary, rolled or suspension (micro-carriers) cell cultures may be used. Inactivated vaccines can be prepared by conventional methods, such as binary ethylenimine or beta-propiolactone inactivation (Park & Bolin, 1987). A variety of adjuvants may be used.

928 2.3.2. Safety requirements

In-vivo tests should be undertaken using a single dose, overdose (for live vaccines only) and repeat doses (taking into account the maximum number of doses for primary vaccination and, if appropriate, the first revaccination/booster vaccination) and contain the maximum permitted antigen load and, depending on the formulation of the vaccine, the maximum number of vaccine strains.

933 i) Target and non-target animal safety

The safety of the final product formulation of both live and inactivated vaccines should be assessed in susceptible young calves that are free of maternally derived antibodies and in pregnant cattle. They should be checked for any local reactions following administration, and, in pregnant cattle, for any effects on the unborn calf. Live attenuated vaccines may contribute to immunosuppression that might increase mortality. It may also contribute to the development of mucosal disease in PI animals that is an animal welfare concern. Therefore vaccination of PI animals with live attenuated vaccines containing cytopathic BVDV should be avoided. Live attenuated vaccines must not be capable of being transmitted to other unvaccinated animals that are in close contact.

ii) Reversion-to-virulence for attenuated/live vaccines and environmental considerations

Virus seeds that have been passaged at least up to and preferably beyond the passage limit specified for the seed should be inoculated into young calves to confirm that there is no evidence of disease. If a live attenuated vaccine has been registered for use in pregnant animals, reversion to virulence tests should also include pregnant animals. Live attenuated vaccines should not be transmissible to unvaccinated 'in-contact' animals.

948 iii) Precautions (hazards)

BVDV is not considered to be a human health hazard. Standard good microbiological practice should be adequate for handling the virus in the laboratory. A live virus vaccine should be identified as harmless for people administering the product. However adjuvants included in either live or inactivated vaccines may cause injury to people. Manufacturers should provide adequate warnings that medical advice should be sought in the case of self-injection (including for adjuvants, oil-emulsion vaccine, preservatives, etc.) with warnings included on the product label/leaflet so that the vaccinator is aware of any danger.

2.3.3. Efficacy requirements

The potency of the vaccine should be determined by inoculation into seronegative and virus negative calves, followed by monitoring of the antibody response. Antigen content can be assayed by ELISA and adjusted as required to a standard level for the particular vaccine. Standardised assay protocols applicable to all vaccines do not exist. Live vaccine batches may be assayed by infectivity titration. Each production batch of vaccine should undergo potency and safety testing as batch release criteria. BVD vaccines must be demonstrated to produce adequate immune responses, as outlined above, when used in their final formulation according to the manufacturer's published instructions.

964

969

970

971

972

973

974

975

976

977

978

979

981

982

983

984 985

986

2.3.4. Vaccines permitting a DIVA strategy (detection of infection in vaccinated animals)

965 966 967 968

To date, there are no commercially available vaccines for BVDV that support use of a true DIVA strategy. Experimental subunit vaccines based on baculovirus-expressed BVD viral glycoprotein E2 have been described but are not available commercially. They offer a future prospect of 'marker vaccines' when used in connection with a complementary serological test. Experimental BVDV E2 DNA vaccines and BVDV E2 subunit vaccines expressed using transgenic plants and alphavirus replicon or chimeric pestivirus vaccines have also been described.

2.3.5. Duration of immunity

There are few published data on the duration of antibody following vaccination with a commercial product. Protocols for their use usually recommend a primary course of two inoculations and boosters at yearly intervals. Only limited data are available on the antibody levels that correlate with protection against respiratory infections (Bolin & Ridpath, 1995; Howard *et al.*, 1989) or *in-utero* infection (Brownlie *et al.*, 1995). However, there are many different commercial formulations and these involve a range of adjuvants that may support different periods of efficacy. Consequently, duration of immunity data must be generated separately for each commercially available product by undertaking challenge tests at the end of the period for which immunity has been claimed.

980 **2.3.6. Stability**

There are no accepted guidelines for the stability of BVD vaccines, but it can be assumed that attenuated virus vaccine (freeze-dried) should remain potent for at least 1 year if kept at 4°C. Inactivated virus vaccine could have a longer shelf life at 4°C. Lower temperatures could prolong shelf life for either type, but adjuvants in killed vaccine may preclude this. Bulk antigens that have not been formulated into finished vaccine can be reliably stored frozen at low temperatures but the antigen quality should be monitored with *in-vitro* assays prior to incorporation into a batch of vaccine.

987

REFERENCES

- BAKER J.C. (1995). The clinical manifestations of bovine viral diarrhea infection. *Vet. Clin. North. Am. Food Animal Practice*, **11**, 425–445.
- BAUERMANN F.V., FLORES E.F. & RIDPATH J.F. (2012). Antigenic relationships between bovine viral diarrhea virus 1 and 2
 and HoBi virus: possible impacts on diagnosis and control. *J. Vet. Diagn. Invest.* (official publication of the American Association of Veterinary Laboratory Diagnosticians), 24, 253–261.
- BAUERMANN F.V., HARMON A., FLORES E.F., FALKENBERG S.M., REECY J.M. & RIDPATH J.F. (2013). *In vitro* neutralization of
 HoBi-like viruses by antibodies in serum of cattle immunized with inactivated or modified live vaccines of bovine viral
 diarrhea viruses 1 and 2. *Vet. Microbiol.*, **166**, 242–245.
- BOLIN S.R. & RIDPATH J.F. (1992). Differences in virulence between two noncytopathic bovine viral diarrhea viruses in calves. *Am. J. Vet. Res.*, **53**, 2157–2163.
- BOLIN S.R. & RIDPATH J.F. (1995). Assessment of protection from systemic infection or disease afforded by low to
 intermediate titers of passively acquired neutralizing antibody against bovine viral diarrhea virus in calves. *Am. J. Vet. Res.*,
 56, 755–759.
- 1001 BROWNLIE J. (1985). Clinical aspects of the bovine virus diarrhoea/mucosal disease complex in cattle. *In Practice*, **7**, 195– 1002 202.
- BROWNLIE J. (1990). The pathogenesis of bovine virus diarrhoea virus infections. *Rev. sci. tech. Off. int. Epiz.*, **9**, 43–59.
- BROWNLIE J., CLARKE M.C., HOOPER L.B. & BELL G.D. (1995). Protection of the bovine fetus from bovine viral diarrhoea virus by means of a new inactivated vaccine. *Vet. Rec.*, **137**, 58–62.
- <u>CHEN M., LIU M., LIU S. & SHANG Y. (2021). HoBi-like pestivirus infection leads to bovine death and severe respiratory</u>
 <u>disease in China. *Transbound. Emerg. Dis.*, 68, 1069–1074.</u>

CORNISH T.E., VAN OLPHEN A.L., CAVENDER J.L., EDWARDS J.M., JAEGER P.T., VIEYRA L.L., WOODARD L.F., MILLER D.R. &
 O'TOOLE D. (2005). Comparison of ear notch immunohistochemistry, ear notch antigen-capture ELISA, and buffy coat virus isolation for detection of calves persistently infected with bovine viral diarrhea virus. *J. Vet. Diagn. Invest.*, **17**, 110–117.

- 1011 DUFFELL S.J. & HARKNESS J.W. (1985). Bovine virus diarrhoea-mucosal disease infection in cattle. *Vet. Rec.*, **117**, 240–245.
- 1012 EDWARDS S. (1990). The diagnosis of bovine virus diarrhoea-mucosal disease in cattle. *Rev. sci. tech. Off. int. Epiz.*, **9**, 1013 115–130.
- 1014 FRAY M.D., MANN G.E., BLEACH E.C.L., KNIGHT P.G., CLARKE M.C. & CHARLESTON B. (2002). Modulation of sex hormone 1015 secretion in cows by acute infection with bovine viral diarrhoea virus. *Reproduction*, **123**, 281-289.

FULTON R.W., SALIKI J.T., BURGE L.J., DOFFAY J.M., BOLIN S.R., MAES R.K., BAKER J.C. & FREY M.L. (1997). Neutralizing
 antibodies to type-1 and type-2 bovine viral diarrhea viruses – detection by inhibition of viral cytopathology and infectivity
 by immunoperoxidase assay. *Clin. Diagn. Lab. Immunol.*, **4**, 380–383.

1019 <u>FUX R. & WOLF G. (2012). Transient elimination of circulating bovine viral diarrhoea virus by colostral antibodies in</u> 1020 persistently infected calves: a pitfall for BVDV-eradication programs? *Vet. Microbiol.*, **161**, 1.

- GIVENS M.D., RIDDELL K.P., WALZ P.H., RHOADES J., HARLAND R., ZHANG Y., GALIK P.K., BRODERSEN B.W., COCHRAN A.M.,
 BROCK K.V., CARSON R.L. & STRINGFELLOW D.A. (2007). Noncytopathic bovine viral diarrhea virus can persist in testicular
 tissue after vaccination of peri-pubertal bulls but prevents subsequent infection. *Vaccine*, 25, 867–876.
- 1024 HOFFMANN B., DEPNER K., SCHIRRMEIER H. & BEER M. (2006). A universal heterologous internal control system for duplex 1025 real-time RT-PCR assays used in a detection system for pestiviruses *J. Virol. Meth*ods, **136**, 200–209.
- HOUE H., BAKER J.C., MAES R.K., RUEGG P.L. & LLOYD J.W. (1995). Application of antibody titers against bovine viral diarrhea virus (BVDV) as a measure to detect herds with cattle persistently infected with BVDV. J. Vet. Diagn. Invest., 7, 327–332.
- HOWARD C.J., CLARKE M.C. & BROWNLIE J. (1989). Protection against respiratory infection with bovine virus diarrhoea virus by passively acquired antibody. *Vet. Microbiol.*, **19**, 195–203.
- 1030 LETELLIER C. & KERKHOFS P. (2003). Real-time PCR for simultaneous detection and genotyping of bovine viral diarrhea 1031 virus. *J. Virol. Methods*, **114**, 21–27.
- 1032 MCGOLDRICK A., BENSAUDE E., IBATA G., SHARP G. & PATON D. J. (1999). Closed one-tube reverse transcription nested 1033 polymerase chain reaction for the detection of pestiviral RNA with fluorescent probes. *J. Virol. Methods*, **79**, 85–95.
- 1034 McGowan M.R. & KIRKLAND P.D. (1995). Early reproductive loss due to bovine pestivirus infection. *Br. Vet. J.*, **151**, 263– 1035 270.
- 1036 MEYLING A. (1984). Detection of BVD virus in viraemic cattle by an indirect immunoperoxidase technique. *In:* Recent 1037 Advances in Virus Diagnosis (CEC Seminar), McNulty M.S. & McFerran J.B., eds. Martinus Nijhoff, Belfast, UK, 37–46.
- MOENNIG V. & LIESS B. (1995). Pathogenesis of intrauterine infections with bovine viral diarrhea virus. *Vet. Clin. North. Am.*,
 11, 477–487.
- MOENNIG V., HOUE H. & LINDBERG A. (2005). BVD control in Europe: current status and perspectives. *Anim. Health Res. Rev.*, **6**, 63–74.
- 1042 NISKANEN R. (1993). Relationship between the levels of antibodies to bovine viral diarrhoea virus in bulk tank milk and the 1043 prevalence of cows exposed to the virus. *Vet. Rec.*, **133**, 341–344.
- 1044 PARK B.K. & BOLIN S.R. (1987). Molecular changes of bovine viral diarrhea virus polypeptides treated with binary 1045 ethylenimine, beta-propiolactone and formalin. *Res. Rep. Rural Dev. Admin. (L&V) Korea,* **29**, 99–103.
- PATON D.J., SANDS J.J. LOWINGS J.P., SMITH J.E., IBATA G., EDWARDS S. (1995). A proposed division of the pestivirus genus
 using monoclonal antibodies, supported by cross-neutralisation assays and genetic sequencing. *Vet. Res.*, 26, 92–109.

1048	Schweizer M., Stalder H., Haslebacher A., Grisiger M., Schwermer H. & Di Labio E. (2021). Eradication of bovine viral
1049	diarrhoea (BVD) in cattle in Switzerland: Lessons taught by the complex biology of the virus. Front. Vet. Sci., 8, 702730.
1050	VILCEK S., PATON D.J., DURKOVIC B., STROJNY L., IBATA G., MOUSSA A., LOITSCH A., ROSSMANITH W., VEGA S., SCICLUNA M.T.
1051	& PALFI V. (2001). Bovine viral diarrhoea virus genotype 1 can be separated into at least eleven genetic groups. Arch. Virol.,
1052	146 , 99–115.
1053	VOGES H., HORNER G.W., ROWE S. & WELLENBERG G.J. (1998). Persistent bovine pestivirus infection localized in the testes
1054	of an immuno-competent, non-viremic bull. <i>Vet. Microbiol.</i> , 61 , 165–175.
1055	ZIMMER G., SCHOUSTRA W. & GRAAT E. A.M. (2002). Predictive values of serum and bulk milk sampling for the presence of
1056	persistently infected BVDV carriers in dairy herds. <i>Res. Vet. Sci.</i> , 72 , 75–82.
1057	*
1058	* *
1059	NB: There are WOAH Reference Laboratories for bovine viral diarrhoea (please consult the WOAH Web site:
1060	https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3)
1061	Please contact the WOAH Reference Laboratories for any further information on diagnostic tests, reason to and vessions for boving viral diagraphics
1062	diagnostic tests, reagents and vaccines for bovine viral diarrhoea
1063	NB: First adopted in 1990. Most recent updates adopted in 2015.

1	Annex 12. Item 5.1 – Chapter 3.4.12 Lumpy Skin Disease
2	MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION
3	Paris, 4–8 September 2023
4	
5	CHAPTER 3.4.12.
6	LUMPY SKIN DISEASE

SUMMARY

8 Description of the disease: Lumpy skin disease (LSD) is a poxvirus disease of cattle characterised by 9 fever, nodules on the skin, mucous membranes and internal organs, emaciation, enlarged lymph nodes, oedema of the skin, and sometimes death. The disease is of economic importance as it can cause a 10 11 temporary reduction in milk production, temporary or permanent sterility in bulls, damage to hides and, occasionally, death. Various strains of capripoxvirus are responsible for the disease. These are antigenically 12 indistinguishable from strains causing sheep pox and goat pox yet distinct at the genetic level. LSD has a 13 14 partially different geographical distribution from sheep and goat pox, suggesting that cattle strains of 15 capripoxvirus do not infect and transmit between sheep and goats. Transmission of LSD virus (LSDV) is 16 thought to be predominantly by arthropods, natural contact transmission in the absence of vectors being inefficient. Lumpy skin disease is endemic in most many African and Middle Eastern countries. Between 17 2012 and 2022, LSD spread into south-east Europe, the Balkans, Russia and Asia as part of the Eurasian 18 19 LSD epidemic.

7

- Pathology: the nodules are firm, and may extend to the underlying subcutis and muscle. Acute histological
 key lesions consist of epidermal vacuolar changes with intracytoplasmic inclusion bodies and dermal
 vasculitis. Chronic key histological lesions consist of fibrosis and necrotic sequestrae.
- 23 Detection of the agent: Laboratory confirmation of LSD is most rapid using a real-time or conventional 24 polymerase chain reaction (PCR) method specific for capripoxviruses in combination with a clinical history 25 of a generalised nodular skin disease and enlarged superficial lymph nodes in cattle. Ultrastructurally, 26 capripoxvirus virions are distinct from those of parapoxvirus, which causes bovine papular stomatitis and 27 pseudocowpox, but cannot be distinguished morphologically from orthopoxvirus virions, including cowpox 28 and vaccinia viruses, both of which can cause disease in cattle, although neither causes generalised infection and both are uncommon in cattle. LSDV will grow in tissue culture of bovine, ovine or caprine origin. 29 30 In cell culture, LSDV causes a characteristic cytopathic effect and intracytoplasmic inclusion bodies that is 31 distinct from infection with Bovine herpesvirus 2, which causes pseudo-lumpy skin disease and produces 32 syncytia and intranuclear inclusion bodies in cell culture. Capripoxvirus antigens can be demonstrated in 33 tissue culture using immunoperoxidase or immunofluorescent staining and the virus can be neutralised using 34 specific antisera.
- A variety of conventional and real-time PCR tests as well as isothermal amplification tests using capripoxvirus-specific primers have been published for use on a variety of samples.
- Serological tests: The virus neutralisation test (VNT) and enzyme-linked immunosorbent assays (ELISAs)
 are widely used and have been validated. The agar gel immunodiffusion test and indirect immunofluorescent
 antibody test are less specific than the VNT due to cross-reactions with antibody to other poxviruses.

40 Western blotting using the reaction between the P32 antigen of LSDV with test sera is both sensitive and 41 specific, but is difficult and expensive to carry out.

Requirements for vaccines: All strains of capripoxvirus examined so far, whether derived from cattle, sheep or goats, are antigenically similar. Attenuated cattle strains, and strains derived from sheep and goats have been used as live vaccines against LSDV.

45

A. INTRODUCTION

46 Lumpy skin disease (LSD) was first seen in Zambia in 1929, spreading into Botswana by 1943 (Haig, 1957), and then into South Africa the same year, where it affected over eight million cattle causing major economic loss. In 1957 it entered 47 48 Kenya, at the same time as associated with an outbreak of sheep pox (Weiss, 1968). In 1970 LSD spread north into the 49 Sudan, by 1974 it had spread west as far as Nigeria, and in 1977 was reported from Mauritania, Mali, Ghana and Liberia. Another epizootic of LSD between 1981 and 1986 affected Tanzania, Kenya, Zimbabwe, Somalia and the Cameroon, with 50 reported mortality rates in affected cattle of 20%. The occurrence of LSD north of the Sahara desert and outside the African 51 continent was confirmed for the first time in Egypt and Israel between 1988 and 1989, and was reported again in 2006 52 (Brenner et al., 2006). In the past decade, LSD occurrences have been reported in the Middle Eastern, European and 53 Asian regions (for up-to-date information, consult WOAH WAHIS interface⁴⁰). Lumpy skin disease outbreaks tend to be 54 sporadic, depending upon animal movements, immune status, and wind and rainfall patterns affecting vector populations. 55 The principal method of transmission is thought to be mechanical by various arthropod vectors (Tuppurainen et al., 2015). 56

57 Lumpy skin disease virus (LSDV) belongs to the family Poxviridae, subfamily Chordopoxvirinae Chordopoxviridae, and genus Capripoxvirus. In common with other poxviruses LSDV replicates in the cytoplasm of an infected cell, forming distinct 58 perinuclear viral factories. The LSD virion is large and brick-shaped measuring 293-299nm (length) and 262-273nm 59 (width). The LSDV genome structure is also similar to other poxviruses, consisting of double-stranded linear DNA that is 60 61 25% GC-rich, approximately 150,000 bp in length, and encodes around 156 open reading frames (ORFs). An inverted terminal repeat sequence of 2200-2300 bp is found at each end of the linear genome. The linear ends of the genome are 62 joined with a hairpin loop. The central region of the LSDV genome contains ORFs predicted to encode proteins required 63 64 for virus replication and morphogenesis and exhibit a high degree of similarity with genomes of other mammalian poxviruses. The ORFs in the outer regions of the LSDV genome have lower similarity and likely encode proteins involved 65 66 in viral virulence and host range determinants.

Phylogenetic analysis shows the majority of LSDV strains group into two monophyletic clusters (cluster 1.1 and 1.2) 67 68 (Biswas et al., 2020; Van Schalkwyk et al., 2021). Cluster 1.1 consists of LSDV Neethling vaccine strains that are based on the LSDV/Neethling/LW-1959 vaccine strain (Kara et al., 2003; Van Rooyen et al., 1959; van Schalkwyk et al., 2020) 69 and historic wild-type strains from South Africa. Cluster 1.2 consists of wild-type strains from southern Africa, Kenya, the 70 northern hemisphere, and the Kenvan KSGP O-240 commercial vaccine. In addition to these two clusters, there have 71 72 recently been recombinant LSDV strains isolated from clinical cases of LSD in the field in Russia and central Asia (Flannery et al., 2021; Sprygin et al., 2018; 2020; Wang et al., 2021). These recombinant viruses show unique patterns of accessory 73 74 gene alleles, consisting of sections of both wild-type and "vaccine" LSDV strains.

75 The severity of the clinical signs of LSD is highly variable and depends on a number of factors, including the strain of capripoxvirus the age of the host, immunological status and breed. Bos taurus is generally more susceptible to clinical disease 76 77 than Bos indicus; the Asian buffalo (Bubalus spp.) has also been reported to be susceptible. Within Bos taurus, the fineskinned Channel Island breeds develop more severe disease, with lactating cows appearing to be the most at risk. However, 78 even among groups of cattle of the same breed kept together under the same conditions, there is a large variation in the 79 80 clinical signs presented, ranging from subclinical infection to death (Carn & Kitching, 1995). There may be failure of the virus to infect the whole group, probably depending on the virulence of the virus isolate, immunological status of the host, host 81 genotype, and vector prevalence. Seroprevalence studies, experimental infections and case reports have provided indications 82 that several wildlife species (e.g. springbok, impala, giraffe, camel, banteng) are susceptible to LSDV infection (Dao et al., 83 84 2022; Hedger & Hamblin, 1983; Kumar et al., 2023; Porco et al., 2023). The scarcity of documented outbreaks in wildlife and 85 the fact that available studies remain limited in number and mostly involve only a few animals, make it difficult to determine the role of wildlife in LSDV epidemiology. This topic deserves further study, especially given the current spread of LSDV in 86 new geographical areas where large numbers of naïve, potentially susceptible wild bovines and other ruminants are present. 87

The incubation period under field conditions has not been reported, but following experimental inoculation is 6–9 days until the onset of fever. In the acutely infected animal, there is an initial pyrexia, which may exceed 41°C and persist for 1 week. All the superficial lymph nodes become enlarged. In lactating cattle there is a marked reduction in milk yield. Lesions develop over the body, particularly on the head, neck, udder, scrotum, vulva and perineum between 7 and 19 days after virus inoculation (Coetzer, 2004). The characteristic integumentary lesions are multiple, well circumscribed to coalescing, 0.5–5 cm in diameter, firm, flat-topped papules and nodules. The nodules involve the dermis and epidermis, and may

⁴⁰ https://www.woah.org/en/what-we-do/animal-health-and-welfare/disease-data-collection/

94 extend to the underlying subcutis and occasionally to the adjacent striated muscle. These nodules have a creamy grey to 95 white colour on cut section, which may initially exude serum, but over the ensuing 2 weeks a cone-shaped central core or sequestrum of necrotic material/necrotic plug ("sit-fast") may appear within the nodule. The acute histological lesions 96 consist of epidermal vacuolar changes with intracytoplasmic inclusion bodies and dermal vasculitis. The inclusion bodies 97 98 are numerous, intracytoplasmic, eosinophilic, homogenous to occasionally granular and they may occur in endothelial cells, fibroblasts, macrophages, pericytes, and keratinocytes. The dermal lesions include vasculitis with fibrinoid necrosis, 99 oedema, thrombosis, lymphangitis, dermal-epidermal separation, and mixed inflammatory infiltrate. The chronic lesions 100 are characterised by an infarcted tissue with a sequestered necrotic core, often rimmed by granulation tissue gradually 101 replaced by mature fibrosis. At the appearance of the nodules, the discharge from the eyes and nose becomes 102 mucopurulent, and keratitis may develop. Nodules may also develop in the mucous membranes of the mouth and 103 alimentary tract, particularly the abomasum and in the trachea and the lungs, resulting in primary and secondary 104 pneumonia. The nodules on the mucous membranes of the eyes, nose, mouth, rectum, udder and genitalia quickly 105 ulcerate, and by then all secretions, ocular and nasal discharge and saliva contain LSD virus (LSDV). The limbs may be 106 oedematous and the animal is reluctant to move. Pregnant cattle may abort, and there is a report of intrauterine 107 108 transmission (Rouby & Aboulsoudb, 2016). Bulls may become permanently or temporarily infertile and the virus can be excreted in the semen for prolonged periods (Irons et al., 2005). Recovery from severe infection is slow; the animal is 109 emaciated, may have pneumonia and mastitis, and the necrotic plugs of skin, which may have been subject to fly strike, 110 111 are shed leaving deep holes in the hide (Prozesky & Barnard, 1982).

The main differential diagnosis is pseudo-LSD caused by bovine herpesvirus 2 (BoHV-2). This is usually a milder clinical condition, characterised by superficial nodules, resembling only the early stage of LSD. Intra-nuclear inclusion bodies and viral syncytia are histopathological characteristics of BoHV-2 infection not seen in LSD. Other differential diagnoses (for integumentary lesions) include: dermatophilosis, dermatophytosis, bovine farcy, photosensitisation, actinomycosis, actinobacillosis, urticaria, insect bites, besnoitiosis, nocardiosis, demodicosis, onchocerciasis, pseudo-cowpox, and cowpox. Differential diagnoses for mucosal lesions include: foot and mouth disease, bluetongue, mucosal disease, malignant catarrhal fever, infectious bovine rhinotracheitis, and bovine papular stomatitis.

119 LSDV is not transmissible to humans. However, all laboratory manipulations must be performed at an appropriate 120 containment level determined using biorisk analysis (see Chapter 1.1.4 *Biosafety and biosecurity: Standard for managing* 121 *biological risk in the veterinary laboratory and animal facilities*).

122

123

B. DIAGNOSTIC TECHNIQUES

	Purpose								
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	Prevalence of infection – surveillance	Immune status in individual animals or populations post- vaccination			
Detection of the agent									
Virus isolation	+	++	+	+++	+	-			
PCR	++	+++	++	+++	+	_			
TEM –		-	_	+	_	_			
		Detecti	on of immune r	response					
VNT	++	++	++	++	++	++			
IFAT	+	+	+	+	+	+			
ELISA	++	++	++	++	++	++			

Table 1. Test methods available for the diagnosis of LSD and their purpose

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; – = not appropriate for this purpose.

PCR = polymerase chain reaction; TEM = Transmission electron microscopy; VNT = virus neutralisation test;

IFAT = indirect fluorescent antibody test; ELISA = enzyme-linked immunosorbent assay.

128 **1. Detection of the agent**

129 **1.1.** Specimen collection, submission and preparation

130 Material for virus isolation and antigen detection should be collected as biopsies or from skin nodules at post-mortem examination. Samples for virus isolation should preferably be collected within the first week of the occurrence of 131 clinical signs, before the development of neutralising antibodies (Davies, 1991; Davies et al., 1971), however virus 132 can be isolated from skin nodules for at least 3-4 weeks thereafter. Samples for genome detection using conventional 133 or real-time polymerase chain reaction (PCR) may be collected when neutralising antibody is present. Following the 134 135 first appearance of the skin lesions, the virus can be isolated for up to 35 days and viral nucleic acid can be demonstrated via PCR for up to 3 months (Tuppurainen et al., 2005; Weiss, 1968). Buffy coat from blood collected 136 137 into heparin or EDTA (ethylene diamine tetra-acetic acid) during the viraemic stage of LSD (before generalisation of 138 lesions or within 4 days of generalisation), can also be used for virus isolation. Samples for histology should include the lesion and tissue from the surrounding (non-lesion) area, be a maximum size of 2 cm³, and be placed immediately 139 140 following collection into ten times the sample volume of 10% neutral buffered formal saline.

- 141Tissues in formalin have no special transportation requirements in regard to biorisks.
Blood samples with
anticoagulant for virus isolation from the buffy coat should be placed immediately on ice after gentle mixing and
processed as soon as possible. In practice, the samples may be kept at 4°C for up to 2 days prior to processing, but
should not be frozen or kept at ambient temperatures. Tissues for virus isolation and antigen detection should be kept
at 4°C, on ice or at -20°C. If it is necessary to transport samples over long distances without refrigeration, the medium
should contain 10% glycerol; the samples should be of sufficient size (e.g. 1 g in 10 ml) that the transport medium
does not penetrate the central part of the biopsy, which should be used for virus isolation.
- 148 Samples for histology should include the lesion and tissue from the surrounding (non-lesion) area, be a maximum size of 2 cm³, and be placed immediately following collection into ten times the sample volume of 10% neutral buffered 149 formaldehyde. Tissues in formalin have no special transportation requirements in regard to biorisks. Material for 150 histology should be prepared using standard techniques and stained with haematoxylin and eosin (H&E) (Burdin, 151 152 1959). Lesion material for virus isolation and antigen detection is minced using a sterile scalpel blade and forceps and then macerated in a sterile steel ball-bearing mixer mill, or ground with a pestle in a sterile mortar with sterile 153 sand and an equal volume of sterile phosphate buffered saline (PBS) or serum-free modified Eagle's medium 154 containing sodium penicillin (1000 international units [IU]/ml), streptomycin sulphate (1 mg/ml), mycostatin 155 156 (100 IU/ml) or fungizone (amphotericin, 2.5 µg/ml) and neomycin (200 IU/ml). The suspension is freeze-thawed three times and then partially clarified using a bench centrifuge at 600 g for 10 minutes. In cases where bacterial 157 contamination of the sample is expected (such as when virus is isolated from skin samples), the supernatant can be 158 filtered through a 0.45 µm pore size filter after the centrifugation step. Buffy coats may be prepared from unclotted 159 blood using centrifugation at 600 g for 15 minutes, and the buffy coat carefully removed into 5 ml of cold double-160 distilled water using a sterile Pasteur pipette. After 30 seconds, 5 ml of cold double-strength growth medium is added 161 and mixed. The mixture is centrifuged at 600 g for 15 minutes, the supernatant is discarded and the cell pellet is 162 suspended in 5 ml growth medium, such as Glasgow's modified Eagle's medium (GMEM). After centrifugation at 163 600 g for a further 15 minutes, the resulting pellet is suspended in 5 ml of fresh GMEM. Alternatively, the buffy coat 164 165 may be separated from a heparinised sample by using a Ficoll gradient.

166 **1.2. Virus isolation on cell culture**

167 LSDV will grow in tissue culture of bovine, ovine or caprine origin. MDBK (Madin–Darby bovine kidney) cells are often 168 used, as they support good growth of the virus and are well characterised (Fay et al., 2020). Primary cells, such as 169 lamb testis (LT) cells also support viral growth, but care needs to be taken to ensure they are not contaminated with viruses such as bovine viral diarrhoea virus. One ml of clarified supernatant or buffy coat is inoculated onto a confluent 170 monolayer in a 25 cm² culture flask at 37°C and allowed to adsorb for 1 hour. The culture is then washed with warm 171 172 PBS and covered with 10 ml of a suitable medium, such as GMEM, containing antibiotics and 2% fetal calf serum. If available, tissue culture tubes containing appropriate cells and a flying cover-slip, or tissue culture microscope slides, 173 174 are also infected.

175 The flasks/tissue culture tubes are examined daily for 7-14 days for evidence of cytopathic effects (CPE). Infected cells develop a characteristic CPE consisting of retraction of the cell membrane from surrounding cells, and eventually 176 rounding of cells and margination of the nuclear chromatin. At first only small areas of CPE can be seen, sometimes 177 as soon as 2 days after infection; over the following 4-6 days these expand to involve the whole cell monolaver-sheet. 178 If no CPE is apparent by day 14, the culture should be freeze-thawed three times, and clarified supernatant inoculated 179 on to a fresh cell monolayer. At the first sign of CPE in the flasks, or earlier if a number of infected cover-slips are 180 being used, a cover-slip should be removed, fixed in acetone and stained using H&E. Eosinophilic intracytoplasmic 181 inclusion bodies, which are variable in size but up to half the size of the nucleus and surrounded by a clear halo, are 182 diagnostic for poxvirus infection. PCR may be used as an alternative to H&E for confirmation of the diagnosis. The 183

- 184 CPE can be prevented or delayed by adding specific anti-LSDV serum to the medium. In contrast, the herpesvirus 185 that causes pseudo-LSD produces a Cowdry type A intranuclear inclusion body. It also forms syncytia.
- 186 An ovine testis cell line (OA3.Te) has been evaluated for the propagation of capripoxvirus isolates (Babiuk *et al.*, 187 2007), however this cell line has been found to be contaminated with pestivirus and should be used with caution.

188 **1.3.** Polymerase chain reaction (PCR)

189 The conventional gel-based PCR method described below is a simple, fast and sensitive method for the detection of 190 capripoxvirus genome in EDTA blood, semen or tissue culture samples (Tuppurainen *et al.*, 2005).

1.3.1. Test procedure

191

192 193

194

195

196

197

198

199

200

201 202

203

204

205 206

207

208 209

210 211

212 213

214

215

216

217

218

219

220 221

222 223

224

225

226

227

228 229

230 231

- The extraction method described below can be replaced using commercially available DNA extraction kits.
- Freeze and thaw 200 µl of blood in EDTA, semen or tissue culture supernatant and suspend in 100 µl of lysis buffer containing 5 M guanidine thiocyanate, 50 mM potassium chloride, 10 mM Tris/HCl (pH 8); and 0.5 ml Tween 20.
- Cut skin and other tissue samples into fine pieces using a sterile scalpel blade and forceps. Grind with a pestle in a mortar. Suspend the tissue samples in 800 μl of the above mentioned lysis buffer.
- iii) Add 2 µl of proteinase K (20 mg/ml) to blood samples and 10 µl of proteinase K (20 mg/ml) to tissue samples. Incubate at 56°C for 2 hours or overnight, followed by heating at 100°C for 10 minutes. Add phenol:chloroform:isoamylalcohol (25:24:1 [v/v]) to the samples in a 1:1 ratio. Vortex and incubate at room temperature for 10 minutes. Centrifuge the samples at 16,060 *g* for 15 minutes at 4°C. Carefully collect the upper, aqueous phase (up to 200 µl) and transfer into a clean 2.0 ml tube. Add two volumes of ice cold ethanol (100%) and 1/10 volume of 3 M sodium acetate (pH 5.3). Place the samples at -20°C for 1 hour. Centrifuge again at 16,060 *g* for 15 minutes at 4°C and discard the supernatant. Wash the pellets with ice cold 70% ethanol (100 µl) and centrifuge at 16,060 *g* for 1 minute at 4°C. Discard the supernatant and dry the pellets thoroughly. Suspend the pellets in 30 µl of nuclease-free water and store immediately at -20°C (Tuppurainen *et al.*, 2005). Alternatively a column-based extraction kit may be used.
- iv) The primers for this PCR assay were developed from the gene encoding the viral attachment protein. The size of the expected amplicon is 192 bp (Ireland & Binepal, 1998). The primers have the following gene sequences:

Forward primer 5'-TCC-GAG-CTC-TTT-CCT-GAT-TTT-TCT-TAC-TAT-3'

Reverse primer 5'-TAT-GGT-ACC-TAA-ATT-ATA-TAC-GTA-AAT-AAC-3'.

- v) DNA amplification is carried out in a final volume of 50 µl containing: 5 µl of 10 × PCR buffer, 1.5 µl of MgCl₂ (50 mM), 1 µl of dNTP (10 mM), 1 µl of forward primer, 1 µl of reverse primer, 1 µl of DNA template (~10 ng), 0.5 µl of *Taq* DNA polymerase and 39 µl of nuclease-free water. The volume of DNA template required may vary and the volume of nuclease-free water must be adjusted to the final volume of 50 µl.
 - vi) Run the samples in a thermal cycler as follows: 2 minutes at 95°C; then 45 seconds at 95°C, 50 seconds at 50°C and 1 minute at 72°C (34 cycles); 2 minutes at 72°C and hold at 4°C until analysis.
- vii) Mix 10 µl of each sample with loading dye and load onto a 1.5% agarose gel in TAE buffer (Tris/acetate buffer containing EDTA). Load a parallel lane with a 100 bp DNA-marker ladder. Electrophoretically separate the products using approximately 8–10 V/cm for 40–60 minutes and visualise with a suitable DNA stain and transilluminator.

Quantitative real-time PCR methods have been described that are reported to be faster and have higher sensitivity than conventional PCRs (Balinsky *et al.*, 2008; Bowden *et al.*, 2008). A real-time PCR method that differentiates between LSDV, sheep pox virus and goat pox virus has been published (Lamien *et al.*, 2011).

232Quantitative real-time PCR assays have been designed to differentiate between Neethling-based233LSDV strains, which are often used for vaccination, and wild-type LSDV strains from cluster 1.2234(Agianniotaki *et al.*, 2017; Pestova *et al.*, 2018; Vidanovic *et al.*, 2016). These "DIVA" assays (DIVA:235differentiation of infected from vaccinated animals) enable, for example, differentiation of "Neethling

response" caused by vaccination with a LSDV Neethling vaccine strain from disease caused by infection with a cluster 1.2 wild-type virus. However these DIVA PCR assays cannot distinguish between a LSDV Neethling vaccine strain and the novel recombinant LSDV strains recently-isolated from disease outbreaks in Asia (Byadovskaya *et al.*, 2021; Flannery *et al.*, 2021). These DIVA assays are also not capable of discriminating between LSDV Neethling vaccine strains and recently characterised (historic) wild-type viruses from South Africa belonging within cluster 1.1 (Van Schalkwyk *et al.*, 2020; 2021). Consequently, in regions where recombinant strains (currently Asia and possibly elsewhere) or wild-type cluster 1.1 strains are circulating (currently South Africa and possibly elsewhere), these DIVA assays are not suitable for distinguishing vaccine and wild-type virus. Thus, in order to overcome these constraints, whole genome sequencing is recommended.

1.4. Transmission electron microscopy

236

237

238

239

240 241

242

243 244

245

246

250 251

252

253

254

255

256 257

258

259

The characteristic poxvirus virion can be visualised using a negative staining preparation technique followed by examination with an electron microscope. There are many different negative staining protocols, an example of which is given below.

1.4.1. Test procedure

Before centrifugation, material from the original biopsy suspension is prepared for examination under the transmission electron microscope by floating a 400-mesh hexagonal electron microscope grid, with pioloform-carbon substrate activated by glow discharge in pentylamine vapour, onto a drop of the suspension placed on parafilm or a wax plate. After 1 minute, the grid is transferred to a drop of Tris/EDTA buffer, pH 7.8, for 20 seconds and then to a drop of 1% phosphotungstic acid, pH 7.2, for 10 seconds. The grid is drained using filter paper, air-dried and placed in the electron microscope. The capripox virion is brick shaped, covered in short tubular elements and measures approximately 290 × 270 nm. A host-cell-derived membrane may surround some of the virions, and as many as possible should be examined to confirm their appearance (Kitching & Smale, 1986).

The capripox-virions of capripoxvirus are indistinguishable from those of orthopoxvirus, but, apart 260 261 from vaccinia virus and cowpox virus, which are both uncommon in cattle and do not cause generalised infection, no other orthopoxvirus is known to cause lesions in cattle. However, vaccinia 262 263 virus may cause generalised infection in young immunocompromised calves. In contrast, orthopoxviruses are a common cause of skin disease in domestic buffalo (Bubalus bubalis) causing 264 265 buffalo pox, a disease that usually manifests as pock lesions on the teats, but may cause skin lesions 266 at other sites, such as the perineum, the medial aspects of the thighs and the head. Orthopoxviruses that cause buffalo pox cannot be readily distinguished from capripoxvirus by electron microscopy. 267 268 The virions of parapoxvirus virions that cause bovine papular stomatitis and pseudocowpox are smaller, oval in shape and each is covered in a single continuous tubular element that appears as 269 striations over the virion. Capripoxvirus virions are also distinct from the herpesvirus that causes 270 pseudo-LSD (also known as "Allerton" or bovine herpes mammillitis). 271

1.5. Fluorescent antibody tests

Capripoxvirus antigen can be identified on infected cover-slips or tissue culture slides using fluorescent antibody tests. Cover-slips or slides should be washed and air-dried and fixed in cold acetone for 10 minutes. The indirect test using immune cattle sera is subject to high background colour and nonspecific reactions. However, a direct conjugate can be prepared from sera from convalescent cattle (or from sheep or goats convalescing from capripox) or from rabbits hyperimmunised with purified capripoxvirus. Uninfected tissue culture should be included as a negative control as cross-reactions can cause problems due to antibodies to cellular components (pre-absorption of these from the immune serum helps solve this issue).

280 **1.6. Immunohistochemistry**

Immunohistochemistry using F80G5 monoclonal antibody specific for capripoxvirus ORF 057 has been described for
 detection of LSDV antigen in the skin of experimentally infected cattle (Babiuk *et al.,* 2008).

1.7. Isothermal genome amplification

284 Molecular tests using loop-mediated isothermal amplification to detect capripoxvirus genomes are reported to provide 285 sensitivity and specificity similar to real-time PCR with a simpler method and lower cost (Das *et al.*, 2012; Murray *et al.*, 2013). Field validation of the Das *et al.* method was reported by Omoga *et al.* (2016).

287 2. Serological tests

298

299

300 301

302

303

304

305 306

307

308

309 310

311

312

313 314

315

316

317

318 319

320

321

322 323

324

325

326

327

328

329 330

331

332 333

334

All the viruses in the genus *Capripoxvirus* share a common major antigen for neutralising antibodies and it is thus not possible to distinguish strains of capripoxvirus from cattle, sheep or goats using serological techniques.

290 **2.1. Virus neutralisation**

A test serum can either be titrated against a constant titre of capripoxvirus (100 TCID_{50} [50% tissue culture infective dose]) or a standard virus strain can be titrated against a constant dilution of test serum in order to calculate a neutralisation index. Because of the variable sensitivity of tissue culture to capripoxvirus, and the consequent difficulty of ensuring the accurate and repeatable seeding of 100 TCID_{50} /well, the neutralisation index is the preferred method in most laboratories, although it does require a larger volume of test sera. The test is described using 96-well flatbottomed tissue-culture grade microtitre plates, but it can be performed equally well in tissue culture tubes with the appropriate changes to the volumes used, although it is more difficult to read an end-point in tubes.

2.1.1. Test procedure

- Test sera, including a negative and a positive control, are diluted 1/5 in Eagle's/HEPES (N-2hydroxyethylpiperazine, N-2-ethanesulphonic acid) buffer and inactivated at 56°C for 30 minutes.
 - ii) Next, 50 µl of the first inactivated serum is added to columns 1 and 2, rows A to H of the microtitre plate. The second serum is placed in columns 3 and 4, the third in columns 5 and 6, the positive control serum is placed in columns 7 and 8, the negative control serum is placed in columns 9 and 10, and 50 µl of Eagle's/HEPES buffer (without serum) is placed in columns 11 and 12, and to all wells in row H.
 - iii) A reference strain of capripoxvirus, usually a vaccine strain known to grow well in tissue culture, with a titre of over log₁₀ 6 TCID₅₀ per ml is diluted in Eagle's/HEPES in bijoux bottles to give a log dilution series of log₁₀ 5.0, 4.0, 3.5, 3.0, 2.5, 2.0, 1.5 TCID₅₀ per ml (equivalent to log₁₀ 3.7, 2.7, 2.2, 1.7, 1.2, 0.7, 0.2 TCID₅₀ per 50 μl).
 - iv) Starting with row G and the most diluted virus preparation, 50 μl of virus is added to each well in that row. This is repeated with each virus dilution, the highest titre virus dilution being placed in row A.
 - v) The plates are covered and incubated for 1 hour at 37°C.
 - vi) An appropriate cell suspension (such as MDBK cells) is prepared from pregrown monolayers as a suspension of 10⁵ cells/ml in Eagle's medium containing antibiotics and 2% fetal calf serum. Following incubation of the microtitre plates, 100 µl of cell suspension is added to all the wells, except wells H11 and H12, which serve as control wells for the medium. The remaining wells of row H are cell and serum controls.
 - vii) The microtitre plates are covered and incubated at 37°C for 9 days.
 - viii) Using an inverted microscope, the monolayers are examined daily from day 4 for evidence of CPE. There should be no CPE in the cells of row H. Using the 0240 KSGP vaccine strain of capripoxvirus, by way of example, the final reading is taken on day 9, and the titre of virus in each duplicate titration is calculated using the Kärber method. If left longer, there is invariably a 'breakthrough' of virus in which virus that was at first neutralised appears to disassociate from the antibody.
 - ix) Interpretation of the results: The neutralisation index is the log titre difference between the titre of the virus in the negative serum and in the test serum. An index of ≥1.5 is positive. The test can be made more sensitive if serum from the same animal is examined before and after infection. Because the immunity to capripoxviruses is predominantly cell mediated, a negative result, particularly following vaccination, after which the antibody response may be low, does not imply that the animal from which the serum was taken is not protected.
- Antibodies to capripoxvirus can be detected from 1 to 2 days after the onset of clinical signs. These remain detectable for about 7 months.

335 2.2. Enzyme-linked immunosorbent assay

Enzyme-linked immunosorbent assays (ELISAs) for the detection of capripoxviral antibodies are widely used and are available in commercial kit form (Milovanovic *et al.*, 2019; Samojlovic *et al.*, 2019).

2.3. Indirect fluorescent antibody test

Capripoxvirus-infected tissue culture grown on cover-slips or tissue culture microscope slides can be used for the indirect fluorescent antibody test. Uninfected tissue culture control, and positive and negative control sera, should be included in the test. The infected and control cultures are fixed in acetone at -20°C for 10 minutes and stored at 4°C. Dilutions of test sera are made in PBS, starting at 1/20 or 1/40, and positive samples are identified using an antibovine gamma-globulin conjugated with fluorescein isothiocyanate. Antibody titres may exceed 1/1000 after infection. Sera may be screened at 1/50 and 1/500. Cross-reactions can occur with orf virus (contagious pustular dermatitis virus of sheep), bovine papular stomatitis virus and perhaps other poxviruses.

346 **2.4. Western blot analysis**

347 Western blotting of test sera against capripoxvirus-infected cell lysate provides a sensitive and specific system for 348 the detection of antibody to capripoxvirus structural proteins, although the test is expensive and difficult to carry out.

- Capripoxvirus-infected LT cells should be harvested when 90% CPE is observed, freeze-thawed three times, and 349 the cellular debris pelleted using centrifugation. The supernatant should be decanted, and the proteins should be 350 351 separated using SDS/PAGE (sodium dodecyl sulphate/polyacrylamide gel electrophoresis). A vertical discontinuous 352 gel system, using a stacking gel made up of acrylamide (5%) in Tris (125 mM), pH 6.8, and SDS (0.1%), and a resolving gel made up of acrylamide (10-12.5%) in Tris (560 mM), pH 8.7, and SDS (0.1%), is recommended for use 353 with a glycine running buffer containing Tris (250 mM), glycine (2 M), and SDS (0.1%). Samples of supernatant should 354 be prepared by boiling for 5 minutes with an appropriate lysis buffer prior to loading. Alternatively, purified virus or 355 recombinant antigens may replace tissue-culture-derived antigen. 356
- Molecular weight markers should be run concurrently with the protein samples. The separated proteins in the 357 358 SDS/PAGE gel should be transferred electrophoretically to a nitrocellulose membrane (NCM). After transfer, the NCM is rinsed thoroughly in PBS and blocked in 3% bovine serum albumin (BSA) in PBS, or 5% skimmed milk powder in 359 PBS, on a rotating shaker at 4 C overnight. The NCM can then be separated into strips by employing a commercial 360 361 apparatus to allow the concurrent testing of multiple serum samples, or may be cut into strips and each strip incubated 362 separately thereafter. The NCM is washed thoroughly with five changes of PBS for 5 minutes on a rotating shaker, and then incubated at room temperature on the shaker for 1.5 hours, with the appropriate serum at a dilution of 1/50 363 in blocking buffer (3% BSA and 0.05% Tween 20 in PBS; or 5% milk powder and 0.05% Tween 20 in PBS). The 364 membrane is again thoroughly washed and incubated (in blocking buffer) with anti-species immunoglobulin 365 366 horseradish-peroxidase-conjugated immunoglobulins at a dilution determined using titration. After further incubation at room temperature for 1.5 hours, the membrane is washed and a solution of diaminobenzidine tetrahydrochloride 367 (10 mg in 50 ml of 50 mM-mm Tris/HCl, pH 7.5, and 20 µl of 30% [v/v] hydrogen peroxide) is added. Incubation is 368 369 then undertaken for approximately 3-7 minutes at room temperature on a shaker with constant observation, and the 370 reaction is stopped by washing the NCM in PBS before excessive background colour is seen. A positive and negative 371 control serum should be used on each occasion.
- Positive test samples and the positive control will produce a pattern consistent with reaction to proteins of molecular weights 67, 32, 26, 19 and 17 kDa – the major structural proteins of capripoxvirus – whereas negative serum samples will not react with all these proteins. Hyperimmune serum prepared against parapoxvirus (bovine papular stomatitis or pseudocowpox virus) will react with some of the capripoxvirus proteins, but not the 32 kDa protein that is specific for capripoxvirus.
- 377

C. REQUIREMENTS FOR VACCINES

1. Background: rationale and intended use of the product

Live attenuated strains of capripoxvirus have been used as vaccines specifically for the control of LSD (Brenner *et al.*,
 2006; Capstick & Coakley, 1961; Carn, 1993). Capripoxviruses are cross reactive within the genus. Consequently, it is
 possible to protect cattle against LSD using strains of capripoxvirus derived from sheep or goats (Coakley & Capstick,
 1961). However, it is recommended to carry out controlled trials, using the most susceptible breeds, prior to introducing a
 vaccine strain not usually used in cattle. The duration of protection provided by LSD vaccination is unknown.

384 Capripoxvirus vaccine strains can produce a large local reaction at the site of inoculation in *Bos taurus* breeds (Davies, 385 1991), which some stock owners find unacceptable. This has discouraged the use of vaccine, even though the 386 consequences of an outbreak of LSD are invariably more severe. Risk-benefit of vaccination should be assessed following 387 stakeholder discussion. 388 <u>Vaccines are a key tool to control LSD. Different types of LSD vaccines have been developed and several are commercially</u>
 389 <u>available (Tuppurainen *et al., 2021).*</u>

390 Live attenuated vaccines (LAV) based on the Neethling LSDV strain (homologous LAV vaccines) have been shown to offer high levels of protection against LSD under experimental conditions (Haegeman et al., 2021) and have been used 391 392 successfully to control the disease in the field, through systematic vaccination of the entire country's cattle population for a number of consecutive years (Klement et al., 2020). Homologous vaccines may induce fever, produce a local reaction 393 394 at the site of inoculation, cause a temporary reduction in milk production and on rare occasions induce a 'Neethling' 395 response (Ben-Gera et al., 2015; Davies, 1991; Haegeman et al., 2021). Such adverse effects, however, usually resolve within a few days and are largely outweighed by the overall benefits of vaccination with homologous vaccines. The duration 396 of immunity induced by good quality live attenuated LSDV vaccines was shown to be at least 18 months (Haegeman et 397 398 al., 2023).

399 As capripox viruses provide cross-reactive protection within the genus, heterologous LAVs comprising sheep pox virus or 400 goat pox virus strains have also been tested and used to protect cattle against LSD. Sheep pox virus-based heterologous vaccines usually contain higher doses of virus than when administered to sheep. Although safe, their effectiveness in 401 402 protecting cattle against LSD is inferior compared to homologous vaccines (Ben-Gera et al., 2015; Zhugunissov et al., 2020). Heterologous vaccines containing goat pox virus strains for use in cattle against LSD have been developed more 403 404 recently. One such vaccine based on the Gorgan strain provided protection under experimental conditions comparable to 405 homologous vaccines (Gari et al., 2015). On the other hand, a goat pox vaccine based on an attenuated Uttarkashi goat 406 pox virus strain performed suboptimally under field conditions in India (Naveem et al., 2023), indicating that further research 407 is warranted before asserting that all goat pox virus-based vaccines induce protection equal to homologous vaccines in 408 cattle against LSD.

In addition, homologous inactivated vaccines against LSD have been developed and tested (Haegeman *et al.*, 2023; Hamdi
 et al., 2020; Wolff *et al.*, 2022). These vaccines are reported to be safe and efficacious. They however require a booster
 vaccination one month after primo-vaccination and then every 6 months thereafter, based on the fact that the duration of
 immunity is shorter than 1 year (Haegeman *et al.*, 2023).

413 <u>None of the commercial vaccines currently available has practical DIVA capacity. This problem may be resolved in the</u>
 414 <u>future by introducing new types of vaccines (e.g. vector-vaccines, subunit vaccines, mRNA vaccines) that are at various</u>
 415 <u>stages of development and evaluation.</u>

416 2. Outline of production of LSD vaccines and minimum requirements for conventional 417 vaccines

418 General requirements set for the facilities used for the production of vaccines and for the documentation and record keeping 419 throughout the whole manufacturing process are described in Chapter 1.1.8 *Principles of veterinary vaccine production.* 420 The documentation should include standard operating procedures (SOP) for the method of manufacture and each step for 421 the testing of cells and reagents used in the process, each batch and the final product.

422 <u>The production of vaccines, including LSD vaccines, starts within research and development (R&D) facilities where vaccine</u> 423 <u>candidates are produced and tested in preclinical studies to demonstrate the quality, safety and efficacy of the product.</u>

Minimum requirements for different production stages of veterinary vaccines are available in different chapters of the
 Terrestrial Manual. These are intended to be used in combination with country-specific regulatory requirements for vaccine
 production and release. Here we outline the most important requirements for the production of live and inactivated LSD
 vaccines. Full requirements are available in Chapter 1.1.8 *Principles of veterinary vaccine production,* Chapter 2.3.3
 Minimum requirements for the organisation and management of a vaccine manufacturing facility and Chapter 2.3.4
 Minimum requirements for the production and quality control of vaccine, and other regulatory documentation.

430 <u>2.1. Quality assurance</u>

431Facilities for manufacturing LSD vaccines should operate in line with the concepts of good laboratory practice (GLP)432and good manufacturing practice (GMP) to produce high quality products. Quality risk management and quality433control with adequate documentation management, as an integral part of the production process, have to be in place.434In case some activities of the production process are outsourced, those should also be appropriately defined.435recorded and controlled.

436The vaccine production process (Outline of Production) should be documented in a series of standard operating437procedures (SOPs), or other documents describing the manufacturing of each batch and the final product (including438starting materials to be used, manufacturing steps, in-process controls and controls on the final product). Detailed439requirements for documentation management in the process of vaccine production are available in Chapter 2.3.3.

A completed Outline of Production is to be enclosed in a vaccine candidate dossier and used for the evaluation of
 the production process and product by regulatory bodies.

442 **2.2. Process validation**

455

443The dossier with the enclosed Outline of Production for the vaccine candidate has to be submitted for regulatory444approval, so it can be assessed and authorised by the competent authority to ensure compliance with local regulatory445requirements. Among others, data on quality, safety, and efficacy will be assessed. The procedures necessary to446obtain these data are described in the subsequent sections.

447 <u>National regulatory authorities might also require official control authority re-testing (check testing) of final products</u>
 448 <u>and batches in government laboratories or an independent batch quality control by a third party.</u>

449 3. Requirements for LSD vaccine candidates and batch production

450 <u>3.1. Requirements for starting materials</u>

- 451 <u>Live attenuated vaccines (LAV) and inactivated vaccines (IV) for LSD are produced using the system of limited and</u> 452 <u>controlled passages of master seed and working seed virus and cell banks with a specified maximum. This approach</u> 453 <u>aims to prevent possible and unwanted drift of properties of seed virus and cells that might arise from repeated</u> 454 <u>passaging.</u>
 - 3.1.1. Characteristics of the seed virus

456 457 458 459	Each seed strain of capripoxvirus used for vaccine production must be accompanied by records clearly and accurately describing its origin, isolation and tissue culture or animal passage history. Preferably, the species and strain of capripoxvirus are characterised using PCR or DNA sequencing techniques.
460 461 462	A quantity of master seed vaccine virus should be prepared, frozen or desiccated and stored at low temperatures such as -80°C and used to produce a consistent working seed for regular vaccine production.
463 464 465	Each master seed strain must be non-transmissible, remain attenuated after further tissue culture passage, and provide complete protection against challenge with virulent field strains for a minimum of 1 year. It must produce a minimal clinical reaction in cattle when given via the recommended route.
466	The necessary safety and potency tests are described in Section C.2.2.4 Final product batch tests.
467 2 .	1.2. Quality criteria (sterility, purity, freedom from extraneous agents)
468 469 470	Each master seed must be tested to ensure its identity and shown to be free from adventitious viruses, in particular pestiviruses, such as border disease and bovine viral diarrhoea virus, and free from contamination with bacteria, fungi or mycoplasmas.
471 472	The general procedures for sterility or purity tests are described in Chapter 1.1.9 Tests for sterility and freedom from contamination of biological materials intended for veterinary use.
473 474 475 476 477 478 479	Master seed virus is a quantity of virus of uniform composition derived from an original isolate, passaged for a documented number of times and distributed into containers at one time and stored adequately to ensure stability (via freezing or lyophilisation). Selection of master seed viruses (MSVs) should ideally be based on their ease of growth in cell culture, virus yield, and in accordance with the regional epidemiological importance. Also, measures to minimise transmissible spongiform encephalopathies (TSE) contamination should be taken into account (see Section C.3.5.1 <i>Purity tests</i>).
480 481	For each seed strain selected for LSD vaccine production, the following information should be provided:
482 483	- <u>Historical record: geographical origin, animal species from which the virus was recovered,</u> isolation procedure, tissue culture or animal passage history
484	- Identity: species and strain identification using DNA sequencing

485 486		- <u>Purity: the absence of bacteria, fungi, mycoplasma, and other viruses (see Chapter 1.1.9 Tests</u> for sterility and freedom from contamination of biological materials intended for veterinary use)
480 487		- Safety (overdose, one/repeated dose tests, and reversion to virulence tests) (see Section C.3.3
488		Vaccine safety)
489		- Efficacy data, linked to a specified (protective) dose (see Section C.3.4 Vaccine efficacy)
490		- <u>Stability</u>
491		Each master seed strain selected for production of live attenuated LSD vaccines must remain
492		attenuated after further passage in animals (see Section C.3.3. Vaccine safety), produce minimal
493		clinical reaction when given via the recommended route, provide complete protection against
494		challenge with virulent field strains, and is ideally not transmissible.
495		A quantity of master seed virus should be prepared and stored to be further used for the preparation
496		of working seeds and production seeds. Working seed viruses may be expanded in one or more (but,
497 498		limited) cell culture passages from the master seed stock and used to produce vaccine batches. This approach and limitation of seed virus passaging will assist in maintaining uniformity and consistency
498 499		in production.
499		
500	<u>3.1.2</u>	. Master cell stocks
501		The production process of LSD vaccines ideally employs an established master cell stock (MCS)
502		system with defined lowest and highest cell passage to be used to grow the vaccine virus. Primary
503		cells derived from normal tissues can be used in the production process, but the use of primary cells
504 505		has an inherently higher risk of introducing extraneous agents compared with the use of established (well characterised) cell lines and should be avoided where alternative methods of producing effective
505 506		vaccines exist. For each MCS, manufacturers should demonstrate:
000		
507		- <u>MCS identity</u>
508		- genetic stability by subculturing from the lowest to the highest passage used for production
509		- <u>stable MCS karyotype with a low level of polyploidy</u>
510 511		 <u>freedom from oncogenicity or tumorigenicity by using <i>in-vivo</i> studies using the highest cell passage that may be used for production</u>
512		- purity of MCSs from extraneous bacteria, fungi, mycoplasma, and viruses
513		- implemented measures to lower TSE contamination risk (see Section C.3.5.1 Purity tests).
514	3.2. Meth	nod of <u>vaccine</u> manufactur <u>ing</u>
515	The method	of manufacture should be documented as the Outline of Production.
516	<u>2.2.1</u>	- Procedure
517	<u>3.2.1</u>	<u>. LSD vaccine batch production</u>
518		Vaccine batches are produced on an appropriate cell line such as MDBK. As already mentioned in
519		the first paragraphs of Section C, all steps undertaken in the production of vaccine batches should
520		be described and documented in the Outline of Production. The production of LAV and IV against
521		LSD starts with the inoculation of the required number of working vials of seed virus is-reconstituted
522		with GMEM or other in appropriate medium and inoculated onto a suitable primary or continuous cell
523		line grown in suspension or monolayer. Cells should be harvested after 1–8 days when they exhibit
524 505		50-70% CPE for maximum in the exponential growth phase. At the time highest viral infectivity, or
525 526		earlier if CPE is extensive and cells appear ready to detach. Techniques such as loads are present, sonication or repeated freeze-thawing are is used to release the intracellular virus from the
520 527		cytoplasm. The lysate may then be clarified <u>using centrifugation</u> to remove cellular debris-(for
528		example by use of centrifugation at 600 g for 20 minutes, with retention of the supernatant). A second
529		passage of the virus may be required to produce sufficient virus for a production batch.
530		An aliquot of the virus suspension is titrated to check the virus titre. For LAV, the virus-containing
531		suspension is diluted to attain the dose at which the vaccine candidate will be evaluated or to at least
532		the determined protective dose for approved vaccines and is then mixed with a suitable protectant
533		such as an equal volume of sterile, chilled 5% lactalbumin hydrolysate and 10% sucrose (dissolved

534 535 536	in double-distilled water or appropriate balanced salt solution), and transferred to individually numbered labelled bottles or bags for storage at low temperatures such as -80°C, or for freeze- drying. A written record of all the procedures followed must be kept for all vaccine batches.
537	2.2.2. Requirements for substrates and media
538 539 540 541	The specification and source of all ingredients used in the manufacturing procedure should be documented and the freedom of extraneous agents (bacteria, fungi, mycoplasma and viruses) should be tested. The detailed testing procedure is described in Chapter 1.1.9. The use of antibiotics must meet the requirements of the licensing authority.
542	2 .2.3. In-process control
543	i) Cells
544 545 546 547 548	ii) Records of the source of the master cell stocks should be maintained. The highest and lowest passage numbers of the cells that can be used for vaccine production must be indicated in the Outline of the Production. The use of a continuous cell line (such as MDBK, etc.) is strongly recommended, unless the virus strain only grows on primary cells. The key advantage of continuous over primary cell lines is that there is less risk of introduction of extraneous agents.
549	iii) Serum
550 551 552	iv) Serum used in the growth or maintenance medium must be free from antibodies to capripoxvirus and free from contamination with pestivirus or other viruses, extraneous bacteria, mycoplasma or fungi.
553	v) Medium
554	vi) Media must be sterile before use.
555	vii) Virus
556 557 558 559 560	viii) Seed virus and final vaccine must be titrated and pass the minimum release titre set by the manufacturer. For example, the minimum recommended field dose of the South African Neethling strain vaccines (Mathijs <i>et al.,</i> 2016) is log ₁₀ 3.5 TCID ₅₀ , although the minimum protective dose is log ₁₀ 2.0 TCID ₅₀ . Capripoxvirus is highly susceptible to inactivation by sunlight and allowance should be made for loss of activity in the field.
561 562 563	ix) The recommended field dose of the Romanian sheep pox vaccine for cattle is log ₁₀ 2.5 sheep infective doses (SID ₅₀), and the recommended dose for cattle of the RM65 adapted strain of Romanian sheep pox vaccine is log ₁₀ 3 TCID ₅₀ (Coakley & Capstick, 1961).
564	3.2.2. Inactivation process for inactivated LSD vaccines
565 566 567 568 569 570 571	Unlike LAV, inactivated LSD vaccines contain inactivated antigens in combination with adjuvants to strengthen the induced immune response after administration. The vaccine evaluation process described below needs to show the amount of antigen necessary to elicit a protective immune response. Currently, literature data indicate that an inactivated vaccine originating from an LSDV virus stock with titre 10 ⁴ cell culture infectious dose ₅₀ (CCID ₅₀)/ml before inactivation can be sufficient to induce an efficient immune response to prevent clinical disease, viremia and virus shedding after challenge of young cattle (Wolf <i>et al.</i> , 2022)
572 573 574 575 576 577 578 579 580	To monitor the inactivation process and the level of antigen inactivation, samples are taken at regular intervals during inactivation and titrated. Inactivation conditions and the length of initial and repeated exposure should be documented in detail since one or more factors during the process could influence the outcomes. The inactivation kinetics should reach a predefined target e.g. one remaining infectious unit per million doses (1×10^{-6} infectious units/dose) as suggested by APHIS (2013). The confirmatory testing of inactivation is performed on each vaccine lot and represents an important part of the inactivation process monitoring. In addition to all the procedures mentioned above, the inactivation procedure and tests demonstrating that antigen inactivation is complete and consistent must additionally be documented in the Outline of Production.

3.3. Vaccine safety 581

- 582 During the vaccine development process, vaccine safety must be evaluated in the target animal (target animal batch 583 safety test -TABST) to demonstrate the safety of the dose intended for registration. The animals used in the safety 584 testing should be representative (species, age and category [calves, heifers, bulls, cows.]) for all the animals for which 585 the vaccine is intended. Vaccinated and control groups are appropriately acclimatised, housed and managed in line with animal welfare standards. Animal suffering has to be eliminated or reduced and euthanasia is recommended in 586 587 moribund animals.
- 588 Essential parameters to be evaluated in safety studies are local and systemic reactions to vaccination, including local 589 reactions at the site of administration, fever, effect on milk production, and induction of a 'Neethling' response. The 590 effect of the vaccine on reproduction needs to be evaluated where applicable.
- 591 A part of the safety evaluation of LAV and IV can be performed during the efficacy trials (see Section C.3.4 Vaccine 592 efficacy) by measuring local and systemic responses following vaccination and before challenge.
- Guidelines for safety evaluation are provided by the European Medicine Agency (EMEA) in VICH GL44: TABST for 593 594 LAV and IV (EMEA, 2009). Safety aspects of LAV and IV against LSD to be evaluated are:

Overdose test for LAV 3.3.1.

Local and systemic responses should be measured following an overdose test whereby 10× the maximum vaccine titre is administered. If the maximum vaccine titre is not specified, 10× the minimum vaccine titre can be applied in multiple injection sites. Ideally, the 10× dose is dissolved in the 1× dose volume of the adjuvants or diluent. Generally, eight animals per group should be used (EMEA, 2009).

<u>3.3.2.</u> One dose and repeat dose test

- This aims to test the safety of the vaccine dose applied in the vaccination regime intended for registration. LAV LSD vaccines require one dose per year, while inactivated LSD vaccines require a booster dose in addition to the primary dose. The minimal recommended interval between administrations is 14 days.
 - Generally, eight animals per group should be used unless otherwise justified (EMEA, 2009). For each target species, the most sensitive breed, age and sex proposed on the label should be used. Seronegative animals should be used. In cases where seronegative animals are not reasonably available, alternatives should be justified.

3.3.3Reversion to virulence tests

- Live attenuated vaccines inherently carry the risk of vaccine virus reverting to virulence when repeated passages in a host species could occur due to shedding and transmission from vaccinated animals to contact animals. LAV LSD vaccines should therefore be tested for non-reversion to virulence by means of passage studies. Vaccine virus (MSV, not the finished vaccine) is inoculated in a group of target animals of susceptible age via the natural route of infection or the route that is most likely to result in infection. The vaccine virus is subsequently recovered from tissues or excretions and is used directly to inoculate a further group of animals. After not less than four passages (see chapter 1.1.8), i.e. use of a total of five groups of animals, the re-isolate must be fully characterised, using the same procedures used to characterise the master seed virus.
- 3.3.4. 620
- 621

595

596

597 598

599

600

601

602 603

604

605

606

607

608

609

610

611

612

613 614

615

616 617

618 619

622

623 624

625 626

Environmental consideration

- This includes the evaluation of the ability of LAV LSD vaccines to be shed, to spread and to infect contact target and non-target animals, and to persist in the environment.

2.2.4. Final product batch tests

- i) Sterility/purity
 - Vaccine samples must be tested for sterility/purity. Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in Chapter 1.1.9.
- 627 iii) Safety and efficacy

600	
628	iv) The efficacy and safety studies should be demonstrated using statistically valid vaccination-
629	challenge studies using seronegative young LSDV susceptible dairy cattle breeds. The group
630	numbers recommended here can be varied if statistically justified. Fifteen cattle are placed in a
631	high containment level large animal unit and serum samples are collected. Five randomly
632	chosen vials of the freeze dried vaccine are reconstituted in sterile PBS and pooled. Two cattle
633	are inoculated with 10 times the recommended field dose of the vaccine, and eight cattle are
634	inoculated with the recommended field dose. The remaining five cattle are unvaccinated control
635	animals. The animals are clinically examined daily and rectal temperatures are recorded. On
636	day 21 after vaccination, the animals are again serum sampled and challenged with a known
637	virulent capripoxvirus strain. The challenge virus solution should also be tested free from
638	extraneous viruses. The clinical response is recorded during the following 14 days. Animals in
639	the unvaccinated control group should develop the typical clinical signs of LSD, whereas there
640	should be no local or systemic reaction in the vaccinates other than a raised area in the skin at
641	the site of vaccination, which should disappear after 4 days. Serum samples are again collected
642	on day 30 after vaccination. The day 21 serum samples are examined for seroconversion to
• •=	
643	selected viral diseases that could have contaminated the vaccine, and the days 0 and
644	30 samples are compared to confirm the absence of antibody to pestivirus. Because of the
645	variable response in cattle to LSD challenge, generalised disease may not be seen in all of the
646	unvaccinated control animals, although there should be a large local reaction.
040	
647	v) Once the efficacy of the particular strain being used for vaccine production has been determined
648	in terms of minimum dose required to provide immunity, it is not necessary to repeat this on the
649	final product of each batch, provided the titre of virus present has been ascertained.
650	vi) Batch potency
651	vii) Potency tests in cattle must be undertaken for vaccine strains of capripoxvirus if the minimum
652	immunising dose is not known. This is usually carried out by comparing the titre of a virulent
653	challenge virus on the flanks of vaccinated and control animals. Following vaccination, the
654	flanks of at least three animals and three controls are shaved of hair. Log ₁₀ dilutions of the
655	challenge virus are prepared in sterile PBS and six dilutions are inoculated intradermally (0.1 ml
656	per inoculum) along the length of the flank; four replicates of each dilution are inoculated down
657	the flank. An oedematous swelling will develop at possibly all 24 inoculation sites on the control
658	animals, although preferably there will be little or no reaction at the four sites of the most dilute
659	inocula. The vaccinated animals may develop an initial hypersensitivity reaction at sites of
660	inoculation within 24 hours, which should quickly subside. Small areas of necrosis may develop
661	
	at the inoculation site of the most concentrated challenge virus. The titre of the challenge virus
662	
	is calculated for the vaccinated and control animals; a difference in titre >log ₁₀ 2.5 is taken as
662	
662	is calculated for the vaccinated and control animals; a difference in titre >log ₁₀ 2.5 is taken as
662 663	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u>
662 663 664	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u>
662 663 664 665	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u> Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for
662 663 664 665 666	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u> <u>Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the</u>
662 663 664 665 666 667	 is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u> <u>Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be</u>
662 663 664 665 666	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u> <u>Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the</u>
662 663 664 665 666 667	 is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. <u>3.4. Vaccine efficacy</u> <u>Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be</u>
662 663 664 665 666 667 668	 is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine
662 663 664 665 666 667 668 669	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production.
662 663 664 665 666 667 668 669 670	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age
662 663 664 665 666 667 668 669 670 671	 is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for
662 663 664 665 666 667 668 669 670	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age
662 663 664 665 666 667 668 669 670 671	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens.
662 663 664 665 666 667 668 669 670 671 672	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens.
662 663 664 665 666 667 668 669 670 671 672 673	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if
662 663 664 665 666 667 668 669 670 671 672 673 674	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two.
662 663 664 665 666 667 668 669 670 671 672 673	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if
662 663 664 665 666 667 668 669 670 671 672 673 674 675	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups:
662 663 664 665 666 667 668 669 670 671 672 673 674	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) – animals inoculated with the vaccine dose and route intended for
662 663 664 665 666 667 668 669 670 671 672 673 674 675	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) – animals inoculated with the vaccine dose and route intended for
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677	is calculated for the vaccinated and control animals; a difference in titre >log10-2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) – animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) – animals inoculated with the vaccine dose and route intended for
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677	is calculated for the vaccinated and control animals; a difference in titre >log10-2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) – animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678	is calculated for the vaccinated and control animals; a difference in titre >log₁₀ 2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) – animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14 days).
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679	 is calculated for the vaccinated and control animals; a difference in titre >log to 2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (<i>n</i>=8) - animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14 days). control group (<i>n</i>=5) - non-vaccinated animals
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679 680	 is calculated for the vaccinated and control animals; a difference in titre >log₁₀ -2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (<i>n</i>=8) – animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14 days). control group (<i>n</i>=5) – non-vaccinated animals
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679 680 681	 is calculated for the vaccinated and control animals; a difference in titre >log 10-2.5 is taken as evidence of protection. 3.4. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (n=8) - animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14 days). control group (n=5) - non-vaccinated animals
662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679 680	 is calculated for the vaccinated and control animals; a difference in titre >log₁₀ -2.5 is taken as evidence of protection. 34. Vaccine efficacy Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for each vaccination regimen that is described in the product label recommendation. This includes studies regarding the onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine production as specified in the Outline of Production. Efficacy (and safety) should be demonstrated in vaccination-challenge studies using representative (by species, age and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for standard viral pathogens. An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two groups: single/repeated dose test group (<i>n</i>=8) – animals inoculated with the vaccine dose and route intended for registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14 days). control group (<i>n</i>=5) – non-vaccinated animals

683 684 685	strain. The challenge virus solution should be of known titre and tested free from extraneous viruses. Experience obtained from previous animal experiments indicates that a dose of challenge virus between 10 ^{4.0} and 10 ^{6.5} TCID ₅₀ produces clinical disease in about half of the susceptible experimental cattle (;.Tuppurainen <i>et al.</i> , 2021).
686 687 688 689 690	The clinical response following challenge is recorded over a period of 14 days. No clinical signs should occur in the vaccinates, other than a local reaction at the site of inoculation. At least 1 animal in the unvaccinated control group should develop the typical clinical signs of LSD. Although a generalised disease with skin nodules may not be seen in all the unvaccinated control animals based on the knowledge that the outcome of a LSDV infection can range from inapparent to severe, at the very least a large local reaction is to be expected.
691 692 693	<u>Clinical and laboratory results will enable assessment of the safety and efficacy of the LSD vaccine candidate and the induced immune responses. Serum samples collected at different time points during the trial can be examined to study seroconversion against selected viral diseases that could have contaminated the vaccine.</u>
694	2.3. Requirements for regulatory approval
695	2.3.1. Safety requirements
696	i) Target and non-target animal safety
697 698 699	ii) The vaccine must be safe to use in all breeds of cattle for which it is intended, including young and pregnant animals. It must also be non-transmissible and remain attenuated after further tissue culture passage.
700 701	iii) Safety tests should be carried out on the final product of each batch as described in Section C.2.2.1.
702	iv) Reversion to virulence for attenuated/live vaccines
703 704	v) The selected final vaccine should not revert to virulence during further passages in target animals.
705	vi) Environmental consideration
706 707	vii) Attenuated vaccine should not be able to perpetuate autonomously in a cattle population. Strains of LSDV are not a hazard to human health.
708	2.3.2. Efficacy requirements
709	i) For animal production
710 711 712 713 714 715 716 717 718	The efficacy of the vaccine must be demonstrated in statistically valid vaccination challenge experiments under laboratory conditions. The group numbers recommended here can be varied if statistically justified. Fifteen cattle are placed in a high containment level large animal unit and serum samples are collected. Five randomly chosen vials of the freeze-dried vaccine are reconstituted in sterile PBS and pooled. Two cattle are inoculated with 10 times the field dose of the vaccine, eight cattle are inoculated with the recommended field dose. The remaining five cattle are unvaccinated control animals. The animals are clinically examined daily and rectal temperatures are recorded. On day 21 after vaccination, the animals are again serum sampled and challenged with a known virulent capripoxvirus strain using intravenous and intradermal inoculation (the challenge virus solution).
719 720 721 722 723 724 725 726 727	should also be tested and shown to be free from extraneous viruses). The clinical response is recorded during the following 14 days. Animals in the unvaccinated control group should develop the typical clinical signs of LSD, whereas there should be no local or systemic reaction in the vaccinates other than a raised area in the skin at the site of vaccination which should disappear after 4 days. Serum samples are again collected on day 30 after vaccination. The day 21 serum samples are examined for seroconversion to selected viral diseases that could have contaminated the vaccine, and the days 0 and 30 samples are compared to confirm the absence of antibody to pestivirus. Because of the variable response in cattle to challenge with LSDV, generalised disease may not be seen in all of the unvaccinated control animals, although there should be a large local reaction.
728 729 730	Once the potency of the particular strain being used for vaccine production has been determined in terms of minimum dose required to provide immunity, it is not necessary to repeat this on the final product of each batch, provided the titre of virus present has been ascertained.

131	in) - r or control and oracleation
732	Vaccination is the only effective way to control LSD outbreaks in endemic countries and recent
733	experiences of the disease in Eastern Europe and the Balkans suggests this is also true for outbreaks
734	in non-endemic countries. Unfortunately, currently no marker vaccines allowing a DIVA strategy are
735	available, although to a limited extent PCR can be used for certain vaccines.
736	The duration of immunity produced by LSDV vaccine strains is currently unknown.
737	2.3.3. Stability
151	
738	All vaccines are initially given a shelf life of 24 months before expiry. Real-time stability studies are
739	then conducted to confirm the appropriateness of the expiry date. Multiple batches of the vaccine
740	should be re-titrated periodically throughout the shelf-life period to determine the vaccine stability.
741	Properly freeze dried preparations of LSDV vaccine, particularly those that include a protectant, such
742	as sucrose and lactalbumin hydrolysate, are stable for over 25 years when stored at
743	-20°C and for 2-4 years when stored at 4°C. There is evidence that they are stable at higher
744	temperatures, but no long term controlled experiments have been reported. No preservatives other
745	than a protectant, such as sucrose and lactalbumin hydrolysate, are required for the freeze-dried
746	preparation.
747	3.5. Batch/serial tests before release for distribution
-	
748	Quality tests on MSV and safety and efficacy tests on vaccine candidates are performed during the evaluation process
749	for new LSD vaccines. Once vaccines are approved to be used in the field, it remains important to verify the quality
750	of each vaccine batch produced. An independent batch quality control assessment may be warranted or requested
751	by national or international regulatory authorities.
752	<u>3.5.1. Purity test</u>
850	Durity is defined by the observes of different contentions. (In the static functions of the
753	Purity is defined by the absence of different contaminants (bacteria, fungi, mycoplasma, and other
754	viruses; see full details in chapter 1.1.9) in the vaccine and its associated diluent/adjuvants. Virus
755	isolation and bacterial culture tests can be used to show freedom from live competent replicating
756	microorganisms, but molecular methods are more rapid and sensitive, but positives can be caused
757	by genome fragments and incompetent replicating microorganisms.
758	Besides the contaminants mentioned above, manufacturers should demonstrate implemented
759	measures to minimise the risk of TSE contamination in ingredients of animal origin such as:
760	- all ingredients of animal origin in production facilities are from countries recognized as having the lowest
760	 all ingredients of animal origin in production facilities are from countries recognised as having the lowest
761	possible risk of bovine spongiform encephalopathy
762	- tissues or other substances used are themselves recognised as being of low or nil risk of containing TSE
763	agents
100	
764	2.5.2 Identity tests
764	3.5.2. Identity tests
765	In addition to identity tests performed on the MSV, the identity tests on final batches aim to
766	demonstrate the presence of only the selected capripoxvirus species and strain in the vaccine as
767	indicated in the Outline of Production and the absence of other strains or members of the genus and
768	any other viral contaminant that might arise during the production process. Identity testing could be
769	assured by using appropriate tests (e.g. PCRs, sanger sequencing, NGS).
770	3.5.3. Potency tests
774	Standard requirements for notancy tasts can be found in CEP. Title 0 part 112 in the European
771	Standard requirements for potency tests can be found in CFR Title 9 part 113, in the European
772	Pharmacopoeia, and in this Terrestrial Manual.
773	<u>3.5.3.1. Live vaccines</u>
	The notanov of LAV against LCD can be measured by means of the first The time films
	The potency of LAV against LSD can be measured by means of virus titration. The virus titre
774	
775	must, as a rule, be sufficiently greater than that shown to be protective in the efficacy test
775 776	must, as a rule, be sufficiently greater than that shown to be protective in the efficacy test for the vaccine candidate. This will ensure that at any time prior to the expiry date, the titre
775	must, as a rule, be sufficiently greater than that shown to be protective in the efficacy test
775 776	must, as a rule, be sufficiently greater than that shown to be protective in the efficacy test for the vaccine candidate. This will ensure that at any time prior to the expiry date, the titre

ii) For control and eradication

731

778		commercial homologous LSD vaccines range between 10 ³ and 10 ⁴ infectious units/dose
779		(Tuppurainen et al., 2021).
780		3.5.3.2. Inactivated LSD vaccines
781		For inactivated LSD vaccines, potency tests are performed using vaccination-challenge
782		efficacy studies in animal hosts (see Section C.3.4. Vaccine efficacy).
783	<u>3.5.4.</u>	<u>Safety/efficacy</u>
784		Safety and efficacy testing is undertaken during the evaluation process of the vaccine candidate, and
785		also needs to be performed on a number of vaccine batches until robust data are generated in line
786		with international and national regulations. Afterwards, when using a seed lot system in combination
787		with strict implementation of GMP standards and depending on local regulations, TABST could be
788		waived as described in VICH50 and VICH55, providing the titer has been ascertained using potency
789		testing. Batches or serials are considered satisfactory if local and systemic reactions to vaccination
790		are in line with those described in the dossier of the vaccine candidate and product literature.
791		3.5.4.1. Field safety/efficacy tests
792		Field testing of two or more batches should be performed on all animal categories for which
793		the product is indicated before release of the product for general use (see chapter 1.1.8).
794		The aim of these studies is to demonstrate the safety and efficacy of the product under
795		normal field conditions of animal care and use in different geographical locations where
796		different factors may influence product performance. A protocol for safety/efficacy testing in
797		the field has to be developed with defined observation and recording procedures. However,
798		it is generally more difficult to obtain statistically significant data to demonstrate efficacy
799		<u>under field conditions. Even when properly designed, field efficacy studies may be</u>
800		inconclusive due to uncontrollable outside influences.
801		3.5.4.2. Duration of Immunity
802		The duration of immunity (DOI) following vaccination should be demonstrated via challenge
803		or the use of a validated serology test. Efficacy testing at the end of the claimed period of
804		protection should be conducted in each species for which the vaccine is indicated or the
805		manufacturer should indicate that the DOI for that species is not known. Likewise, the
806		manufacturer should demonstrate the effectiveness of the recommended booster regime in
807		line with these guidelines, usually by measuring the magnitude and kinetics of the
808		serological response observed.

809 3. Vaccines based on biotechnology

A new generation of capripox vaccines is being developed that uses the LSDV as a vector for the expression and delivery
 of immuno-protective proteins of other ruminant pathogens with the potential for providing dual protection (Boshra *et al.*,
 2013; Wallace & Viljoen, 2005), as well as targeting putative immunomodulatory genes for inducing improved immune
 responses (Kara *et al.*, 2018).

814 4. Post-market studies

815 <u>4.1. Stability</u>

Stability testing shall be carried out as specified in Annex II of Regulation (EU) 2019/6 and in the Ph. Eur. 0062:
 Vaccines for veterinary use, on not fewer than three representative batches providing this mimics the full-scale
 production described in the application. At the end of shelf-life, sterility has to be re-evaluated using sterility testing
 or by showing container closure integrity. Multiple batches of the vaccine should be re-titrated periodically throughout
 the shelf-life period to determine the vaccine stability.

821 <u>4.2. Post-marketing surveillance</u>

After release of a vaccine, its performance under field conditions should continue to be monitored by competent authorities and by the manufacturer itself. Not all listed adverse effects may show up in the clinical trials performed to assess safety and efficacy of the vaccine candidate due to the limited number of animals used. Post-marketing surveillance studies can also provide information on vaccine efficacy when used in normal practice and husbandry conditions, on duration of induced immunity, on ecotoxicity, etc. 832

827

828

829

830

831

REFERENCES

First, a reliable reporting system should be in place to collect consumer complaints and notifications of adverse

AGIANNIOTAKI E.I., CHAINTOUTIS S.C., HAEGEMAN A., TASIOUDI K.E., DE LEEUW I., KATSOULOS P.D., SACHPATZIDIS A., DE CLERCQ
 K., ALEXANDROPOULOS T., POLIZOPOULOU Z.S., CHONDROKOUKI E.D. & DOVAS C.I. (2017). Development and validation of a
 TaqMan probe-based real-time PCR method for the differentiation of wild type lumpy skin disease virus from vaccine virus
 strains. J. Virol. Methods, 249, 48–57.

- BABIUK S., BOWDEN T.R., PARKYN G., DALMAN B., MANNING L., NEUFELD J., EMBURY-HYATT C., COPPS J. & BOYLE D.B. (2008).
 Quantification of lumpy skin disease virus following experimental infection in cattle. *Transbound. Emerg. Dis.*, 55, 299–307.
- BABIUK S., PARKYN G., COPPS J., LARENCE J.E., SABARA M.I., BOWDEN T.R., BOYLE D.B. & KITCHING R.P. (2007). Evaluation
 of an ovine testis cell line (OA3.Ts) for propagation of capripoxvirus isolates and development of an immunostaining
 technique for viral plaque visualization. *J. Vet. Diagn.Invest.*, **19**, 486–491.
- 843 <u>BALINSKY C.A, DELHON G, SMOLIGA G, PRARAT M, FRENCH R.A, GEARY S.J, ROCK D.L</u> & <u>RODRIGUEZ L.L</u>. (2008). Rapid 844 preclinical detection of sheep pox virus by a real-time PCR assay. *J. Clin. Microbiol.*, **46**, 438–442.
- BEN-GERA J., KLEMENT E., KHINICH E, STRAM Y. & SHPIGEL N. (2015). Comparison of the efficacy of Neethling lumpy skin disease virus and x10RM65 sheep-pox live attenuated vaccines for the prevention of lumpy skin disease – The results of a randomized controlled field study. *Vaccine*, 33, 4837–4842.
- BISWAS S., NOYCE R.S., BABIUK L.A., LUNG O., BULACH D. M., BOWDEN T.R., BOYLE D. B., BABIUK S. & EVANS D.H. (2020).
 Extended sequencing of vaccine and wild-type capripoxvirus isolates provides insights into genes modulating virulence and host range. *Transbound. Emerg. Dis.*, 67, 80–97.
- BOSHRA H., TRUONG T., NFON C., GERDTS V., TIKOO S., BABIUK L.A., KARA P., MATHER A., WALLACE D. & BABIUK S. (2013).
 Capripoxvirus vectored vaccines against livestock diseases in Africa. *Antiviral Res.*, 98, 217–227.
- BOWDEN, T.R, BABIUK S.L, PARKYN G.R., COPPS J.S. & BOYLE D.B. (2008). Capripoxvirus tissue tropism and shedding: A quantitative study in experimentally infected sheep and goats. *Virology*, **371**, 380–393.
- BRENNER J., HAIMOVITZ M., ORON E., STRAM Y., FRIDGUT O., BUMBAROV V., KUZNETZOVA L., OVED Z., WASERMAN A., GARAZZI
 S., PERL S., LAHAV D., EDERY N. & YADIN H. (2006). Lumpy skin sease (LSD) in a large dairy herd in Israel. *Isr. J. Vet. Med.*,
 61, 73–77.
- BURDIN M.L. (1959). The use of histopathological examination of skin material for the diagnosis of lumpy skin disease in Kenya. *Bull. Epizoot. Dis. Afr.*, **7**, 27–36.
- BYADOVSKAYA O., PESTOVA Y., KONONOV A., SHUMILOVA I., KONONOVA S., NESTEROV A., BABIUK S. & SPRYCIN A. (2021).
 Performance of the currently available DIVA real-time PCR assays in classical and recombinant lumpy skin disease
 viruses. Transbound. Emerg. Dis., 68, 3020-3024. doi: 10.1111/tbed.13942. Epub 2021 May 13. PMID: 33253485
- CAPSTICK P.B. & COAKLEY W. (1961). Protection of cattle against lumpy skin disease. Trials with a vaccine against Neethling
 type infection. Res. Vet. Sci., 2, 362–368
- 865 CARN V.M. (1993). Control of capripoxvirus infections. Vaccine, 11, 1275–1279.
- CARN V.M. & KITCHING, R.P. (1995). The clinical response of cattle following infection with lumpy skin disease (Neethling)
 virus. Arch. Virol., 140, 503–513.
- 868 COAKLEY W. & CAPSTICK P.B. (1961). Protection of cattle against lumpy skin disease. Factors affecting small scale
 869 production of tissue culture propagated virus vaccine. *Res. Vet. Sci.*, 2, 369–371.

- COETZER J.A.W. (2004). Lumpy skin disease. *In:* Infectious Diseases of Livestock, Second Edition Coetzer J.A.W. & Justin
 R.C., eds. Oxford University Press, Cape Town, South Africa, 1268–1276.
- BAO T.D., TRAN L.H., NGUYEN H.D., HOANG T.T., NGUYEN G.H., TRAN K.V.D., NGUYEN H.X., VAN DONG H., BULA.N. & BULV.N.
 (2022). Characterization of Lumpy skin disease virus isolated from a giraffe in Vietnam. *Transbound. Emerg. Dis.*,
 69(5):e3268-e3272. doi: 10.1111/tbed.14583. Epub 2022 May 10. PMID: 35502589
- DAS A., BABIUK S. & MCINTOSH M.T. (2012). Development of a loop-mediated isothermal amplification assay for rapid detection of capripoxviruses. *J. Clin. Microbiol.*, **50**, 1613–1620.
- 877 DAVIES F.G. (1991). Lumpy Skin Disease, a Capripox Virus Infection of Cattle in Africa. FAO, Rome, Italy.
- DAVIES F.G., KRAUSS H., LUND L.J. & TAYLOR M. (1971). The laboratory diagnosis of lumpy skin disease. *Res. Vet. Sci.*, 12, 123–127.
- FAY P.C., COOK C.G., WIJESIRIWARDANA N., TORE G., COMTET L., CARPENTIER A., SHIH B., FREIMANIS G., HAGA I.R. & BEARD
 P.M. (2020). Madin-Darby bovine kidney (MDBK) cells are a suitable cell line for the propagation and study of the bovine
 poxvirus lumpy skin disease virus. J. Virol. Methods, 285, 113943. doi: 10.1016/j.jviromet.2020.113943.
- FLANNERY J., SHIH B., HAGA I.R., ASHBY M., CORLA A., KING S., FREIMANIS G., POLO N., TSE A.C., BRACKMAN C.J., CHAN J., PUN
 P., FERGUSON A.D., LAW A., LYCETT S., BATTEN C. & BEARd P.M. (2021). A novel strain of lumpy skin disease virus causes
 clinical disease in cattle in Hong Kong. *Transbound. Emerg. Dis.*, doi: 10.1111/tbed.14304
- <u>GARI G., ABIE G., GIZAW D., WUBETE A., KIDANE M., ASGEDOM H., BAYISSA B., AYELET G., OURA C.A.L., ROGER F. &</u>
 <u>TUPPURAINEN E.S.M. (2015). Evaluation of the safety, immunogenicity and efficacy of three capripoxvirus vaccine strains</u>
 against lumpy skin disease virus. *Vaccine*, **33**, 3256–3261. doi: 10.1016/j.vaccine.2015.01.035.
- 889 HAIG D. (1957). Lumpy skin disease. Bull. Epizoot. Dis. Afr., 5, 421–430.
- HEDGER R.S. & HAMBLIN C. (1983). Neutralising antibodies to lumpy skin disease virus in African wildlife. *Comp. Immunol. Microbiol. Infect. Dis.*, 6, 209–213. doi: 10.1016/0147-9571(83)90012-7. PMID: 6627909 DOI: 10.1016/0147-9571(83)90012-7.
- HAEGEMAN A., DE LEEUW I., MOSTIN L., VAN CAMPE W., AERTS L., VENTER E., TUPPURAINEN E., SAEGERMAN C. & K. DE CLERCO
 (2021). Comparative Evaluation of Lumpy Skin Disease Virus-Based Live Attenuated Vaccines. Vaccines, 9, 473.
 https://doi.org/10.3390/vaccines9050473
- HAEGEMAN A., DE LEEUW I., MOSTIN L., VAN CAMPE W., PHILIPS W., ELHARRAK M., DE REGGE N. & DE CLERCO K. (2023).
 Duration of immunity induced after vaccination of cattle with a live attenuated or inactivated Lumpy skin disease virus vaccine. *Microorganisms*, **11**, 210. https://doi.org/10.3390/microorganisms11010210
- HAMDI J., BOUMART Z., DAOUAM S., EL ARKAM A., BAMOUH Z., JAZOULI M., OMARI TADLAOUI K., FASSI FIHRI O., GAVRILOV B. & EL
 HARRAK M. (2020). Development and Evaluation of an Inactivated Lumpy Skin Disease Vaccine for Cattle. *Vet. Microbiol.*,
 245, 108689. doi: 10.1016/j.vetmic.2020.108689.
- IRELAND D.C. & BINEPAL Y.S. (1998). Improved detection of capripoxvirus in biopsy samples by PCR. J. Virol. Methods, 74,
 1–7.
- 904 IRONS P.C., TUPPURAINEN E.S.M. & VENTER E.H. (2005). Excretion of lumpy skin disease virus in bull semen. 905 Theriogenology, 63, 1290–1297.
- KARA P.D., AFONSO C.L., WALLACE D.B., KUTISH G.F., ABOLNIK C., LU Z., VREEDE F.T., TALJAARD L.C., ZSAK A., VILJOEN G.J. &
 ROCK D.L. (2003). Comparative sequence analysis of the South African vaccine strain and two virulent field isolates of
 Lumpy skin disease virus. *Arch. Virol.*, **148**, 1335–1356.
- KARA P.D., MATHER A.S., PRETORIUS A., CHETTY T., BABIUK S. & WALLACE D.B. (2018). Characterisation of putative
 immunomodulatory gene knockouts of lumpy skin disease virus in cattle towards an improved vaccine. Vaccine, 36, 4708 4715. doi: 10.1016/j.vaccine.2018.06.017.
- KITCHING R.P. & SMALE C. (1986). Comparison of the external dimensions of capripoxvirus isolates. *Res. Vet. Sci.*, 41, 425–
 427.

- 914 KLEMENT E., BROGLIA A., ANTONIOU S.-E., TSIAMADIS V., E. PLEVRAKI, PETROVIĆ T., POLAČEK V., DEBELJAK Z., MITEVA A.,
- ALEXANDROV T., MAROJEVIC D., PITE L., KONDRATENKO V., ATANASOV Z., GUBBINS S., STEGEMAN A. & CORTIÑAS ABRAHANTES J.
 (2020). Neethling vaccine proved highly effective in controlling lumpy skin disease epidemics in the Balkans. *Prev. Vet.* Med., 181, 104595 https://doi.org/10.1016/j.prevetmed.2018.12.001
- MUMAR R., GODARA B., CHANDER Y., KACHHAWA J.P., DEDAR R.K., VERMA A., RIYESH T., PAL Y., BARUA S., TRIPATHI B.N. &
 KUMAR N. (2023). Evidence of lumpy skin disease virus infection in camels. *Acta Trop.*, 242, 106922. doi: 10.1016/j.actatropica.2023.106922. Epub 2023 Apr 7. PMID: 37031926
- LAMIEN C.E., LELENTA M., GOGER W., SILBER R., TUPPURAINEN E., MATIJEVIC M., LUCKINS A.G. & DIALLO A. (2011). Real time
 PCR method for simultaneous detection, quantitation and differentiation of capripoxviruses *J. Virol. Methods*, **171**, 134–140.

MATHIJS E., VANDENBUSSCHE F., HAECEMAN A., KINC A., NTHANCENI B., POTCIETER C., MAARTENS L., VAN BORM S. & DE CLERCQ
 K. (2016). Complete Genome Sequences of the Neethling Like Lumpy Skin Disease Virus Strains Obtained Directly from
 Three Commercial Live Attenuated Vaccines. *Genome Announc.*, 4 (6). pii: e01255-16. doi: 10.1128/genomeA.01255-16.

- MILOVANOVIC M., DIETZE K., MILICEVIC V., RADOJICIC S., VALCIC M., MORITZ T. & HOFFMANN B. (2019). Humoral immune
 response to repeated lumpy skin disease virus vaccination and performance of serological tests. *BMC Vet. Res.*, **15**, 80.
 doi: 10.1186/s12917-019-1831-y.
- MURRAY L., EDWARDS L., TUPPURAINEN E.S., BACHANEK-BANKOWSKA K., OURA C.A., MIOULET V. & KING D.P. (2013). Detection of capripoxvirus DNA using a novel loop-mediated isothermal amplification assay. *BMC Vet. Res.*, **9**, 90.
- OMOGA D.C.A., MACHARIA M., MAGIRI E., KINYUA J., KASIITI J. & HOLTON T. (2016). Molecular based detection, validation of a
 LAMP assay and phylogenetic analysis of capripoxvirus in Kenya. J. Adv. Biol. Biotech., 7, 1–12.
- PESTOVA Y., BYADOVSKAYA O., KONONOV A. & SPRYGIN A. (2018). A real time high-resolution melting PCR assay for detection
 and differentiation among sheep pox virus, goat pox virus, field and vaccine strains of lumpy skin disease virus. *Mol. Cell. Probes*, 41, 57–60.
- PORCO A., CHEA S., SOURS S., NOU V., GROENENBERG M., AGGER C., TUM S., CHHUON V., SORN S., HONG C., DAVIS B., DAVIS
 S., KEN S., OLSON S.H. & FINE A.E. (2023). Case report: Lumpy skin disease in an endangered wild banteng (Bos javanicus)
 and initiation of a vaccination campaign in domestic livestock in Cambodia. *Front. Vet. Sci.*, 10:1228505. doi:
 10.3389/fvets.2023.1228505
- PROZESKY L. & BARNARD B.J.H. (1982). A study of the pathology of lumpy skin disease in cattle. Onderstepoort J. Vet. Res.,
 941 49, 167–175.
- ROUBY S. & ABOULSOUDB E. (2016). Evidence of intrauterine transmission of lumpy skin disease virus. *Vet. J.*, 209, 193–
 195.
- SAMOJLOVIC M., POLACEK V., GURJANOV V., LUPULOVIC D., LAZIC G., PETROVIĆ T., LAZIC S. (2019). Detection of antibodies against lumpy skin disease virus by virus neutralization test and ELISA methods. *Acta Vet.*, **69**, 47–60.
- SPRYGIN A., BABIN Y., PESTOVA Y., KONONOVA S., WALLACE D.B., VAN SCHALKWYK A., BYADOVSKAYA O., DIEV V., LOZOVOY D. &
 KONONOV A. (2018). Analysis and insights into recombination signals in lumpy skin disease virus recovered in the field.
 PLoS One, **13**, e0207480.
- SPRYGIN A., VAN SCHALKWYK A., SHUMILOVA I., NESTEROV A., KONONOVA S., PRUTNIKOV P., BYADOVSKAYA O. & KONONOV A.
 (2020). Full-length genome characterization of a novel recombinant vaccine-like lumpy skin disease virus strain detected during the climatic winter in Russia, 2019. *Arch. Virol.*, **165**, 2675–2677.
- TUPPURAINEN E., DIETZE K., WOLFF J., BERGMANN H., BELTRAN-ALCRUDO D., FAHRION A., LAMIEN EULOGE C., BUSCH F., SAUTER LOUIS C., CONRATHS F.J., DE CLERCQ K., HOFFMANN B. & SASCHA K. (2021). Review: Vaccines and Vaccination against
 Lumpy Skin Disease. Vaccines (Basel). 9, 1136. doi: 10.3390/vaccines9101136
- TUPPURAINEN E.S.M., VENTER E.H. & COETZER J.A.W. (2005). The detection of lumpy skin disease virus in samples of experimentally infected cattle using different diagnostic techniques. *Onderstepoort J. Vet. Res.*, **72**, 153–164.
- TUPPURAINEN E.S., VENTER E.H., COETZER J.A. & BELL-SAKYI L. (2015). Lumpy skin disease: attempted propagation in tick cell lines and presence of viral DNA in field ticks collected from naturally-infected cattle. *Ticks Tick Borne Dis.*, **6**, 134–140.

- VAN ROOYEN P.J., KOMM N.A.L., WEISS K.E. & ALEXANDER R.A. (1959). A preliminary note on the adaptation of a strain of lumpy skin disease virus to propagation in embryonated eggs. *Bull. Epizoot. Dis. Afr.*, **7**, 79–85.
- VAN SCHALKWYK A., BYADOVSKAYA O., SHUMILOVA I., WALLACE D.B. & SPRYGIN A. (2021). Estimating evolutionary changes
 between highly passaged and original parental lumpy skin disease virus strains. *Transbound. Emerg. Dis.*, doi:
 10.1111/tbed.14326.
- VAN SCHALKWYK A., KARA P., EBERSOHN K., MATHER A., ANNANDALE C.H., VENTER E.H. & WALLACE D.B. (2020). Potential link
 of single nucleotide polymorphisms to virulence of vaccine-associated field strains of lumpy skin disease virus in South
 Africa. *Transbound. Emerg. Dis.*, 67, 2946–2960.
- VIDANOVIC D., SEKLER M., PETROVIC T., DEBELJAK Z., VASKOVIC N., MATOVIC K. & HOFFMAN B. (2016). Real time PCR assay
 for the specific detection of field Balkan strain of lumpy skin disease virus. *Acta Vet. Brno*, 66, 444–454.
- WANG Y., ZHAO L., YANG J., SHI M., NIE F., LIU S., WANG Z., HUANG D., WU H., LI D., LIN H. & LI Y. (2021). Analysis of vaccine like lumpy skin disease virus from flies near the western border of China. *Transbound. Emerg. Dis.*, doi:
 10.1111/tbed.14159
- WALLACE D.B. & VILJOEN G.J. (2005). Immune responses to recombinants of the South African vaccine strain of lumpy skin
 disease virus generated by using thymidine kinase gene insertion. Vaccine, 23, 3061–3067.
- 974 WEISS K.E. (1968). Lumpy skin disease. *Virol. Monogr.*, **3**, 111–131.

987

- 975 WOLFF J., BEER M. & HOFFMANN B. (2022). High Efficiency of Low Dose Preparations of an Inactivated Lumpy Skin Disease
 976 Virus Vaccine Candidate. Vaccines (Basel), **10**, 1029. doi: 10.3390/vaccines10071029.
- 2HUGUNISSOV K., BULATOV Y., ORYNBAYEV M., KUTUMBETOV L., ABDURAIMOV Y., SHAYAKHMETOV Y., TARANOV D., AMANOVA Z.,
 MAMBETALIYEV M., ABSATOVA Z., AZANBEKOVA M., KHAIRULLIN B., ZAKARYA K. & TUPPURAINEN E. (2020). Goatpox virus (G20 LKV) vaccine strain elicits a protective response in cattle against lumpy skin disease at challenge with lumpy skin disease
 virulent field strain in a comparative study. Vet Microbiol., 245, 108695. doi: 10.1016/j.vetmic.2020.108695.
- 981
 *

 982
 *

 983
 NB: There are WOAH Reference Laboratories for lumpy skin disease (please consult the WOAH Web site:

 984
 https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).

 985
 Please contact WOAH Reference Laboratories for any further information on diagnostic tests, reagents and vaccines for lumpy skin disease
 - **NB:** FIRST ADOPTED IN 1989. MOST RECENT UPDATES ADOPTED IN 2021.

1	Annex 13. Item 5.1. – Chapter 3.6.9. Equine rhinopneumonitis (infection with equid herpesvirus-1)
2	MEETING OF THE BIOLOGICAL STANDARDS COMMISSION
3	Paris, 4–8 September 2023
4	
5	
6	CHAPTER 3.6.9.
7	EQUINE RHINOPNEUMONITIS (INFECTION WITH
8	EQUID HERPESVIRUS-1-AND-4)

SUMMARY

Equine rhinopneumonitis (ER) is a collective term for any one of several contagious, clinical disease entities of equids that may occur as a result of infection by either of two closely related herpesviruses, equid <u>alpha</u>herpesvirus-1 and -4 (EHV-1 and EHV-4). Infection with EHV-1 is listed by WOAH and is therefore <u>the focus of this chapter.</u> EHV-1 is and EHV-4 are endemic in most domestic equine populations worldwide.

9

10

11 12

13

14 Primary infection by either EHV-1-or EHV-4-is characterised by upper respiratory tract disease of varying severity that is related to the age and immunological status of the infected animal. Following viraemia EHV-15 1 also causes the more serious complications of abortion, perinatal foal death, or paralytic neurological 16 disease (equine herpesvirus myeloencephalopathy). EHV 4 has been associated with sporadic cases of 17 18 abortion, but rarely multiple abortions and not the large outbreaks associated with EHV-1. Like other 19 herpesviruses, EHV-1 and 4-induces long-lasting latent infections and can be reactivated following stress-or pregnancy. Furthermore, most horses are likely to be re-infected multiple times during their lifetime, often 20 mildly or subclinically. Detection of viral DNA or anti-EHV antibodies should therefore be interpreted with 21 22 care.

- 23 Identification of the agent: The standard method of identification of EHV-1 and EHV-4 from appropriate 24 clinical or necropsy material is by polymerase chain reaction (PCR), followed by laboratory isolation of the 25 virus in cell culture.
- Positive identification of viral isolates as EHV-1 or <u>EHV-4</u> can be achieved by type-specific PCR or sequencing. Viruses can be isolated in equine-cell culture from nasal or nasopharyngeal swab extracts taken from horses during the febrile stage of <u>with acute</u> respiratory tract infection, from the placenta, from and liver, lung, spleen, <u>adrenal glands</u> or thymus of aborted fetuses and early foal deaths, and from the leukocyte fraction of the blood of animals with acute during the febrile stage of EHV-1 infection. Unlike EHV 4, EHV 1 will also grow in various non-equine cell types such as the RK-13 cell line and this property can be used to distinguish between the two viruses.
- A rapid presumptive diagnosis of abortion induced by EHV-1 or (infrequently) EHV-4 can be achieved by
 direct immunofluorescent detection of viral antigen in cryostat sections of placenta and tissues from aborted
 fetuses, using a conjugated polyclonal antiserum.
- Post-mortem demonstration of histopathological lesions of EHV-1 in placenta and tissues from aborted
 fetuses, cases of perinatal foal death or in the central nervous system of neurologically affected animals
 complements <u>other diagnostic techniques the laboratory diagnosis</u>.
- Serological tests: Most horses possess some level of antibody to EHV-1/4, the demonstration of specific
 antibody in the serum collected from a single blood sample is therefore not confirmation of a positive
 diagnosis of recent infection. Paired, acute and convalescent sera from animals suspected of being infected

42 with EHV-1 or EHV-4 should be tested for a four-fold or greater rise in virus-specific antibody titre by either 43 virus neutralisation (VN) or complement fixation (CF) tests. Neither of these assays is type-specific but both 44 have proven useful for diagnostic purposes especially since the CF antibody response to recent infection is 45 relatively short-lived. Limited use has also been made of a type-specific enzyme-linked immunosorbent 46 assay-(Crabb et al., 1995; Hartley et al., 2005).

47 **Requirements for vaccines:** Both live attenuated and inactivated viral vaccines are available for use in 48 assisting in the control of EHV-1/4. Vaccination is helpful in reducing the severity of respiratory infection in 49 young horses and the incidence of abortion in mares, however current vaccines are not licenced to protect 50 against neurological disease. Vaccination should not be considered a substitute for sound management 51 practices known to reduce the risk of infection. Revaccination at frequent intervals is recommended in the 52 case of each of the products, as the duration of vaccine-induced immunity is relatively short.

53 Standards for production and licensing of both attenuated and inactivated EHV-1/4-vaccines are established 54 by appropriate veterinary regulatory agencies in the countries of vaccine manufacture and use. A single set 55 of internationally recognised standards for EHV vaccines is not available. In each case, however, vaccine 56 production is based on the system of a detailed outline of production employing a well characterised cell line 57 and a master seed lot of vaccine virus that has been validated with respect to virus identity, safety, virological 58 purity, immunogenicity and the absence of extraneous microbial agents.

59

A. INTRODUCTION

60 Equine rhinopneumonitis (ER) is a historically-derived term that describes a constellation of several disease entities of horses that may include respiratory disease, abortion, neonatal foal pneumonitis, or myeloencephalopathy (Allen & Bryans, 61 1986; Allen et al., 1999; Bryans & Allen, 1988; Crabb & Studdert, 1995). The disease has been is recognised for over 60 62 years as a threat to the international horse industry, and is caused by either of two members of the Herpesviridae family, 63 64 equid alphaherpesvirus-1 and -4 (EHV-1 and EHV-4). EHV-1 and EHV-4 are closely related alphaherpesviruses of horses with nucleotide sequence identity within individual homologous genes ranging from 55% to 84%, and amino acid sequence 65 identity from 55% to 96% (Telford et al., 1992; 1998). The two herpesviruses With the exception of EHV-1 in Iceland 66 (Thorsteinsdóttir et al., 2021), the two herpesviruses are considered endemic enzotic in all countries in which large 67 populations of horses are maintained as part of the cultural tradition or agricultural economy. There is no recorded evidence 68 that the two herpesviruses of ER pose any health risks to humans working with the agents. Infection with EHV-1 is listed 69 by WOAH and is therefore the focus of this chapter. 70

Viral transmission to cohort animals occurs by inhalation of aerosols of virus-laden respiratory secretions. Morbidity tends to be highest in young horses sharing the same air space. Aborted tissues and placental fluids from infected mares can contain extremely high levels of live virus and represent a major source of infection. Extensive use of vaccines has not eliminated EHV-1 infections, and the world-wide-annual financial impact from this these-equine pathogens is immense considerable.

In horses under 3 years of age, clinical ER usually takes the form of an acute, febrile respiratory illness that spreads rapidly 76 through the group of animals. The viruses infects and multiplies multiply in epithelial cells of the respiratory mucosa. Signs 77 78 of infection become apparent 2-8 days after exposure to virus, and are characterised by fever, inappetence, depression, and nasal discharge. The severity of respiratory disease varies with the age of the horse and the level of immunity resulting 79 from previous vaccination or natural exposure. Bi-phasic fever, viraemia and complications are more likely with EHV-1 than 80 81 EHV-4. Subclinical infections with EHV-1/4-are common, even in young animals. Although mortality from uncomplicated 82 ER is rare and complete recovery within 1-2 weeks is the normal outcome, respiratory infection is a frequent and significant cause of interrupted schedules among horses assembled for training, racing, or other equestrian events. Fully protective 83 84 immunity resulting from infection is of short duration, and convalescent animals are susceptible to reinfection by EHV-1/4 85 after several months. Although reinfections by the two herpesviruses cause less severe or clinically inapparent respiratory disease, the risks of subsequent abortion or neurological disease remain. Like other herpesviruses, EHV-1/4 causes long-86 lasting latent infections and latently infected horses represent a potential infection risk for other horses. Virus can be 87 88 reactivated as a result of stress or pregnancy. The greatest clinical threats to individual breeding, racing, or pleasure horse operations posed by ER are the potential abortigenic and neurological sequelae of EHV-1 respiratory infection. ER 89 abortions occur annually in horse populations worldwide and may be sporadic or multiple. Foals infected in utero may be 90 born alive and die within a few days of birth. EHV-1 neurological disease is less common than abortions but has been 91 92 recorded all over the world with associated fatalities. Outbreaks result in movement restrictions and, sometimes, 93 cancellation of equestrian events (Couroucé et al., 2023; FEI, 2021).

94 Neurological disease, also known as equine herpesvirus myeloencephalopathy, remains an infrequent but serious

95 complication of EHV-1 infection. A single mutation in the DNA polymerase gene (ORF30) has been associated with 96 increased risk of neurological disease, however strains without this change can also cause paralysis (Goodman *et al.*,

2007; Nugent *et al.*, 2006). Strain typing techniques have been employed to identify viruses carrying the neuropathic

marker, and it can be helpful to be aware of an increased risk of neurological complications. However, for practical purposes
 strain typing is not relevant for agent identification, or international trade. Strain typing may be beneficial for implementation
 of biosecurity measures in the management of outbreaks of equine herpesvirus myeloencephalopathy.

101 <u>Strain typing has been shown to be not reliable for predicting the clinical outcome of EHV-1 infection but can be useful in</u> 102 <u>epidemiological investigations (Garvey *et al.*, 2019; Nugent *et al.*, 2006; Sutton *et al.*, 2019).</u>

B. DIAGNOSTIC TECHNIQUES

Both-EHV-1 and EHV-4 is transmitted by the respiratory route and has have the potential to be highly contagious viruses 104 105 particularly where large numbers of horses are housed in the same air space. EHV1 and the former can cause explosive outbreaks of abortion or neurological disease. Rapid diagnostic methods are therefore essential useful for managing the 106 disease. Real-time polymerase chain reaction (PCR) assays are widely routinely used by diagnostic laboratories worldwide 107 and are both rapid and sensitive. Real-time PCR assays that allow simultaneous testing for EHV-1 and EHV-4 and 108 109 quantification of viral load have been developed. Virus isolation has been replaced by real-time PCR as the frontline 110 diagnostic test in the majority of laboratories but can also be useful, particularly for the detection of viraemia. This is also true of for EHV-1 associated abortions and neonatal foal deaths, when the high level of virus in the tissues usually produces 111 a cytopathic effect in 1-3 days. Immunchistochemical or immunofluorescent approaches are employed in some 112 laboratories can be extremely useful for rapid diagnosis of EHV-induced abortion from fresh or embedded tissue-and are 113 114 relatively straightforward. Several other techniques based on enzyme-linked immunosorbent assay (ELISA) or nucleic acid hybridisation probes have also been described, however their use is often restricted to specialised laboratories and they 115 are not included here. Virus neutralisation (VN) and complement fixation (CF) are the most frequently used serological 116 tests, and seroconversion in paired samples is considered indicative of exposure to virus by natural infection or by 117 vaccination. 118

119

103

Table 1. Test methods available for the diagnosis of equine rhinopneumonitis and their purpose

	Purpose								
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	al Contribute to Confirmation from eradication of clinical cases prior policies ^(a) cases cases		Prevalence of infection - surveillance	Immune status in individual animals or populations post-vaccination			
Identification of the agent ^(b)									
Virus isolation	-	- ++_+							
PCR	-	+++	-	+++	-	-			
<u>Direct</u> immunofluorescence	Ξ	Ξ	Ξ	<u>++</u>	Ξ	≡			
Detection of immune response									
VN	*±	+±	<u></u> _+	+++	+++	+++			
ELISA	+	 <u></u> = +	 <u></u> <u></u> +	+_+	++_+	+			
CFT	_	- <u>++</u>	_	+++	_	<u>+++</u>			

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

- = suitable in very limited circumstances; = not appropriate for this purpose.
- PCR = polymerase chain reaction; VN = virus neutralisation; ELISA = enzyme-linked immunosorbent assay; CFT = complement fixation test.
 - ^(a)No eradication policies exist for equine rhinopneumonitis.

^(b)A combination of agent identification methods applied on the same clinical sample is recommended.

126 1. Identification Detection of the agent

127 **1.1. Collection and preparation of specimens**

128 Nasal/nasopharyngeal swabs: swab extract can be used for DNA extraction and subsequent virus detection by PCR using one of a variety of published techniques or commercially available kits (see below). Virus isolation can also be 129 130 attempted from the swab extracts. To increase the chances of isolating live virus, swabs are best obtained from horses during the very early, febrile stages acute stage of the respiratory disease, and are collected via the nares by 131 sampling the area with a swab of an appropriate size and length for horses. After collection, the swab should be 132 133 removed and transported immediately to the virology laboratory in 3 ml of cold (not frozen) virus transport medium (e.g. phosphate buffered saline [PBS] or serum-free MEM [minimal essential medium] with antibiotics). Virus 134 135 infectivity can be prolonged by the addition of bovine serum albumin, fetal calf serum or gelatine to 0.1% (w/v).

136 Tissue samples: total DNA can be extracted using a number of commercially available kits and used in PCR to detect viral DNA (described below in Section B.1.2.1). Virus isolation from placenta and fetal tissues from suspect cases of 137 EHV-1 abortion is most successful when performed on aseptically collected samples of placenta, liver, lung, thymus. 138 adrenal glands and spleen. Virus may be isolated from post-mortem cases of EHV-1 neurological disease by culture 139 140 of samples of brain and spinal cord but such attempts to isolate virus are often unsuccessful; however, they may be 141 useful for PCR testing and pathological examination. Tissue samples should be transported to the laboratory and held at 4°C until inoculated into tissue culture. Samples that cannot be processed within a few hours should be stored 142 at -70°C. 143

Blood: for virus <u>detection by PCR or</u> isolation from blood leukocytes, collect a <u>10–</u>20 ml sample of blood, using an
 aseptic technique in-citrate, heparin or EDTA [ethylene diamine tetra-acetic acid] anticoagulant. EDTA is the preferred
 anticoagulant for PCR testing <u>in some laboratories as heparin may inhibit DNA polymerase</u>. The samples should be
 transported without delay to the laboratory on ice, but not frozen.

148 <u>Cerebrospinal fluid: the detection of EHV-1 DNA in cerebrospinal fluid has been reported in cases of neurological</u>
 149 <u>disease.</u>

150 **1.2. Virus detection by polymerase chain reaction**

PCR has become the primary diagnostic method for the detection of EHV-1 and EHV-4 in clinical specimens, paraffin-151 embedded archival tissue, or inoculated cell cultures (Borchers & Slater, 1993; Lawrence et al., 1994; O'Keefe et al., 152 1994; Varrasso et al., 2001). A variety of type-specific PCR primers have been designed to distinguish between the 153 154 presence of EHV 1 and EHV 4. The correlation between PCR and virus isolation techniques for diagnosis of EHV 1 or EHV 4 is high (Varrasso et al., 2001). Diagnosis by PCR is rapid, sensitive, and does not depend on the presence 155 of infectious virus in the clinical sample. For diagnosis of active infection by EHV, PCR methods are routinely used 156 157 to detect EHV-1 DNA in nasopharyngeal swabs and tissue samples most reliable with tissue samples from aborted fetuses and placental tissue or from nasopharyngeal swabs of foals and yearlings. They are particularly useful in 158 explosive outbreaks of abortion, respiratory or neurological disease in which a rapid identification and monitoring of 159 160 the virus spread is critical for guiding management strategies, including movement restrictions. PCR examination of spinal cord and brain tissue, as well as peripheral blood mononuclear cells (PBMC), are important in seeking a 161 diagnosis on a horse with neurological signs (Pronost et al., 2012). 162

163 Several PCR assays have been published. A nested PCR procedure can be used to distinguish between EHV-1 and EHV-1. A sensitive protocol suitable for clinical or pathological specimens (nasal secretions, blood leukocytes, brain 164 and spinal cord, fetal tissues, etc.) has been described by Borchers & Slater (1993). However, nested PCR methods 165 have a high risk of laboratory cross-contamination, and sensitive rapid one-step PCR tests to detect EHV-1 and EHV-166 4 (e.g. Lawrence et al., 1994) are preferred. The WOAH Reference Laboratories use quantitative real-time PCR 167 assays such as those targeting heterologous sequences of major glycoprotein genes to distinguish between EHV 1 168 and EHV-1. A multiplex real-time PCR targeting glycoprotein B gene of EHV-1 and EHV-4 was described by Diallo 169 170 et al. (2007). PCR protocols have been developed that can differentiate between EHV 1 strains carrying the ORF30 neuropathogenic marker, using both restriction enzyme digestion of PCR products (Fritsche & Borchers, 2011) or by 171 172 quantitative real-time PCR (Allen et al, 2007, Smith et al., 2012). Methods have also been developed to type strains for epidemiological purposes, based on the ORF68 gene (Nugent et al., 2006). The WOAH Reference Laboratories 173 employ in-house methods for strain typing, however these protocols have not yet been validated between different 174 175 laboratories at an international level.

176 Real-time (or quantitative) PCR has become the method of choice for <u>many_the majority of</u> diagnostic_<u>tests</u> 177 <u>laboratories</u> and provides rapid and sensitive detection of viral DNA. Equine post-mortem tissues from newborn and 178 adult animals or equine fetal tissue from abortions (tissues containing lung, liver, spleen, thymus, adrenal gland and 179 placental tissues) can be used. For respiratory samples, equine nasopharyngeal swabs or deep nasal swabs (submitted in a suitable viral transport medium), <u>buffy coat</u>, tracheal wash (TW) or broncho-alveolar lavage (BAL) are
 all suitable. DNA should be extracted using an appropriate kit or robotic system.

182There is no internationally standardised real-time PCR method for EHV-1 but Table 2 summarises the primer and183probe sequences for some of the most widely used assays. Type-specific PCR primers have been designed to184distinguish between the presence of EHV-1 and EHV-4. The optimised thermocycler times and temperatures are185documented in the publications cited.

<u>Primer</u>	Primer sequence (5' to 3')	<u>Target</u>	<u>Reference</u>
Forward	CAT-GTC-AAC-GCA-CTC-CCA		
<u>Reverse</u>	<u>GGG-TCG-GGC-GTT-TCT-GT</u>	EHV-1 gB	<u>Diallo <i>et al.,</i> 2006</u>
<u>Probe</u>	FAM-CCC-TAC-GCT-GCT-CC-MGB-NFQ		
Forward	CAT-ACG-TCC-CTG-TCC-GAC-AGA-T		
Reverse	<u>GGTACTCGGCCTTTGACGAA</u>	EHV-1 gB	<u>Hussey <i>et al.,</i> 2006</u>
<u>Probe</u>	FAM-TGA-GAC-CGA-AGA-TCT-CCT-CCA-CCG-A-BHQ1		
<u>Forward</u>	TAT-ACT-CGC-TGA-GGA-TGG-AGA-CTT-T		
<u>Reverse</u>	TTG-GGG-CAA-GTT-CTA-GGT-GGT-T	EHV-1 gB	<u>Pusterla <i>et al.,</i> 2009</u>
<u>Probe</u>	6FAM-ACA-CCT-GCC-CAC-CGC-CTA-CCG		
<u>Forward</u>	GCG-GGC-TCT-GAC-AAC-ACA-A		ISO 17025 accredited for the
<u>Reverse</u>	TTG-TGG-TTT-CAT-GGG-AGT-GTG-TA	EHV-1 gC	detection of EHV-1 at WOAH
<u>Probe</u>	FAM-TAA-CGC-AAA-CGG-TAC-AGA-A-BHQ1		Reference Laboratory

Table 2. Primer and probe sequences for EHV1/4 detection by real-time PCR

*This multiplex real-time PCR test has been validated to ISO 17025 and is designed for use in a 96-well format. This 187 188 can be readily combined with automatic nucleic acid extraction methods. Discrimination between EHV-1 and EHV-4 is carried out by the incorporation of type specific dual labelled probes based on methods published by Hussey et al. 189 (2006) and Lawrence et al. (1994). To establish such a real-time PCR assay for diagnostic purposes, validation 190 191 against blinded samples is required. Sensitivity and specificity for the assay should be determined against each target. Support for development of assays and appropriate sample panels can be obtained from the WOAH Reference 192 Laboratories. Reference material and sample panels for real-time PCR can be obtained from the WOAH Reference 193 194 Laboratories.

195 • **Point**

196

197

198

199

200

201

202

203 204

205

206

186

Point of care (POC) molecular tests

Loop-mediated isothermal amplification (LAMP) assays for the detection of EHV-1 have been described (Nemoto *et al.*, 2011). An evaluation of a hydrolysis probe-based insulated isothermal PCR (iiPCR) assay for the detection of EHV-1 showed it to have a high sensitivity and specificity compared with real-time PCR (Balasuriya *et al.*, 2017). However further validation of POC tests in the field is required.

Molecular characterisation

Allelic discrimination real-time PCR assays identifying a single nucleotide polymorphism that was originally suggested to distinguish between neuropathogenic and non-neuro-pathogenic EHV-1 strains have been developed (Smith *et al.*, 2012). However, investigations in many countries worldwide demonstrated that the nucleotide substitution was not a reliable predictor of enhanced neuropathogenicity. Multilocus typing and whole genome sequencing are useful for molecular epidemiological studies (Garvey *et al.*, 2019). Nugent *et al.*, 2006; Sutton *et al.*, 2019).

1.3. Virus isolation

<u>Virus isolation is no longer a routine test used for EHV-1 detection in the majority of diagnostic laboratories but is</u>
 <u>more often conducted for surveillance and research purposes.</u> A number of cell types may be used for isolation of
 EHV-1 (e.g. rabbit kidney [RK-13 (AATC-CCL37)], baby hamster kidney [BHK-21], Madin–Darby bovine kidney
 [MDBK], pig kidney [PK-15], etc.). <u>RK13 cells are commonly used for this purpose.</u> For efficient primary isolation of

212 EHV-4 from horses with respiratory disease, equine-derived cell cultures must be used. Both EHV-1 and EHV-4 may 213 be isolated from nasopharyngeal samples using primary equine fetal kidney cells or equine fibroblasts derived from dermal (E-Derm) or lung tissue. EHV-1 can be isolated on other cell types, as will be discussed later. The 214 215 nasopharyngeal swab and its accompanying transport medium are transferred into the barrel of a sterile 10 ml syringe. Using the syringe plunger, the fluid is squeezed from the swab into a sterile tube. A portion of the expressed 216 217 fluid can be filtered through a sterile, 0.45 µm membrane syringe filter unit into a second sterile tube if heavy bacterial contamination is expected, but this may also lower virus titre. Recently prepared cell monolayers in tissue culture 218 flasks are inoculated with the filtered, as well as the unfiltered, nasopharyngeal swab extract. Cell monolayers in 219 multiwell plates incubated in a 5% CO2-environment may also be used. Virus is allowed to attach by incubating the 220 inoculated monolayers at 37°C. Monolayers of uninoculated control cells should be incubated in parallel. 221

- 222 At Recently prepared cell monolayers in tissue culture flasks or plates are inoculated with nasopharyngeal swab extract or homogenised tissue: approximately 10% (w/v) pooled tissue homogenates of liver, lung, thymus, adrenal 223 and spleen (from aborted fetuses/neonatal foals) or of brain and spinal cord (from cases of neurological disease). 224 225 Virus is allowed to attach by incubating the end of the attachment period, inoculated monolayers at 37°C for 1 hour after which the inocula are removed and the monolayers are rinsed twice with PBS to remove virus-neutralising 226 227 antibody that may or maintenance medium. Monolayers of uninoculated control cells should be present in the nasopharyngeal secretions-incubated in parallel. After addition of supplemented maintenance medium (MEM 228 229 containing 2% fetal calf serum [FCS] and twice the standard concentrations of antibiotics/antifungals [penicillin, 230 streptomycin, gentamicin, and amphotericin B]), the flasks are incubated at 37°C in a 5% CO₂ environment.
- 231 The use of a positive control virus samples of relatively low titre may be used to validate the isolation procedure carries the risk that this may lead but should be processed separately to eventual avoid contamination of diagnostic 232 specimens. This risk can be minimised by using routine precautions and good laboratory technique, including the use 233 of biosafety cabinets, inoculating positive controls after the diagnostic specimens, decontaminating the surfaces in 234 the hood while the inoculum is adsorbing and using a positive control of relatively low titre. Inoculated flasks should 235 be inspected daily by microscopy for the appearance of characteristic herpesvirus cytopathic effect (CPE) (focal 236 237 rounding, increase in refractility, and detachment of cells). Cultures exhibiting no evidence of viral CPE after 1 week 238 of incubation should be blind-passaged into freshly prepared monolayers of cells, using small aliquots of both media and cells as the inoculum. Further blind passage is usually not productive. 239

240 It can be useful to inoculate samples into both non equine and equine cells in parallel to distinguish between EHV 1 and EHV-1, since EHV-1 can cause sporadic cases of abortion. Around 10% (w/v) pooled tissue homogenates of 241 liver, lung, thymus, and spleen (from aborted fetuses) or of central nervous system tissue (from cases of neurological 242 disease) are used for virus isolation. These are prepared by first mincing small samples of tissue into 1 mm cubes in 243 244 a sterile Petri dish with dissecting scissors, followed by macerating the tissue cubes further in serum-free culture medium with antibiotics using a homogeniser or mechanical tissue grinder. After centrifugation at 1200 g for 245 10 minutes, the supernatant is removed and 0.5 ml is inoculated into duplicate cell monolayers in tissue culture flasks. 246 Following incubation of the inoculated cells at 37°C for 1.5-2 hours, the inocula are removed and the monolavers are 247 248 rinsed twice with PBS or maintenance medium. After addition of 5 ml of supplemented maintenance medium, the flasks are incubated at 37°C for up to 1 week or until viral CPE is observed. Cultures exhibiting no evidence of viral 249 250 CPE after 1 week of incubation should be passaged a second time into freshly prepared monolayers of cells, using 251 small aliquots of both media and cells as the inoculum.

252 Blood samples: EHV-1 and, infrequently, EHV-4-can be isolated from PBMC. Buffy coats may be prepared from unclotted (heparinised) blood by centrifugation at 600-525 g for 15-5 minutes, and. The buffy coat is taken after the 253 plasma has been carefully removed. The buffy coat is then layered onto a PBMC separating solution (Ficoll; density 254 255 1077 g/ml, commercially available) and centrifuged at 400 g for 20 minutes. The PBMC interface (without most granulocytes) is and washed twice in PBS (300 g for 10 minutes) and resuspended in 1 ml three times in 3 ml MEM 256 257 containing 2% FCS. As a quicker alternative method, PBMC may be collected by centrifugation directly from plasma. (525 g for 5 minutes). Following the third wash, the buffy coat is harvested and resuspended in 2.5 ml MEM containing 258 2% FCS. An aliquot of the rinsed cell suspension is added to each of the duplicate monolayers of equine fibroblast, 259 equine fetal or RK-13 cell monolayers in 25 cm² flasks containing 8-10 ml freshly added maintenance medium. The 260 flasks can be used for DNA extraction. For virus isolation, the resuspended cells (1 ml) are co-cultivated with freshly 261 262 prepared primary equine lung or RK-13 cell suspensions (5 ml) in 25 cm² flasks. Confluent cell monolayers are not 263 used. The flasks are incubated at 37°C in a 5% CO2 environment for 3 days or until the cells have reached 90% confluence. The monolayers are then rinsed three times with 1 × PBS and supplemented with 5 ml MEM containing 264 2% FCS. They are incubated at 37°C for 7 days; either with or without removal of the inoculum. If PBMCs are not 265 removed prior to incubation, CPE may be difficult to detect in the presence of the massive inoculum of leukocytes: 266 267 each flask of cells is freeze-thawed after 7 for a further 4 days of incubation and the contents centrifuged at 300 g 268 for 10 minutes. Finally, 0.5 ml of the cell free, culture medium supernatant is transferred to freshly made cell monolayers that are just subconfluent. These are incubated and observed daily for viral CPE for at least 5-6 days. 269 270 Again, samples. Samples exhibiting no evidence of viral CPE after 1 week of incubation should be passaged a second 271 time before discarding as negative.

Virus identity may be confirmed by PCR or by immunofluorescence with specific antisera. Virus isolates from positive cultures should be submitted to a WOAH Reference Laboratory <u>for strain characterisation and</u> to maintain a geographically diverse archive. Further strain characterisation for surveillance purposes or detection of the neurological marker can be provided at some laboratories.

1.4. Virus detection by direct immunofluorescence

- 277 Direct immunofluorescent detection of EHV<u>-1</u> antigens in samples of post-mortem tissues collected from aborted 278 equine fetuses and the placenta provides a rapid preliminary diagnosis of herpesvirus abortion (Gunn, 1992). The 279 diagnostic reliability of this technique approaches that of virus isolation attempts from the same tissues.
- In the United States of America (USA), potent polyclonal antiserum to EHV-1, prepared in swine and conjugated with FITC, is available to veterinary diagnostic laboratories for this purpose from the National Veterinary Services Laboratories of the United States Department of Agriculture (USDA). The antiserum cross-reacts with EHV-4 and hence is not useful for serotyping, however virus typing can be conducted on any virus positive specimens by PCR.
- Freshly dissected samples (5 × 5 mm pieces) of fetal tissue (lung, liver, thymus, and spleen) are frozen, sectioned on a cryostat at -20°C, mounted on to microscope slides, and fixed with 100% acetone. After air-drying, the sections are incubated at 37°C in a humid atmosphere for 30 minutes with an appropriate dilution of the conjugated swine antibody to EHV-1. Unreacted antibody is removed by two washes in PBS, and the tissue sections are then covered with aqueous mounting medium and a cover-slip, and examined for fluorescent cells indicating the presence of EHV antigen. Each test should include a positive and negative control consisting of sections from known EHV-1 infected and uninfected fetal tissue.

291 **1.5. Virus detection by immunoperoxidase staining**

292 Immunohistochemical (IH) staining methods, such as immunoperoxidase, have been developed for detecting EHV-1 antigen in fixed tissues of aborted equine fetuses, placental tissues or neurologically affected horses (Schultheiss et 293 al., 1993; Whitwell et al., 1992). Such techniques can be used as an alternative to immunofluorescence described 294 above and can also be readily applied to archival frozen or fixed tissue samples. Immunohistochemical staining for 295 EHV-1 is particularly useful for the simultaneous evaluation of morphological lesions and the identification of the virus. 296 297 Immunoperoxidase staining for EHV-1/4 may also be carried out on infected cell monolayers-(van Maanen et al., 2000). Adequate controls must be included with each immunoperoxidase test run for evaluation of both the method 298 specificity and antibody specificity. In one WOAH Reference Laboratory, this method is used routinely for frozen or 299 fixed tissue, using If non-specific rabbit-polyclonal sera is used raised against EHV 1. This staining method is not 300 type-specific and therefore the staining method needs to be combined with virus isolation or PCR to discriminate 301 between EHV-1 and EHV-4, however it provides a useful method for rapid diagnosis of EHV-induced abortion. 302

303 **1.6. Histopathology**

Histopathological examination of sections of fixed placenta and lung, liver, spleen, adrenal and thymus from aborted fetuses and brain and spinal cord from neurologically affected horses should be carried out. In aborted fetuses, eosinophilic intranuclear inclusion bodies present within bronchiolar epithelium or in cells at the periphery of areas of hepatic necrosis are consistent with a diagnosis of herpesvirus infection. The characteristic microscopic lesion associated with EHV-1 neuropathy is a degenerative thrombotic vasculitis of small blood vessels in the brain or spinal cord (perivascular cuffing and infiltration by inflammatory cells, endothelial proliferation and necrosis, and thrombus formation).

311 2. Serological tests

EHV-1 and EHV-4 are endemic in most parts of the world and seroprevalence is high, however serological testing of paired
sera can be useful for diagnosis of ER in horses. A positive diagnosis is based on the demonstration of significant increases
(four-fold or greater) in antibody titres in paired sera taken during the acute and convalescent stages of the disease. The
results of tests performed on sera from a single collection date are, in most cases, impossible to interpret with any degree
of confidence. The initial (acute phase) serum sample should be taken as soon as possible after the onset of clinical signs,
and the second (convalescent phase) serum sample should be taken 2–4 weeks later.

'Acute phase' sera from mares after abortion or from horses with EHV-1 neurological disease may already contain maximal
 titres of EHV-1 antibody, with no increase in titres detectable in sera collected at later dates. In such cases, serological
 testing of paired serum samples from clinically unaffected cohort members of the herd may prove useful for retrospective
 diagnosis of ER within the herd.

322 Finally, the serological detection of antibodies to EHV-1 in heart or umbilical cord blood or other fluids of equine fetuses 323 can be of diagnostic value in cases of abortion especially when the fetus is virologically negative. The EHV 1/4 nucleic acid 324 may be identified from these tissues by PCR.

Serum antibody levels to EHV-1/4 may be determined by virus neutralisation (VN) (Thomson et al., 1976), complement 325 326 fixation (CF) tests (Thomson et al., 1976) or enzyme-linked immunosorbent assay (ELISA) (Crabb & Studdert, 1995). There 327 are no internationally recognised reagents or standardised techniques for performing any of the serological tests for 328 detection of EHV-1/4 antibody; titre determinations on the same serum may differ from one laboratory to another. Furthermore, The CF and VN tests detect antibodies that are cross-reactive between EHV-1 and EHV-4. Nonetheless, the 329 demonstration of a four-fold or greater rise in antibody titre to EHV-1 or EHV-4 during the course of a clinical illness provides 330 331 serological confirmation of recent infection with one of the viruses. Commercial ELISAs that distinguish EHV-1 and EHV-332 4 antibodies are available but less widely used than the CF and VN tests. Unlike other alphaherpesviruses, DIVA⁴¹ ELISAs, 333 which have been very useful in eradication programmes for bovine rhinotracheitis and pseudorabies (Aujeszky's disease), 334 have not been developed for EHV-1/4.

The microneutralisation test is a <u>VN and the CF tests are</u> widely used and sensitive serological assays for detecting EHV-1/4 antibody and will thus be described here.

337 **2.1. Virus neutralisation test**

338 This test is most commonly performed in flat-bottom 96-well microtitre plates (tissue culture grade) using a constant dose of virus and doubling dilutions of equine test sera. At least two-three replicate wells for each serum dilution are 339 340 required. Heat-inactivated maintenance medium with a concentration of 2% FCS (HIMM) Serum free MEM is used 341 throughout as a diluent. Virus stocks of known titre are diluted just before use to contain 100 TCID₅₀ (50% tissue culture infective dose) in 25 µl. Monolayers of E-Derm or RK-13 cells are prepared monodispersed with EDTA/trypsin 342 and resuspended at a concentration of 5 × 10⁵/ml. Note that RK-13 cells can be used with EHV-1 but do not show 343 344 CPE with EHV-4. Antibody positive and negative control equine sera and controls for cell viability, virus infectivity, and test serum cytotoxicity, must be included in each assay. End-point VN titres of antibody are calculated by 345 determining the reciprocal of the highest serum dilution that protects ≥75% 100% of the cell monolayer from virus 346 destruction in both of the replicate wells. 347

- 348 Serum toxicity may be encountered in samples from horses repeatedly vaccinated with a commercial vaccine 349 prepared from EHV-1 grown up in RK-13 cells. This can give rise to difficulties in interpretation of test reactions at 350 lower serum dilutions. The problem can be overcome using E-Derm or other non-rabbit kidney derived cell line.
- 351 2.1.1. Test procedure

352 353

354

355

356

357

358 359

360

361 362

363

364

365 366

367

368 369

370 371

- A suitable test procedure is as follows:
- i) <u>Prepare semi-confluent monolayers in tissue culture microtitre plates.</u>
 - ii) Inactivate test and control sera for 30 minutes in a water bath at 56°C.
 - iii) Add <u>40</u> 25 µl of <u>HIMM</u> serum free MEM to all wells of the microtitre assay plates.
 - iv) <u>For test sample titration</u>, pipette 25 40 µl of each test serum into duplicate triplicate wells of both rows A and B of the plate. The first two rows serve as the dilution of the test serum and the third row serves as the serum toxicity control-and the second row as the first dilution of the test. Make doubling dilutions of each serum starting with row B and proceeding to the bottom of the plate by sequential mixing and transfer of 25 40 µl to each subsequent row of wells. Six sera can be assayed in each plate. Add 40µl of HIMM to the serum control rows.
- Add <u>40</u> 25–µl of the appropriately diluted EHV-1 or EHV-4 virus stock to each <u>all</u> wells (100 TCID₅₀/well) of the test plate except those of row A, which are the serum controls wells. Note that the final serum dilutions, after addition of virus, run from <u>a starting dilution of 1/4-te</u> 1/256. A separate control plate should include titration of both a negative and positive (<u>high and low</u>) horse serum sera of known titre, cell control (no virus), and a back titration of virus control (no serum), and a virus titration using six wells per log dilution (100 TCID₅₀ to 0.01 TCID₅₀/well) calculate the actual amount of virus used in the test
- vi) Incubate the plates for 1 hour at 37°C in 5% CO₂ atmosphere. Add 50 μl of the prepared E-Derm or RK-13 cell suspension (5 × 10⁵ cells/ml) in MEM/10% FCS to each well.
 - vii) Transfer 50 µl from each well of the test and control plates to the tissue culture microtitre plates.

⁴¹ DIVA: detection of infection in vaccinated animals

372	viii)	Incubate the plates for 2-4–5 days at 37°C in an atmosphere of 5% CO ₂ in air.
	,	
373	ix)	Examine the plates microscopically for CPE and record the results on a worksheet. <u>Confirm the</u>
374		validity of the test by establishing that the working dilution of stock virus is at 100 TCID ₅₀ /well,
375		that the (high and low) positive control sera are within one well of their pre-determined titre and
376		that the negative control serum is negative at a 1/4 dilution. This takes approximately 72 hours.
377		If at this stage the antigen is too weak the virus concentration may be increased by extending
378		the incubation period up to 5 days. If the antigen is too strong the test must be repeated.
379		Wells are scored as positive for neutralisation of virus if ≥ 75% of the cell monolayer remains
380		intact. The highest dilution of serum resulting in ≥ 75% neutralisation of virus (<25% CPE) in
381		replicate wells is the end-point titre for that serum. Examine the plates microscopically for CPE
382		and record the results on a worksheet.
383	x)	Alternatively, the cell monolayers can be scored for CPE after fixing and staining as follows:
384	,	after removal of the culture fluid, immerse the plates for 15 minutes in a solution containing 2
385		mg/ml crystal violet, 10% formalin, 45% methanol, and 45% water. Then, rinse the plates
386		vigorously under a stream of running tap water. Wells containing intact cell monolayers stain
387		blue, while monolayers destroyed by virus do not stain. Verify that the cell control, positive
388		serum control, and serum cytotoxicity control wells stain blue, that the virus control and negative
389		serum control wells are not stained, and that the actual amount of virus added to each well is
390		between 10 ^{1.5} and 10 ^{2.5} TCID ₅₀ . Wells are scored as positive for neutralisation of virus if 100%
		of the cell monolayer remains intact. The highest dilution of serum resulting in complete
391 200		neutralisation of virus (no CPE) in both duplicate wells is the end-point titre for that serum.
392		
393	xi)	Calculate the neutralisation titre for each test serum, and compare acute and convalescent
394		phase serum titres from each animal for a four-fold or greater increase.
395 396	The CF test can be u	nt fixation test used for the detection and quantification of antibodies against EHV-1. The test determines whether
397		ntibody are capable of forming a complex. The presence of an immune complex is revealed by
398		which consists of guinea-pig complement and sensitised sheep red blood cells (SRBCs) coated
399		lytic serum (haemolysin). In the absence of antibodies against equine herpesvirus, no
400		mplex is formed, the complement remains free in the solution and the sensitised SRBCs become
401		nce of antibodies against equine herpesvirus, an antibody/antigen complex is formed, the
402		es fixed and is therefore unable to lyse the SRBCs. They subsequently form a button at the bottom
403	<u>of the test well.</u>	
404	Guinea-pig compler	nent, rabbit haemolytic serum, complement fixation diluent (CFD) and bovine serum albumin
405		ned commercially. The dilution of guinea-pig complement that has activity at 3 HD (haemolytic
406		ce of sensitised SRBCs should be optimised. The recommended dilution of rabbit haemolytic
407		ng dilution) is sometimes provided by the supplier. However, the optimal dilution of haemolysin
408		d with the in use reagents (complement etc.) so that the test can be performed reproducibly. The
409		ion of antigen to be used in the test should be determined using an antigen versus antibody
410		ique and by testing a panel of known positive sera.
411		d in U bottomed microtitre plates. Paired sera should be assayed on the same plate. An antibody
412		Id be included as a control on each plate. All sera are tested on a second plate containing all
413		virus to check for anti-complementary activity. A back titration of the working dilution (3 HD) of
414		0, 1 HD, 0.5 HD is set up in duplicate wells on the complement control plate (eight wells in total).
415	An SRBC control is	<u>set up in eight wells.</u>
416	<u>2.2.1. Pre</u>	paration of samples
417	i)	Samples and controls are prepared by adding 4 volumes (600 µl) of CFD to 1 volume (150 µl)
418	,	of test sera to give a 1/5 dilution.
419	ii)	Diluted serum is inactivated for 30 minutes at 60°C to destroy the naturally occurring
419	")	complement.
421	<u>2.2.2. Tes</u>	t procedure

422 423	i)	<u>Prepare the test plate and anti-complementary plate by adding 25 µl 0.05% BSA/CFD to all wells except the first column (H).</u>
424	ii)	Add 50 µl of 0.05% BSA/CFD to the eight wells of the complement control (back titration).

425		iii)	Add 75 µl of 0.05% BSA/CFD to eight wells of cell control.
426		iv)	Add 50 µl of the diluted inactivated test serum and controls to the first well of each row on both
427 428			the test and anti-complementary plates. Serial doubling dilutions are then made by transferring 25 µl across the plate and discarding the final 25 ml.
429		v)	Place the microtitre plates on ice for addition of antigen and complement.
430		vi)	Add 25 µl of antigen (diluted to working strength in 0.05% BSA/CFD) to the test plates.
431 432		vii)	Add 25 µl of 0.05% BSA/CFD to all wells of the anti-complementary plate to compensate for lack of antigen.
432		viii)	Add 25 µl of guinea-pig complement diluted in 0.05% BSA/CFD to 3 HD to all wells except the
433 434		viii)	complement control and SRBC control.
435 436		ix)	Back titrate the working dilution of 3 HD complement to 2 HD, 1 HD and 0.5 HD in 200 µl volumes. Add 25 µl of each dilution to the appropriate wells.
437		x)	Incubate all plates at 4°C overnight.
438	<u>2.2.3.</u>	Pre	paration and addition of sheep blood
439		i)	SRBCs collected into Alsever's solution are washed twice in 0.05% BSA/PBS solution.
440 441		ii)	Gently resuspend the SRBCs in 5–10 ml 0.05% BSA/CFD solution. Dilute to 2% SRBCS (v/v packed cells) in BSA/CFD solution.
442		iii)	Mix the 2% SRBCs with an equal volume of BSA/CFD solution containing haemolysin at its
443 444			optimal sensitising concentration to give a 1% SRBC solution. Prepare an appropriate volume of this solution by allowing 3 ml per microtitre plate.
445		iv)	Incubate at 37°C for 10 minutes. Store the 1% sensitised SRBCs overnight at 4°C.
446		v)	The following day, incubate the 1% sensitised SRBCs at 37°C for 30 minutes. During the final
447		•)	20 minutes of this incubation, transfer the test plates from 4°C to 37°C.
448 449		vi)	At the end of the 30-minute incubation, add 25 ml of 1% sensitised SRBCs to all plates. Mix on a plate shaker for 30 seconds.
450 451		vii)	Incubate the plates at 37°C for 30 minutes. Shake the plates after 15 minutes and at the end of this incubation (a total of three times).
452		viii)	Incubate the plates at 4°C for 2 hours to allow the cells to settle.
453		ix)	Read and record the test results after 2 hours.
454	<u>2.2.4.</u>	Rea	ding results
455		i)	Confirm the validity of the test by establishing that the working dilution of complement is at 3 HD:
456 457			<u>100% lysis at 3 HD and 2 HD, and 50% lysis at 1 HD. Distinct buttons should be visible in the eight wells of the SRBC control.</u>
458 459		ii)	There must be 100% lysis observed at the 1/5 dilution for the negative control (<5). The antibody titre of the positive control serum must read within one well of its predetermined titre.
460		iii)	Confirm that there are no buttons visible on the anti-complementary plates. Buttoning indicates
461 462			either the presence of residual native complement in the sample or that there is a non-specific complement fixing effect occurring. Sera that show anti-complementary activity should be
463			retested and treated as described below.
464		iv)	In the test wells, buttoning indicates the presence of antibodies in the serum. The antibody titre
465			is the dilution at which there is 50% buttoning and 50% lysis observed.
466	<u>2.2.5.</u>	Trea	atment of samples showing anti-complementary activity
467 468		i)	Add 50 µl of guinea-pig complement to 150 µl of the serum showing anti-complementary activity.
469		ii)	Incubate the sample at 37°C for 30 minutes.
470		iii)	Add 550 µl of CFD (1:5 dilution).
471		iv)	Heat inactivate at 60°C for 30 minutes.

C. REQUIREMENTS FOR VACCINES

Background 473 1.

472

474 Both live attenuated and inactivated vaccines are available for use in horses as licensed, commercially prepared products for use in reducing the impact of disease in horses caused by EHV-1/4 infection. The products contain different 475 permutations of EHV-1 and EHV-4 and some also include equine influenza virus. 476

477 Clinical experience has demonstrated that vaccination can be useful for reducing clinical signs of respiratory disease and 478 incidence of abortion, however none of the vaccines protects against neurological disease. Multiple doses repeated 479 annually, of each of the currently marketed ER vaccines are recommended by their respective manufacturers. Vaccination schedules vary with a particular vaccine. 480

The indications stated on the product label for use of several available vaccines for ER are either as a preventative of 481 482 herpesvirus-associated respiratory disease, or as an aid in the prevention of abortion, or both. A minority of Only four 483 vaccine products have met the regulatory requirements for claiming efficacy in providing protection from herpesvirus abortion as a result of successful vaccination and challenge experiments in pregnant mares. None of the vaccine products 484 have been demonstrated to prevent the occurrence of neurological disease sometimes associated with EHV-1 infection. 485

486 Guidelines for the production of veterinary vaccines are given in Chapter 1.1.8 Principles of veterinary vaccine production. 487 The guidelines given here and in chapter 1.1.8 are intended to be general in nature and may be supplemented by national and regional requirements. 488

2. Outline of production and minimum requirements for vaccines 489

490 2.1. Characteristics of the seed

491 The master seed virus (MSV) for ER vaccines must be prepared from strains of EHV-1 and/or EHV-4 that have been positively and unequivocally identified by both serological and genetic tests. Seed virus must be propagated in a cell 492 line approved for equine vaccine production by the appropriate regulatory agency. A complete record of original 493 source (including isolate number, location, year of isolation), passage history, medium used for propagation, etc., 494 495 shall be kept for the master seed preparations of both the virus(es) and cell stock(s) intended for use in vaccine production. 496

- 497
- 498 499

500 501

502

503 504

505

506

507 508

509

510

511

2.1.1. Biological characteristics of the master seed

- Permanently stored stocks of both MSV and master cell stock (MCS) used for vaccine production must be demonstrated to be pure, safe and, in the case of MSV, also immunogenic.
 - Generally, the fifth passage from the MSV and the twentieth passage from the MCS are the highest allowed for vaccine production. Results of all quality control tests on master seeds must be recorded and made a part of the licensee's permanent records.

2.1.2. Quality criteria

Tests for master seed purity include prescribed procedures that demonstrate the virus and cell seed stocks to be free from bacteria, fungi, mycoplasmas, and extraneous viruses. Special tests must be performed to confirm the absence of equine arteritis virus, equine infectious anaemia virus, equine influenza virus, equine herpesvirus-2, -3, and -5, equine rhinitis A and B viruses, the alphaviruses of equine encephalomyelitis, bovine viral diarrhoea virus (BVDV - common contaminant of bovine serum), and porcine parvovirus (PPV - potential contaminant of porcine trypsin). The purity check should also include the exclusion of the presence of EHV-1 from EHV-4 MSV and vice versa.

2.1.3. Validation as a vaccine strain

Tests for immunogenicity of the EHV-1/4 MSV stocks should be performed in horses on an 512 experimental test vaccine prepared from the highest passage level of the MSV allowed for use in 513 vaccine production. The test for MSV immunogenicity consists of vaccination of horses with low 514 antibody titres to EHV-1/4, with doses of the test vaccine that will be recommended on the final 515 product label. Second serum samples should be obtained and tested for significant increases in 516 neutralising antibody titre against the virus, 21 days after the final dose. 517 Samples of each lot of MSV to be used for preparation of live attenuated ER vaccines must be tested 518 for safety in horses determined to be susceptible to the virulent wild-type virus, including pregnant 519

mares in the last 4 months of gestation. Vaccine safety must be demonstrated in a 'safety field trial' in horses of various ages from three different geographical areas. The safety trial should be conducted by independent veterinarians using a prelicensing batch of vaccine. EHV-1 vaccines making a claim for efficacy in controlling abortion must be tested for safety in a significant number of late gestation pregnant mares, using the vaccination schedule that will be recommended by the manufacturer for the final vaccine product.

526 2.2. Method of manufacture

520

521 522

523

524 525

527 528

529

530

531

532

533 534

535

536

537

538 539

540

541 542

543

544

546

547

550

553

555

556 557

558

559

560

561 562

563

564

567

568

2.2.1. Procedure

A detailed protocol of the methods of manufacture to be followed in the preparation of vaccines for ER must be compiled, approved, and filed as an Outline of Production with the appropriate licensing agency. Specifics of the methods of manufacture for ER vaccines will differ with the type (live or inactivated) and composition (EHV-1 only, EHV-1 and EHV-4, EHV-4 and equine influenza viruses, etc.) of each individual product, and also with the manufacturer.

2.2.2. **Requirements for ingredients**

Cells, virus, culture medium, and medium supplements of animal origin that are used for the preparation of production lots of vaccine must be derived from bulk stocks that have passed the prescribed tests for bacterial, fungal, and mycoplasma sterility; nontumorgenicity; and absence of extraneous viral agents.

2.2.3. Final product batch tests

i) Sterility

Samples taken from each batch of completed vaccine are tested for bacteria, fungi, and mycoplasma contamination. Procedures to establish that the vaccine is free from extraneous viruses are also required; such tests should include inoculation of cell cultures that allow detection of the common equine viruses, as well as techniques for the detection of BVDV and PPV in ingredients of animal origin used in the production of the batch of vaccine.

ii) Identity 545

> Identity tests shall demonstrate that no other vaccine strain is present when several strains are propagated in a laboratory used in the production of multivalent vaccines.

iii) Safety 548

549 Safety tests shall consist of detecting any abnormal local or systemic adverse reactions to the vaccine in the host species by all vaccination route(s). Tests to assure safety of each production batch of ER 551 vaccine must demonstrate complete inactivation of virus (for inactivated vaccines) as well as a level 552 of residual virus-killing agent that does not exceed the maximal allowable limit (e.g. 0.2% for formaldehyde).

iv) Batch potency 554

> Batch potency is examined on the final formulated product. Batch control of antigenic potency for EHV-1 vaccines only may be tested by measuring the ability of dilutions of the vaccine to protect hamsters from challenge with a lethal dose of hamster adapted EHV 1 virus. Although potency testing on production batches of ER vaccine may also-be performed by vaccination of susceptible horses followed by assay for seroconversion, the recent availability of virus type-specific MAbs has permitted development of less costly and more rapid in-vitro immunoassays exist for antigenic potency. The basis for such in-vitro assays for ER vaccine potency is the determination, by use of the specific MAb, of the presence of at least the minimal amount of viral antigen within each batch of vaccine that correlates with the required level of protection (or seroconversion rate) in a standard animal test for potency.

- 2.3. Requirements for authorisation/registration/licencing 565
- 566

Manufacturing process 2.3.1.

For registration of vaccine, all relevant details concerning manufacture of the vaccine and quality control testing (see Sections C.2.1 and C.2.2) should be submitted to the authorities. This information 569 shall be provided from three consecutive vaccine batches with a volume not less than 1/3 of the typical industrial batch volume. 570 2.3.2 571 Safety requirements Vaccine safety should be evaluated in vaccinated animals using different assays (see Section 572 573 2.2.3.iii). 2.3.3 Efficacy requirements 574 Vaccine efficacy (protection) is estimated in vaccinated animals directly by evaluating their resistance 575 to live pathogen challenge. 576 2.3.4 **Duration of immunity** 577 As part of the licensing or marketing authorisation procedure, the manufacturer may be required to 578 demonstrate the duration of immunity (DOI) of a given vaccine by either challenge or alternative test 579 at the end of the claimed period of protection. 580 Tests to establish the duration of immunity to EHV-1/4 or EHV1/4 achieved by immunisation with 581 each batch of vaccine are not required. The results of many reported observations indicate that 582 immunity induced by vaccination-against EHV-1 or EHV induced immunity to EHV-1/4 is not more 583 584 than a few months in duration; these observations are reflected in the frequency of revaccination recommended on ER vaccine product labels. 585 2.3.5 Stability 586 As part of the licensing or marketing authorisation procedure, the manufacturer will be required to 587 588 demonstrate the stability of all the vaccine's properties at the end of the claimed shelf-life period. 589 Storage temperature shall be indicated and warnings should be given if product is damaged by freezing or ambient temperature. 590 591 At least three production batches of vaccine should be tested for shelf life before reaching a conclusion on the vaccine's stability. When stored at 4°C, inactivated vaccine products generally 592 593 maintain their original antigenic potency for at least 1 year. Lyophilised preparations of the live virus vaccine are also stable during storage for 1 year at 4°C. Following reconstitution, live virus vaccine 594 595 is unstable and cannot be stored without loss of potency.

596 **Note:** current vaccines are authorised for prevention of respiratory disease or as an aid in the prevention of abortion. 597 Unless the vaccine's ability to prevent neurological disease is under investigation, the virus used in the challenge 598 experiments should not be a strain with a history of inducing neurological disease.

599

REFERENCES

600 ALLEN G.P. (2007). Development of a real-time polymerase chain reaction assay for rapid diagnosis of neuropathogenic 601 strains of equine herpesvirus-1. *J Vet. Diagn. Invest.,* **19**, 69–72.

ALLEN G.P. & BRYANS J.T. (1986). Molecular epidemiology, pathogenesis and prophylaxis of equine herpesvirus-1
 infections. *In:* Progress in Veterinary Microbiology and Immunology, Vol. 2, Pandey R., ed. Karger, Basel, Switzerland &
 New York, USA, 78–144.

ALLEN G.P., KYDD J.H., SLATER J.D. & SMITH K.C. (1999). Recent advances in understanding the pathogenesis, epidemiology, and immunological control of equid herpesvirus-1 (EHV-1) abortion. *Equine Infect. Dis.*, **8**, 129–146.

ALLKOFER A., GARVEY M., RYAN E., LYONS R., RYAN M., LUKASEVICIUTE G., WALSH C., VENNER M. & CULLINANE A. (2021).
 Primary vaccination in foals: a comparison of the serological response to equine influenza and equine herpesvirus vaccines
 administered concurrently or 2 weeks apart. *Arch. Virol.*, **166**, 571–579.

ATASEVEN V.S., DAĞALP S.B., GÜZEL M., BAŞARAN Z., TAN M.T. & GERAGHTY B. (2009). Prevalence of equine herpesvirus-1
 and equine herpesvirus-4 infections in equidae species in Turkey as determined by ELISA and multiplex nested PCR. *Res. Vet. Sci.*, 86, 339–344.

- BALASURIYA U.B.R., LEE P.-Y.A., TSAI Y.-L., TSAI C.-F., SHEN Y.-H., CHANG H.-F. G., SKILLMAN A., WANG H.-T.T., PRONOST S.
 & ZHANG Y. (2017). Translation of a laboratory-validated equine herpesvirus-1 specific real-time PCR assay into an
- insulated isothermal polymerase chain reaction (iiPCR) assay for point-of-need diagnosis using POCKIT[™] nucleic acid
 analyzer. J. Virol. Methods, 241, 58–63.
- 617 BANNAI H., MAE N., ODE H., NEMOTO M., TSUJIMURA K., YAMANAKA T., KONDO T. & MATSUMURA T. (2014). Successful control 618 of winter pyrexias caused by equine herpesvirus type 1 in Japanese training centers by achieving high vaccination 619 coverage. *Clin. Vaccine Immunol.*, **21**, 1070–1076.
- BANNAI H., TSUJIMURA K., NEMOTO M., OHTA M., YAMANAKA T., KOKADO H. & MATSUMURA T. (2019). Epizootiological investigation of equine herpesvirus type 1 infection among Japanese racehorses before and after the replacement of an inactivated vaccine with a modified live vaccine. *BMC Vet. Res.*, **15**, 280.
- 623 BARBIC L., LOJKIC I., STEVANOVIC V., BEDEKOVIC T., STARESINA V., LEMO N., LOJKIC M. & MADIC J. (2012). Two outbreaks of 624 neuropathogenic equine herpesvirus type 1 with breed-dependent clinical signs. *Vet. Rec.*, **170**, 227.
- BRESGEN C., LÄMMER M., WAGNER B., OSTERRIEDER N. & DAMIANI A.M. (2012). Serological responses and clinical outcome
 after vaccination of mares and foals with equine herpesvirus type 1 and 4 (EHV-1 and EHV-4) vaccines. *Vet. Microbiol.*,
 160, 9–16.
- 628 <u>COUROUCÉ A., NORMAND C., TESSIER C., POMARES R., THÉVENOT J., MARCILLAUD-PITEL C., LEGRAND L., PITEL P.H., PRONOST</u>
 629 <u>S. & LUPO C. (2023). Equine Herpesvirus-1 Outbreak During a Show-Jumping Competition: A Clinical and Epidemiological</u>
 630 <u>Study. J. Equine Vet. Sci., 128, 104869.</u>
- 631 BORCHERS K. & SLATER J. (1993). A nested PCR for the detection and differentiation of EHV 1 and EHV 4. J. Virol. Methods, 632 **45**, 331–336.
- BRYANS J.T. & ALLEN G.P. (1988). Herpesviral diseases of the horse. *In:* Herpesvirus Diseases of Animals, Wittman G., ed.
 Kluwer, Boston, USA, 176–229.
- 635 CRABB B.S., MACPHERSON C.M., REUBEL G.H., BROWNING G.F., STUDDERT M.J. & DRUMMER H.E. (1995). A type-specific 636 serological test to distinguish antibodies to equine herpesviruses 4 and 1. *Arch. Virol.*, **140**, 245–258.
- CRABB B.S. & STUDDERT M.J. (1995). Equine herpesviruses 4 (equine rhinopneumonitis virus) and 1 (equine abortion virus).
 Adv. Virus Res., 45, 153–190.
- DIALLO I.S., HEWITSON G., WRIGHT L., RODWELL B.J. & CORNEY B.G. (2006). Detection of equine herpesvirus type 1 using a
 real-time polymerase chain reaction. *J. Virol. Methods*, **131**, 92–98.
- 641 DIALLO I.S., HEWITSON G., WRICHT L.L., KELLY M.A., RODWELL B.J. & CORNEY B.G. (2007). Multiplex real time PCR for 642 detection and differentiation of equid herpesvirus 1 (EHV-1) and equid herpesvirus 4 (EHV-4). *Vet. Microbiol.*, **123**, 93– 643 403.
- EL BRINI Z., FASSI FIHRI O. & PAILLOT R. (2021). Seroprevalence of Equine Herpesvirus 1 (EHV-1) and Equine Herpesvirus
 4 (EHV-4) in the Northern Moroccan Horse Populations. *Animals* (*Basel*), **11**, 2851.
- <u>FÉDÉRATION EQUESTRE INTERNATIONALE (FEI) (2021), Management of an EHV-1 outbreak at FEI events and its international</u>
 <u>impact. Vet. Rec.</u>, **189**, e905. https://doi.org/10.1002/vetr.905
- FOOTE C.E., GILKERSON J.R., WHALLEY J.M. & LOVE D.N. (2003). Seroprevalence of equine herpesvirus 1 in mares and foals
 on a large Hunter Valley stud farm in years pre- and postvaccination. *Aust. Vet. J.*, 81, 283–288.
- 650 <u>GARVEY M., LYONS R. & HECTOR R.D. (2019). Molecular Characterisation of Equine Herpesvirus 1 Isolates from Cases of</u> 651 <u>Abortion, Respiratory and Neurological Disease in Ireland between 1990 and 2017. *Pathogens*, **8**, 7.</u>
- 652 <u>GILDEA S., SANCHEZ HIGGINS M.J., JOHNSON G., WALSH C. & CULLINANE A. (2016). Concurrent vaccination against equine</u> 653 <u>influenza and equine herpesvirus – a practical approach. *Influenza Other Respir. Viruses*, **10**, 433–437.</u>
- <u>GILKERSON J. R., WHALLEY J. M., DRUMMER H. E., STUDDERT M.J. & LOVE D.N. (1999). Epidemiology of EHV-1 and EHV-4 in</u>
 the mare and foal populations on a Hunter Valley stud farm: are mares the source of EHV-1 for unweaned foals. *Vet.* <u>Microbiol.</u>, 68, 27–34.

- 657 <u>GRYSPEERDT A., VANDERKERCKHOVE A., VAN DOORSSELAERE J., VAN DE WALLE G., & NAUWYNCK H. (2011). Description of an</u>
 658 <u>unusually large outbreak of nervous system disorders caused by equine herpesvirus 1 (EHV1) in 2009 in Belgium. Vlaams</u>
 659 <u>Diergeneeskundig Tijdschrif, 80, 147–153.</u>
- 660 FRITSCHE A.K. & BORCHERS K. (2011). Detection of neuropathogenic strains of equid herpesvirus 1 (EHV-1) associated with 661 abortions in Germany. *Vet. Microbiol.*, **147**, 176-180.
- 662 GUNN H.M. (1992). A direct fluorescent antibody technique to diagnose abortion caused by equine herpesvirus. *Irish Vet.* 663 *J.*, **44**, 37–40.
- HELDENS J.G., HANNANT D., CULLINANE A.A., PRENDERGAST M.J., MUMFORD J.A., NELLY M., KYDD J.H., WESTSTRATE M.W. &
 VAN DEN HOVEN R.(2001). Clinical and virological evaluation of the efficacy of an inactivated EHV1 and EHV4 whole virus
 vaccine (Duvaxyn EHV1,4). Vaccination/challenge experiments in foals and pregnant mares. *Vaccine*, **19**, 4307–4317.
- HENNINGER R.W., REED S.M., SAVILLE W.J., ALLEN G.P., HASS G.F., KOHN C.W. & SOFALY C. (2007). Outbreak of neurologic
 disease caused by equine herpesvirus-1 at a university equestrian center. J. Vet. Intern. Med., 21, 157–165.

GOODMAN L.B., LORECIAN A., PERKINS G.A., NUCENT J., BUCKLES E.L., MERCORELLI B., KYDD J.H., PALÙ G., SMITH K.C.,
 OSTERRIEDER N. & DAVIS-POYNTER N. (2007). A point mutation in a herpesvirus polymerase determines neuropathogenicity.
 PLoS Pathog., 3 (11), e160.

- 672 HARTLEY C.A., WILKS C.R., STUDDERT M.J. & GILKERSON J.R. (2005). Comparison of antibody detection assays for the 673 diagnosis of equine herpesvirus 1 and 4 infections in horses. *Am. J. Vet. Res.*, **66**, 921–928.
- HUSSEY S.B., CLARK R., LUNN K.F., BREATHNACH C., SOBOLL G., WHALLEY J.M. & LUNN D.P. (2006). Detection and
 quantification of equine herpesvirus-1 viremia and nasal shedding by real-time polymerase chain reaction. *J. Vet. Diagn. Invest.*, **18**, 335–342.
- <u>KYDD J. H., WATTRANG E. & HANNANT D. (2003). Pre-infection frequencies of equine herpesvirus-1 specific, cytotoxic T</u>
 <u>lymphocytes correlate with protection against abortion following experimental infection of pregnant mares. Vet. Immunol.</u>
 <u>Immunopathol.</u>, 96, 207–217.
- LAWRENCE G.L., GILKERSON J., LOVE D.N., SABINE M. & WHALLEY J.M. (1994). Rapid, single-step differentiation of equid
 herpesvirus 1 and 4 from clinical material using the polymerase chain reaction and virus-specific primers. *J. Virol. Methods*,
 47, 59–72.
- 683 <u>MCCARTAN C.G., RUSSELL M.M., WOOD J.L. & MUMFORD J.A. (1995). Clinical, serological and virological characteristics of</u> 684 <u>an outbreak of paresis and neonatal foal disease due to equine herpesvirus-1 on a stud farm. *Vet. Rec.*, **136**, 7–12.</u>
- MOORE, S.E., Strang, C.L., Marr, C.M., Newton, R. and Cameron, I.J. (2019), Management of an outbreak of multiple
 equine herpesvirus type 1 abortions among vaccinated mares on a large UK Thoroughbred stud farm. Vet Rec Case Rep,
 7: e000799.
- NEMOTO M., OHTA M., TSUJIMURA K., BANNAI H., YAMANAKA T., KONDO T. & MATSUMURA T. (2011). Direct detection of equine
 herpesvirus type 1 DNA in nasal swabs by loop-mediated isothermal amplification (LAMP). J. Vet. Med. Sci., 73, 1225–
 1227.
- NUGENT J., BIRCH-MACHIN I., SMITH K.C., MUMFORD J.A., SWANN Z., NEWTON J.R., BOWDEN R.J., ALLEN G.P. & DAVIS-POYNTER
 N. (2006). Analysis of equid herpesvirus 1 strain variation reveals a point mutation of the DNA polymerase strongly
 associated with neuropathogenic versus nonneuropathogenic disease outbreaks. J. Virol., 80, 4047–4060.
- 694 <u>PRONOST S., LEGRAND L., PITEL P.H., WEGGE B., LISSENS J., FREYMUTH F., RICHARD E. & FORTIER G. (2012). Outbreak of</u>
 695 <u>equine herpesvirus myeloencephalopathy in France: a clinical and molecular investigation. *Transbound. Emerg. Dis.*, 59,
 696 256–263.
 </u>
- PUSTERLA N., KALSCHEUR M., PETERS D., BIDWELL L., HOLTZ S., BARNUM S. & LAWTON K. (2023). Investigation of the Frequency
 of Detection of Common Respiratory Pathogens in Nasal Secretions and Environment of Healthy Sport Horses Attending
 a Multi-Week Show Event during the Summer Months. *Viruses*, **15**, 1225.
- PUSTERLA N., SANDLER-BURTNESS E., BARNUM S., HILL L.A., MENDONSA E., KHAN R., PORTENER D., RIDLAND H. & SCHUMACHER
 S. (2022). Frequency of Detection of Respiratory Pathogens in Nasal Secretions From Healthy Sport Horses Attending a
 Spring Show in California. J. Equine Vet. Sci, 117, 104089.

- PUSTERLA N., WILSON W.D., MAPES S., FINNO C., ISBELL D., ARTHUR R.M. & FERRARO G.L. (2009). Characterization of viral loads, strain and state of equine herpesvirus-1 using real-time PCR in horses following natural exposure at a racetrack in California. Vet. J., **179**, 230–239.
- 706 O'KEEFE J.S., JULIAN A., MORIARTY K., MURRAY A. & WILKS C.R. (1994). A comparison of the polymerase chain reaction with 707 standard laboratory methods for the detection of EHV-1 and EHV-4 in archival tissue samples. *N.Z. Vet. J.*, **42**, 93–96.
- SCHULTHEISS P.C., COLLINS J.K. & CARMAN J. (1993). Use of an immunoperoxidase technique to detect equine herpesvirus 1 antigen in formalin-fixed paraffin-embedded equine fetal tissues. J. Vet. Diagn. Invest., 5, 12–15.
- Singh B.K., Ahuja S. & Gulati B.R. (2004). Development of a neutralizing monoclonal antibody-based blocking ELISA for
 detection of equine herpesvirus 1 antibodies. *Vet. Res .Commun.*, 28, 437–446.
- SMITH F.L., WATSON J.L., SPIER S.J., KILCOYNE I., MAPES S., SONDER C. & PUSTERLA N. (2018). Frequency of shedding of respiratory pathogens in horses recently imported to the United States. J. Vet. Intern. Med., 32, 1436–1441.
- SMITH K.L., LI Y., BREHENY P., COOK R.F., HENNEY P.J., SELLS S., PRONOST S., LU Z., CROSSLEY B.M., TIMONEY P.J. &
 BALASURIYA U.B. (2012). Development and validation of a new and improved allelic discrimination real-time PCR assay for
 the detection of equine herpesvirus-1 (EHV-1) and differentiation of A2254 from G2254 strains in clinical specimens. *J. Clin. Microbiol.*, **50**, 1981–1988.
- <u>STRANG C. & NEWTON R. (2017). Control and disease clearance after neurological EHV-1 in the UK. Vet. Rec., 181, 678–</u>
 <u>679.</u>

SUTTON G., GARVEY M., CULLINANE A., JOURDAN M., FORTIER C., MOREAU P., FOURSIN M., GRYSPEERDT A., MAISONNIER V.,
 MARCILLAUD-PITEL C., LEGRAND L., PAILLOT R. & PRONOST S. (2019). Molecular Surveillance of EHV-1 Strains Circulating in
 France during and after the Major 2009 Outbreak in Normandy Involving Respiratory Infection, Neurological Disorder, and
 Abortion. Viruses, 11, 916.

- TELFORD E.A.R., WATSON M.S., McBride K. & Davison A.J. (1992). The DNA sequence of equine herpesvirus 1. *Virology*,
 189, 304–316.
- TELFORD E.A.R., WATSON M.S., PERRY J., CULLINANE A.A. & DAVISON A.J. (1998). The DNA sequence of equine herpesvirus 4. *J. Gen. Virol.*, **79**, 1197–1203.
- THOMSON G.R., MUMFORD J.A., CAMPBELL J., GRIFFITHS L. & CLAPHAM P. (1976). Serological detection of equid herpesvirus 1 infections of the respiratory tract. *Equine Vet. J.*, **8**, 58–65.
- THORSTEINSDÓTTIR L., GUÐMUNDSSON G., JENSSON H., TORSTEINSDÓTTIR S. & SVANSSON V. (2021). Isolation of equid
 alphaherpesvirus 3 from a horse in Iceland with equine coital exanthema. *Acta Vet. Scand.*, 63, 6.
- YAN MAANEN C., VREESWIJK J., MOONEN P., BRINKHOF J. DE BOER-LUIJTZE E. & TERPSTRA C. (2000). Differentiation and genomic and antigenic variation among fetal, respiratory, and neurological isolates from EHV1 and EHV4 infections in The Netherlands. *Vet. Q.*, 22, 88–93.
- VAN MAANEN C., WILLINK D.L., SMEENK L.A., BRINKHOF J. & TERPSTRA C. (2000). An equine herpesvirus 1 (EHV1) abortion
 storm at a riding school. Vet. Q., 22, 83–87.
- VARRASSO A., DYNON K., FICORILLI N., HARTLEY C.A., STUDDERT M.J. & DRUMMER H.E. (2001). Identification of equine
 herpesviruses 1 and 4 by polymerase chain reaction. *Aust. Vet. J.*, **79**, 563–569.
- WHITWELL K.E., GOWER S.M. & SMITH K.C. (1992). An immunoperosidase method applied to the diagnosis of equine herpesvirus abortion, using conventional and rapid microwave techniques. *Equine Vet. J.*, **24**, 10–12.
- YASUNAGA S., MAEDA K., MATSUMURA T., KONDO T. & KAI K. (2000). Application of a type-specific enzyme-linked
 immunosorbent assay for equine herpesvirus types 1 and 4 (EHV-1 and -4) to horse populations inoculated with inactivated
 EHV-1 vaccine. J. Vet. Med. Sci., 62, 687–691.

* *

744 745

 746 NB: There are WOAH Reference Laboratories for equine rhinopneumonitis (please consult the WOAH Web site: <u>https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3</u>).

 748 Please contact the WOAH Reference Laboratories for any further information on

749 750	diagnostic tests, reagents and vaccines for equine rhinopneumonitis and to submit strains for further characterisation.
751	NB: First adopted in 1990. Most recent updates adopted in 2017.

1

6

7

8

9

Annex 14. Item 5.1. – Chapter 3.8.1. Border disease

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023

SECTION 3.8.

OVIDAE AND CAPRINAE

CHAPTER 3.8.1.

BORDER DISEASE

SUMMARY

10 Border disease (BD) is a viral disease of sheep and goats first reported in sheep in 1959 from the border region of England and Wales, and since recorded world-wide. Prevalence rates in sheep vary from 5% to 11 12 50% between countries and from region to region within countries. Clinical signs include barren ewes, 13 abortions, stillbirths and the birth of small weak lambs. Affected lambs can show and a fine tremor, abnormal 14 body conformation and hairy fleeces (so-called 'hairy-shaker' or 'fuzzy' lambs). Consequently, the disease has sometimes been referred to as 'hairy shaker disease'. Vertical transmission plays an important role in 15 the epidemiology of the disease. Infection of fetuses can result in the birth of persistently infected (PI) lambs. 16 17 These PI lambs are viraemic, antibody negative and constantly excrete virus. The virus spreads from sheep 18 to sheep, with PI animals being the most potent source of infection. Infection in goats is less common with 19 abortion being the main presenting sign.

- 20 BD is caused by the Pestivirus border disease virus (BDV), but in some parts of the world, especially where there is close contact between sheep or goats and cattle, the same clinical signs may be caused by infection 21 22 with bovine viral diarrhoea virus (BVDV). Therefore the genetic and antigenic differences between BDV and 23 BVDV need to be taken into consideration when investigating disease outbreaks or certifying animals or germplasm for international movement. It is important to identify the viraemic PI animals so that they will not 24 25 be used for breeding or trading purposes. Serological testing is insufficient. However, it is generally considered that serologically positive, nonviraemic sheep are 'safe', do not present a risk as latent infections 26 are not known to occur in recovered animals. Pregnant seropositive, nonviraemic animals may, however, 27 present a risk by carrying a PI fetus that cannot be detected until after parturition. 28
- Identification of the agent: BDV is a <u>species of</u> Pestivirus (<u>Pestivirus ovis</u>) in the family Flaviviridae and is
 closely related to classical swine fever virus (<u>Pestivirus suis</u>) and BVDV <u>viruses, which are classified in the</u>
 <u>distinct species Pestivirus bovis (commonly known as BVDV type 1)</u>, <u>Pestivirus tauri (formerly BVDV type 2)</u>
 <u>and Pestivirus brazilense (BVDV type 3 or Hobi-like pestivirus</u>). Nearly all isolates of BDV are
 noncytopathogenic in cell culture. There are no defined serotypes but virus isolates exhibit considerable
 antigenic diversity. A number of separate genotypes, have been identified.
- Apparently healthy PI sheep resulting from congenital infection can be identified by direct detection of virus or nucleic acid in blood or tissues or by virus isolation in cell culture followed by immunostaining to detect the noncytopathogenic virus.

- 38 Diagnostic methods: The demonstration of virus by culture and antigen detection may be less reliable in 39 lambs younger than 2 months that have received colostral antibody. Acute infection is usually subclinical 40 and viraemia is transient and difficult to detect. The isolation of virus from tissues of aborted or stillborn 41 lambs is often difficult but virus can be detected by sensitive reverse transcriptase polymerase chain reaction 42 methods that are able to detect residual nucleic acid. However, tissues and blood from PI sheep more than a few months old contain high levels of virus, which can be easily identified by isolation and direct methods 43 to detect antigens or nucleic acids. As sheep may be infected with BVDV, it is preferable to use diagnostic 44 assays that are 'pan-pestivirus' reactive and will readily detect all strains of BDV and BVDV. 45
- 46 **Serological tests:** Acute infection with BDV is best confirmed by demonstrating seroconversion using paired 47 or sequential samples from several animals in the group. The enzyme-linked immunosorbent assay and 48 virus neutralisation test (VNT) are the most commonly used antibody detection methods. Due to the antigenic 49 differences between BDV and BVDV, assays for the detection of antibodies to BDV, especially by VNT, 50 should preferably be based on a strain of BDV.
- 51 **Requirements for vaccines:** There is no standard vaccine for BDV, but a commercial killed whole-virus 52 vaccine has been produced. Ideally, such a vaccine should be suitable for administration to females before 53 breeding for prevention of transplacental infection. The use of BVDV vaccines has been advocated, but the 54 antigenic diversity of BD viruses must be considered. <u>In many instances, the antigenic diversity of BDV</u> 55 <u>strains is sufficiently different to BVDV that a BVDV vaccine is unlikely to provide protection.</u>
- 56 BD viruses have contaminated several modified live veterinary vaccines produced in sheep cells or 57 containing sheep serum. This potential hazard should be recognised by manufacturers of biological 58 products.

A. INTRODUCTION

59

60 Border disease virus (BDV) is a Pestivirus of the family Flaviviridae and is closely related to classical swine fever virus (CSFV) and bovine viral diarrhoea virus (BVDV). There are four number of officially recognised species, namely - BDV 61 (Pestivirus ovis) CSFV (Pestivirus suis), BVDV types 1 and 2 (taxonomically known as Pestivirus bovis and Pestivirus tauri, 62 respectively) and BDV (ICTV, 2016) BVDV 3 or Hobi-like pestivirus (Pestivirus brazilense), but a number of other 63 64 pestiviruses that are considered to be distinct species have been reported. While CSF viruses are predominantly restricted to pigs, examples of there are situations where the other three species have all-been recovered from sheep. While the 65 majority of isolates have been identified as BD viruses in areas where sheep or goats are raised in isolation from other 66 species (Vilcek et al., 1997), in regions where there is close contact between small ruminants and cattle. BVDV may be 67 68 frequently identified (Carlsson, 1991). Nearly all virus isolates of BDV are noncytopathogenic, although occasional cytopathic viruses have been isolated (Vantsis et al., 1976). BDV spreads naturally among sheep by the oro-nasal route 69 and by vertical transmission. It is principally a cause of congenital disease in sheep and goats, but can also cause acute 70 71 and persistent infections. Infection is less common in goats, in which persistent infection is rare as abortion is the main presenting sign. Pigs may also be infected by pestiviruses other than CSFV and antibodies to BDV in pigs may interfere 72 73 with tests for the diagnosis of CSF (Oguzoglu et al., 2001). Several genotypes of BD viruses from sheep, goats and Pyrenean chamois (Rupicapra pyrenaica pyrenaica) have been described. Phylogenetic analysis using computer-assisted 74 nucleotide sequence analysis suggests that genetic variability among BD viruses is greater than within each of the other 75 76 Pestivirus species. Four distinguishable genogroups of BDV have been described as well as putative novel Pestivirus genotypes from Tunisian sheep and a goat (Becher et al., 2003; Vilcek & Nettleton, 2006). The chamois BD virus is similar 77 to isolates from sheep in the Iberian Peninsula (Valdazo-Gonzalez et al., 2007). This chapter describes BDV infection in 78 sheep. Chapter 3.4.7 Bovine viral diarrhoea should also be consulted for related diagnostic methods. 79

80 1. Acute infections

Healthy newborn and adult sheep exposed to BDV usually experience only mild or inapparent disease. Slight fever and a mild leukopenia are associated with a short-lived viraemia detectable between days 4 and 11 post-infection, after which virus neutralising antibody appears in the serum (Thabti *et al.*, 2002).

Acute infections are best diagnosed serologically using paired sera from a representative number of sheep. Occasional BDV isolates have been shown to produce high fever, profound and prolonged leukopenia, anorexia, conjunctivitis, nasal discharge, dyspnoea and diarrhoea, and 50% mortality in young lambs. One such isolate was recovered from a severe epidemic of BD among dairy sheep in 1984 (Chappuis *et al.*, 1986). A second such isolate was a BDV contaminant of a live CSEV vaccine (Wensvoort & Terrstra, 1988)

88 live CSFV vaccine (Wensvoort & Terpstra, 1988).

89 2. Fetal infection

The main clinical signs of BD are seen following the infection of pregnant ewes. While the initial maternal infection is 90 subclinical or mild, the consequences for the fetus are serious. Fetal death may occur at any stage of pregnancy, but is 91 92 more common in fetuses infected early in gestation. Small dead fetuses may be resorbed or their abortion may pass unnoticed as the ewes continue to feed well and show no sign of discomfort. As lambing time approaches, the abortion of 93 larger fetuses, stillbirths and the premature births of small, weak lambs will be seen. Confirmation that an abortion or 94 stillbirth is due to BDV is often difficult to establish, but virus may be isolated from fetal tissues in some cases. The use of 95 96 an appropriate real-time reverse-transcription polymerase chain reaction (RT-PCR) assay may give a higher level of 97 success because of the advantages of high sensitivity and the ability to detect genome from non-infectious virus. In aborted fetuses, it is also possible to detect virus by immunohistochemistry of brain, thyroid and other tissues (Thur et al., 1997). 98 Samples of fetal fluids or serum should be tested for BDV antibody. 99

100 During lambing, an excessive number of barren ewes will become apparent, but it is the diseased live lambs that present 101 the main clinical features characteristic of BD. The clinical signs exhibited by BD lambs are very variable and depend on 102 the breed of sheep, the virulence of the virus and the time at which infection was introduced into the flock. Affected lambs are usually small and weak, many being unable to stand. Nervous signs and fleece changes are often apparent. The 103 nervous signs of BD are its most characteristic feature. The tremor can vary from violent rhythmic contractions of the 104 muscles of the hindlegs and back, to barely detectable fine trembling of the head, ears, and tail. Fleece abnormalities are 105 106 most obvious in smooth-coated breeds, which develop hairy fleeces, especially on the neck and back. Abnormal brown or black pigmentation of the fleece may also be seen in BD-affected lambs. Blood samples to be tested for the presence of 107 BDV or antibody should be collected into anticoagulant from suspect lambs before they have received colostrum. Once 108 lambs have ingested colostrum, it is difficult to isolate virus until they are 2 months old and maternal antibody levels have 109 110 waned. However, during this period, it may be possible to detect viral antigen in skin biopsies, by immunohistochemistry, in washed leukocytes by enzyme-linked immunosorbent assay (ELISA) or by real-time RT-PCR. ELISAs directed at 111 detection of the Erns antigen appear to be less prone to interference by maternal antibodies and can often be used to 112 113 detect antigen in serum.

114 With careful nursing, a proportion of BD lambs can be reared, although deaths may occur at any age. The nervous signs 115 gradually decline and can disappear by 3–6 months of age. Weakness, and swaying of the hind-quarters, together with 116 fine trembling of the head, may reappear at times of stress. Affected lambs often grow slowly and under normal field 117 conditions many will die before or around weaning time. In cases where losses at lambing time have been low and no 118 lambs with obvious signs of BD have been born, this can be the first presenting sign of disease.

Some fetal infections occurring around mid-gestation can result in lambs with severe nervous signs, locomotor disturbances and abnormal skeletons. Such lambs have lesions of cerebellar hypoplasia and dysplasia, hydranencephaly and porencephaly resulting from necrotising inflammation. The severe destructive lesions appear to be immune mediated, and lambs with such lesions frequently have high titres of serum antibody to BDV. Most lambs infected in late gestation are normal and healthy and are born free from virus but with BDV antibody. Some such lambs can be weak and may die in early life (Barlow & Patterson, 1982).

125 3. Persistent viraemia

When fetuses survive an infection that occurs before the onset of immune competence, they are born with a persistent 126 viraemia. The ovine fetus can first respond to an antigenic stimulus between approximately 60 and 85 days of its 150-day 127 128 gestation period. In fetuses infected before the onset of immune competence, viral replication is uncontrolled and 50% 129 fetal death is common. In lambs surviving infection in early gestation, virus is widespread in all organs. Such lambs appear 130 to be tolerant of the virus and have a persistent infection, usually for life. A precolostral blood sample will be virus positive and antibody negative. Typically, there is no inflammatory reaction and the most characteristic pathological changes are 131 in the central nervous system (CNS) and skin. Throughout the CNS, there is a deficiency of myelin, and this causes the 132 133 nervous signs. In the skin, primary wool follicles increase in size and the number of secondary wool follicles decreases, causing the hairy or coarse fleece. 134

Persistently viraemic sheep can be identified by the detection of viral antigens, nucleic acids or infectious virus in a blood 135 sample. Viraemia is readily detectable by testing of serum at any time except within the first 2 months of life, when virus 136 may be masked by colostral antibody and, possibly, in animals older than 4 years, some of which develop low levels of 137 anti-BDV antibody (Nettleton et al., 1992). Methods other than virus isolation may be preferred to avoid interference from 138 antibodies. When the presence of colostral antibodies is suspected, the virus may be detected in washed leukocytes and 139 140 in skin by using sensitive ELISAs. Although virus detection in blood during an acute infection is difficult, persistent viraemia should be confirmed by retesting animals after an interval of at least 3 weeks. The use of real-time RT-PCR should be 141 142 considered at all times and for any sample type due to its high analytical sensitivity and the lack of interference from antibodies in a sample. 143

144 Some viraemic sheep survive to sexual maturity and are used for breeding. Lambs born to these infected dams are always

persistently viraemic. Persistently viraemic sheep are a continual source of infectious virus to other animals and their

identification is a major factor in any control programme. Sheep being traded should be screened for the absence of BDV

147 viraemia.

Usually persistently infected (PI) rams have poor quality, highly infective semen and reduced fertility. All rams used for breeding should be screened for persistent BDV infection on a blood sample. Semen samples can also be screened for virus, but virus isolation is much less satisfactory than from blood because of the toxicity of semen for cell cultures. Realtime RT-PCR for detection of pestivirus nucleic acid would usually overcome toxicity problems, and thus this assay should be useful for testing semen from rams.

153 4. Late-onset disease in persistently viraemic sheep

Some PI sheep housed apart from other animals spontaneously develop intractable diarrhoea, wasting, excessive ocular and nasal discharges, sometimes with respiratory distress. At necropsy such sheep have gross thickening of the distal ileum, caecum and colon resulting from focal hyperplastic enteropathy. Cytopathic BDV can be recovered from the gut of these lambs. With no obvious outside source of cytopathic virus, it is most likely that such virus originates from the lamb's own virus pool, similar to what occurs with BVDV. Other PI sheep in the group <u>do-may</u> not develop the disease. This syndrome, which has been produced experimentally and recognised in occasional field outbreaks of BD, has several similarities with bovine mucosal disease (Nettleton *et al.*, 1992).

161

B. DIAGNOSTIC TECHNIQUES

162

Table 1.	. Test methods	available for	diagnosis	of border	disease	and their	purpose

	Purpose								
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	Prevalence of infection – surveillance	Immune status in individual animals or populations post- vaccination			
Identification of the agent ^(a)									
Virus isolation	+	++	++	+++	-	-			
Antigen detection by ELISA	+	++	+++	+++	_	-			
NA detection by real-time RT-PCR	+++	+++	+++	+++	+++	-			
NA detection by ISH	_	_	-	+	_	-			
Detection of immune response									
ELISA	++	++	++	+	++	++			
VN	+++	+++	++	+++	+++	+++			

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; – = not appropriate for this purpose.

ELISA = enzyme-linked immunosorbent assay; IHC = immunohistochemistry; NA = nucleic acid; RT-PCR = reverse-transcription polymerase chain reaction; ISH = *in-situ* hybridisation; VN = virus neutralisation.

^(a)A combination of agent identification methods applied on the same clinical sample is recommended.

168 **1. Identification of the agent**

There is no designated WOAH Reference Laboratory for BDV, but the reference laboratories for BVDV or CSFV will be able to provide advice⁴². One of the most sensitive proven methods for identifying BDV remains virus isolation. However, a broadly reactive real-time RT-PCR assay (preferably pan-pestivirus reactive) will usually provide higher analytical sensitivity than virus isolation, can be used to test samples that are difficult to manage by virus isolation and can be performed in a few hours. Antigen-detection ELISA and immunohistochemical techniques on tissue sections are also valuable methods for identifying BDV-infected animals.

175 **1.1. Virus isolation**

176 It is essential that laboratories undertaking virus isolation have a guaranteed supply of pestivirus-free susceptible cells and bovine serum, or equivalent, that contain no anti-pestivirus activity and no contaminating virus. It is important 177 178 that a laboratory quality assurance programme be in place. Chapter 3.4.7 provides detailed methods for virus isolation 179 in either culture tubes or microplates for the isolation of pestiviruses from sheep or goat samples, including serum, whole blood, semen and tissues. The principles and precautions outlined in that chapter for the selection of cell 180 181 cultures, medium components and reagents are equally relevant to this chapter. Provided proven pan-pestivirus 182 reactive reagents (e.g. monoclonal antibodies [MAbs], primers and probes for real-time RT-PCR) are used for antigen 183 or nucleic acid detection, the principal difference is the selection of appropriate cell cultures.

- 184 BD virus can be isolated in a number of primary or secondary ovine cell cultures (e.g. kidney, testes, lung). Ovine cell lines for BDV growth are rare. Semicontinuous cell lines derived from fetal lamb muscle (FLM), whole embryo (Thabti 185 et al., 2002) or sheep choroid plexus can be useful, but different lines vary considerably in their susceptibility to the 186 virus. Ovine cells have been used successfully for the isolation and growth of BD viruses and BVDV types 1 and 2 187 from sheep. In regions where sheep may become infected with BVD viruses from cattle, a virus isolation system using 188 both ovine and bovine cells could be optimal. However, bovine cells have lower sensitivity for the primary isolation 189 and growth of some BD viruses, so reliance on bovine cells alone is inadvisable. Details of suitable bovine cell cultures 190 are provided chapter 3.4.7. The precautions outlined in that chapter for the establishment of cells and medium 191 192 components that are free from contamination with either pestiviruses or antibodies, and measures to ensure that the cells are susceptible to a wide range of local field strains are equally relevant to systems for detection of BDV. 193
- From live animals, serum is the most frequently used sample to be tested for the presence of infectious virus. 194 However, for difficult cases, the most sensitive way to confirm pestivirus viraemia is to wash leukocytes repeatedly 195 (at least three times) in culture medium before co-cultivating them with susceptible cells in either cell culture tubes or 196 197 microplates. After culture for 5-7 days, the cultures should be frozen and thawed once and an aliquot of diluted culture fluid passaged onto further susceptible cells grown in microplates or on chamber slides to allow antigen detection by 198 immunocytochemistry. Staining for noncytopathic pestiviruses will usually detect virus at the end of the primary 199 200 passage, but to detect slow-growing viruses in poorly permissive cells two passages are desirable. It is recommended that the culture supernatant used as inoculum for the second passage is diluted approximately 1/100 in new culture 201 medium because some high titred field isolates will replicate poorly if passaged undiluted (i.e. at high multiplicity of 202 infection - moi). 203
- Tissues should be collected from dead animals in virus transport medium. In the laboratory, the tissues are ground to give a 10–20% (w/v) suspension, centrifuged to remove debris, and the supernatant passed through 0.45 μm filters. Spleen, lung, thyroid, thymus, kidney, brain, lymph nodes and gut lesions are the best organs for virus isolation.
- Semen can be examined for the presence of BDV, but raw semen is strongly cytotoxic and must be diluted, usually 207 at least 1/10 in culture medium. As the major threat of BDV-infected semen is from PI rams, blood is a more reliable 208 clinical sample than semen for identifying such animals. There are many variations in virus isolation procedures. All 209 should be optimised for maximum sensitivity using a standard reference virus preparation and, whenever possible, 210 recent BDV field isolates. Most of the limitations of virus isolation for the detection of BDV in serum or blood, tissues 211 212 or semen can be overcome by the use of a proven, sensitive pan-pestivirus reactive real-time RT-PCR. Some laboratories screen samples by real-time RT-PCR and undertake virus isolation on positive samples to collect BDV 213 strains for future reference or research purposes. 214
- For specific technical details of virus isolation procedures, including immunoperoxidase staining, refer to chapter 3.4.7.

⁴² Please consult the WOAH Web site: https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3

217 **1.2.** Nucleic acid detection methods

The complete genomic sequences of three BD viruses have been determined and compared with those of other 218 pestiviruses (Becher et al., 1998; Ridpath & Bolin, 1997). Phylogenetic analysis shows BD viruses to be more closely 219 related to CSFV than to BVDV (Becher et al., 2003; Van Rijn et al., 1997; Vilcek & Nettleton, 2006; Vilcek et al., 220 1997). Real-time RT-PCR for diagnosing pestivirus infection is now used widely and a number of formats have been 221 222 described. Real-time RT-PCR assays have the advantages of being able to detect both infectious virus and residual 223 nucleic acid, the latter being of value for investigating abortions and lamb deaths. Furthermore, the presence of virus-224 specific antibodies in a sample will have no adverse effect on the sensitivity of the real-time RT-PCR assay. These assays are also useful for screening semen and, when recommended nucleic acid extraction protocols are followed, 225 are less affected by components of the semen compared with virus isolation. Because of the potential for small 226 ruminants to be infected with genetically different strains of BDV or with strains of BVDV, a proven-pan-pestivirus 227 reactive real-time RT-PCR with proven high sensitivity should be used. To ensure that the genetic spectrum of BDV 228 229 strains is sufficiently covered, it may be necessary to apply a broadly reactive BDV specific real time RT-PCR in 230 parallel to maximise diagnostic sensitivity. Suitable protocols for both nucleic acid extraction as well as the real-time RT-PCR are described in chapter 3.4.7. All precautions to minimise laboratory contamination should be followed 231 232 closelv.

233 After testing samples in a pan-pestivirus reactive assay, samples giving positive results can any level of reactivity should be investigated further by the application of a BDV-specific real-time RT-PCR (Willoughby et al., 2006). It is 234 important to note however that different genotypes of BDV may be circulating in some populations, especially wild 235 ruminants such as chamois and deer, and may be transferred to sheep. An assay that is specific for the detection of 236 237 BDV should be used with some caution as variants or previously unrecognised genotypes may not be detected. hence the value of initially screening samples with a pan-pestivirus reactive real-time RT-PCR. Nevertheless, there 238 are also situations where a pan-pestivirus reactive real-time RT-PCR may have lower analytical sensitivity. 239 Consequently, in any situation where BDV infection is suspected, the application of several diagnostic methods is 240 241 recommended. Maternal serology can also play an important role as negative results should exclude the potential involvement of a pestivirus. 242

243 **1.3. Enzyme-linked immunosorbent assay for antigen detection**

244 ELISAs for the direct detection of pestivirus antigen in blood and tissues of infected animals have proven to be extremely useful for the detection of PI animals and the diagnosis of disease. The first ELISA for pestivirus antigen 245 detection was described for detecting viraemic sheep and was later modified into a double MAb capture ELISA for 246 247 use in sheep and cattle (Entrican et al., 1994). The test is most commonly employed to identify PI viraemic sheep 248 using washed, detergent-lysed blood leukocytes. The sensitivity is close to that of virus isolation and it is a practical method for screening large numbers of blood samples. As with virus isolation, high levels of colostral antibody can 249 250 mask persistent viraemia. The ELISA is more effective than virus isolation in the presence of antibody, but may give false-negative results in viraemic lambs younger than 2 months old. The ELISA is usually not sensitive enough to 251 252 detect acute BDV infections on blood samples. As well as for testing leukocytes, the antigen ELISA can also be used 253 on tissue suspensions, especially spleen, from suspected PI sheep and, as an alternative to immunofluorescence and immunoperoxidase methods, on cell cultures. Several pestivirus ELISA methods have been published but there 254 are at present no commercially available kits that have been fully validated for detecting BDV. Prior to use for 255 regulatory purposes, these kits should be validated in the region where they are to be used to ensure that a wide 256 257 range of field strains of BDV can be detected and that they are suitable for the sample types to be tested.

258 **1.4. Immunohistochemistry**

Viral antigen demonstration is possible in most of the tissues of PI animals (Braun *et al.*, 2002; Thur *et al.*, 1997) although this is not a method that is routinely used for diagnostic purposes. This should be done on acetone-fixed frozen tissue sections (cryostat sections) or paraffin wax embedded samples using appropriate antibodies. Panpestivirus reactive antibodies with NS2-3 specificity are suitable. Tissues with a high amount of viral antigen are brain, thyroid gland, lung and oral mucosa. Skin biopsies have been shown to be useful for *in-vivo* diagnosis of persistent BDV infection.

265 2. Serological tests

Antibody to BDV is usually detected in sheep sera using VN or an ELISA. The less sensitive agar gel immunodiffusion test is not recommended. Control positive and negative reference sera must be included in every test. These should give results within predetermined limits for the test to be considered valid. Single sera can be tested to determine the prevalence of BDV in a flock, region or country. For diagnosis, however, acute and convalescent sera are the best samples for confirming acute BDV infection. Repeat sera from one animal should always be tested alongside each other on the same plate to provide a reliable comparison of titres.

272 **2.1. Virus neutralisation test**

282

283

284

285

286

287

288 289

290 291

292 293

294

295

296

297

298

299

300

301

302 303

304

305

306 307

308

309

310 311

312

317

Due to antigenic diversity among pestiviruses the choice of test virus is difficult (Dekker *et al.*, 1995; Nettleton *et al.*, 1998). No single strain of BDV is ideal. A local strain that gives the highest antibody titre with a range of positive sheep sera should be used.

Because there are few cytopathogenic strains of BDV available, <u>to achieve optimal analytical sensitivity</u>, it is more usual to employ a representative local non-cytopathogenic strain and read the assay after immunoperoxidase staining of the cells. Proven highly sensitive, pestivirus-free sheep cells such as lamb testis or kidney cells are suitable and can be maintained as cryogenically frozen stocks for use over long periods of time. The precautions outlined for selection of pestivirus-free medium components are equally applicable to reagents to be used in VN tests. A recommended procedure follows.

2.1.1. Test procedure

- i) The test sera are heat-inactivated for 30 minutes at 56°C.
- ii) From a starting dilution of 1/4, serial twofold dilutions of the test sera are made in a cell-culture grade flat-bottomed 96-well microtitre plate, using cell culture medium as diluent. For each sample, three or four wells are used at each dilution depending on the degree of precision required. Also, for each sample and at each serum dilution, one well is left without virus to monitor for evidence of sample toxicity that could mimic viral cytopathology or interfere with virus replication. Control positive and negative sera should also be included in each batch of tests.
- iii) An equal volume (e.g. 50 μl) of a stock of BDV containing 100 TCID₅₀ (50% tissue culture infective dose) is added to each well. A back titration of virus stock is also done in some spare wells to check the potency of the virus (acceptance limits 30-80-300 TCID₅₀).
 - iv) The plate is incubated for 1 hour at 37°C.
 - A flask of suitable cells (e.g. ovine testis or kidney cells) is trypsinised and the cell concentration is adjusted to 2 × 10⁵/ml. 100 μl of the cell suspension is added to each well of the microtitre plate.
 - vi) The plate is incubated at 37°C for 4–5 days, either in a 5% CO₂ atmosphere or with the plate sealed.
- vii) The wells are examined microscopically to ensure that there is no evidence of toxicity or cytopathic effect (CPE), then fixed and stained by immunoperoxidase staining using an appropriate MAb. The VN titre for each serum is the dilution at which the virus is neutralised in 50% of the wells. This can be calculated by the Spearman–Kärber or Reed Muench methods. A seronegative animal will show no neutralisation at the lowest dilution <u>of serum</u> (i.e. 1/4), equivalent to a final dilution of 1/8. For accurate comparison of antibody titres, and particularly to demonstrate significant (more than fourfold) changes in titre, samples should be tested in parallel in the same test
 - viii) Occasionally there may be a need to determine whether antibody in a flock is against a virus belonging to a particular *Pestivirus* serogroup. A differential VN test can be used in which sera are titrated out against representative viruses from each of the four *Pestivirus* groups, i.e. BDV, BVDV types 1 and 2, and CSFV. Maximum titre will identify the infecting serotype and the spectrum of cross-reactivity with the other serotypes will also be revealed.

313 2.2. Enzyme-linked immunosorbent assay

An MAb-capture ELISA for measuring BDV antibodies has been described. Two pan-pestivirus MAbs that detect different epitopes on the immunodominant nonstructural protein NS 2/3 are used to capture detergent-lysed cellculture grown antigen. The results correlate qualitatively with the VN test (Fenton *et al.*, 1991).

2.2.1. Antigen preparation

318Use eight 225 cm² flasks of newly confluent FLM cells; four flasks will be controls and four will be319infected. Wash the flasks and infect four with a 0.01-0.1 m.o.i. of Moredun cytopathic BDV. Allow the320virus to adsorb for 2 hours at 37° C. Add maintenance media containing 2% FBS (free from BDV321antibody), and incubate cultures for 4–5 days until CPE is obvious. Pool four control flask322supernatants and separately pool four infected flask supernatants. Centrifuge at 3000 g for32315 minutes to pellet cells. Discard the supernatants. Retain the cell pellets. Wash the flasks with

324 325 326 327 328 329		cont Rep on th infec	In of PBS and repeat the centrifugation step as above. Pool all the control cell pellets in 8 ml PBS and repeat the centrifugation step as above. Pool all the control cell pellets in 8 ml PBS and return 2 ml to each control flask to lyse the remaining attached cells. eat for infected cells. Keep the flasks at 4° C for at least 2 hours agitating the small volume of fluid the cells vigorously every 30 minutes to ensure total cell detachment. Centrifuge the control and cted antigen at 12,000 <i>g</i> for 5 minutes to remove the cell debris. Supernatant antigens are stored 70°C in small aliquots.
330	2.2.2.	Test	t procedure
331 332 333		i)	The two MAbs are diluted to a predetermined dilution in 0.05 M bicarbonate buffer, pH 9.6. All wells of a suitable ELISA-grade microtitre plate (e.g. Nunc maxisorb, Greiner 129b) are coated overnight at 4°C.
334 335		ii)	After washing three times in PBST, a blocking solution of PBST containing 10% horse serum (PBSTH) is added to all wells, which are incubated at 37°C for 1 hour.
336 337 338		iii)	The antigen is diluted to a predetermined dilution in PBSTH and alternate rows of wells are coated with virus and control antigens for 1 hour at 37°C. The plates are then washed three times in PBST before addition of test sera.
339 340		iv)	Test sera are diluted 1/50 in PBSTH and added to duplicate virus and duplicate control wells for 1 hour at 37°C. The plates are then washed three times in PBST.
341 342		v)	Anti-ovine IgG peroxidase conjugate is diluted to a predetermined dilution in PBSTH and added to all wells for 1 hour at 37°C. The plates are washed three times in PBST.
343 344 345 346 347 348 349		vi)	A suitable activated enzyme substrate/chromogen, such as ortho-phenylene diamine (OPD) or tetramethyl benzidine (TMB), is added). After colour development, the reaction is stopped with sulphuric acid and the absorbance read on an ELISA plate reader. The mean value of the two control wells is subtracted from the mean value of the two virus wells to give the corrected absorbance for each serum. Results are expressed as corrected absorbance with reference to the corrected absorbance of known positive and negative sera. Alternatively, ELISA titres can be extrapolated from a standard curve of a dilution series of a known positive reference serum.
350 351 352 353 354		vii)	If antigens of sufficient potency can be produced the MAb capture stage can be omitted. In this case alternate rows of wells are coated with virus and control antigen diluted to a predetermined dilution in 0.05 M bicarbonate buffer, pH 9.6, overnight at +4°C. The plates are washed and blocked as in step ii above. After washing, diluted test sera are added and the test proceeds from step iv as above.

355

C. REQUIREMENTS FOR VACCINES

356 **1. Background**

To be useful, a BDV vaccine should be effective when administered to female sheep before breeding to prevent transplacental infection. Experimental and commercial inactivated whole virus BDV vaccines have been produced in Europe (Brun *et al.*, 1993; Vantsis *et al.*, 1980). Unlike vaccines for BVDV, there is limited demand for vaccines against BDV and those produced have only been inactivated products. No live attenuated or recombinant subunit vaccines for BDV have been produced commercially.

362 Pestivirus contaminants of modified live virus vaccines have been found to be a cause of serious disease following their use in pigs, cattle, sheep and goats. Contaminated vaccines have included those used for the control of Aujesky's disease. 363 CSF, rotavirus, coronavirus, rinderpest, sheep pox and contagious pustular dermatitis. The insidious ability of pestiviruses 364 365 to cross the placenta, and thus establish PI animals, gives them the potential to contaminate vaccines through cells, serum used as medium supplement, or seed stock virus. As nearly all isolates of pestiviruses are noncytopathic, they will remain 366 367 undetected unless specific tests are carried out. Although such contamination should be less likely to be a problem with an inactivated vaccine, nevertheless steps should be taken to ensure that materials used in production are not 368 369 contaminated.

370 **1.1. Characteristics of a target product profile**

Traditionally, pestivirus vaccines fall into two classes: modified live or inactivated virus vaccines. The essential requirement for both types is to afford a high level of fetal infection. Only inactivated vaccines have been produced for BDV. Properly formulated inactivated vaccines are very safe to use but, to obtain satisfactory levels of immunity, they usually require booster vaccinations, which may be inconvenient. Because of the propensity for antigenic variability, the vaccine should contain strains of BDV that are closely matched to viruses found in the area in which they are used. This may present particular challenges with BDV in regions where several antigenic types have been
 found. Due to the need to customise vaccines for the most commonly encountered strains within a country or region,
 it is not feasible to produce a vaccine antigen bank that can be drawn upon globally

379 Guidance for the production of veterinary vaccines is given in Chapter 1.1.8 *Principles of veterinary vaccine* 380 *production*. The guidelines given here and in chapter 1.1.8 are intended to be general in nature and may be 381 supplemented by national and regional requirements.

382 2. Outline of production and minimum requirements for vaccines

383 2.1. Characteristics of the seed

384 An ideal vaccine should contain a strain or strains of virus that give protection against all sheep pestiviruses. This may be challenging however, because of the range of pestiviruses with which sheep can be infected. There is 385 considerable antigenic variation across these viruses - both between viruses that have been classified in the BDV 386 genogroup as well as between viruses in the BVDV1 and BVDV2 genotypes (Becher et al., 2003; Vilcek & Nettleton, 387 2006; Wensvoort et al., 1989). Infection of sheep with the putative BVDV-3 genotype has also been described (Decaro 388 et al., 2012). It is likely that the antigenic composition of a vaccine will vary from region to region to provide an 389 adequate antigenic match with dominant virus strains. Cross-neutralisation studies are required to establish optimal 390 combinations. Nevertheless it would appear that any BDV vaccine should contain at least a representative of the BDV 391 392 and BVDV (type 1) groups. Characterisation of the biologically cloned vaccine viruses should include typing with MAbs and genotyping (Paton et al., 1995). 393

394

411

412

2.1.1. Quality criteria (sterility, purity, freedom from extraneous agents)

- It is crucial to ensure that all materials used in the preparation of the bulk antigens have been 395 396 extensively screened to ensure freedom from extraneous agents. This should include master and 397 working seeds, the cell cultures and all medium supplements such as bovine serum. Some bovine viruses and particularly BVDV can readily infect small ruminants such as sheep. Therefore, it is 398 particularly important to ensure that any serum used that is of bovine origin is free of both adventitious 399 400 BVDV and antibodies against BVDV strains because low levels of either virus or antibody can mask the presence of the other. Materials and vaccine seeds should be tested for sterility and freedom 401 from contamination with other agents, especially viruses as described in the chapter 1.1.8 and 402 403 Chapter 1.1.9 Tests for sterility and freedom from contamination of biological materials intended for veterinary use. 404
- 405If a vaccine passes basic tests, the efficacy of vaccination should ultimately be measured by the
capacity to prevent transplacental transmission. Effective challenge of vaccinated pregnant ewes at
50–60 days gestation has been achieved by intranasal installation of virus or by mixing with PI sheep
(Brun *et al.*, 1993). Usually this reliably produces persistently viraemic offspring in non-immune ewes.
In regions where multiple genotypes of BDV viruses are commonly encountered, efficacy in protecting
against multiple strains should be measured.

2.2. Method of manufacture

2.2.1. Procedure

413 Inactivated vaccines have been prepared using conventional laboratory techniques with stationary or rolled cell cultures. Inactivants have included formalin and beta-propriolactone. Adjuvants have 414 included aluminium hydroxide and oil (Brun et al., 1993; Vantsis et al., 1980). Optimal yields depend 415 on the cell type and isolate used. A commercial BDV vaccine containing two strains of virus has been 416 prepared on ovine cell lines (Brun et al., 1993). Cells must be produced according to a seed-lot 417 system from a master cell seed (MCS) that has been shown to be free from all contaminating 418 microorganisms. Vaccine should only be produced in cells fewer than 20 passages from the MCS. 419 Control cells from every passage should be checked for pestivirus contamination. Standard 420 procedures may be used, with the expectation for harvesting noncytopathic virus on days 4-7 after 421 inoculation of cultures. The optimal yield of infectious virus will depend on several factors, including 422 423 the cell culture, isolate used and the initial seeding rate of virus. These factors should be taken into 424 consideration and virus replication kinetics investigated to establish the optimal conditions for largescale virus production. Whether a live or inactivated vaccine, the essential aim will be to produce a 425 high-titred virus stock. This bulk antigen preparation can subsequently be prepared according to the 426 type of vaccine being considered. 427

428 2.2.2. Requirements for ingredients

BDV vaccines have usually been grown in cell cultures of ovine origin that are frequently supplemented with medium components of animal origin. The material of greatest concern is bovine serum due to the potential for contamination with BVD viruses and antibodies to these viruses. These adventitious contaminants not only affect the efficiency of production but also may mask the presence of low levels of infectious BVDV that may have undesirable characteristics. In addition to the virus seeds, all materials should be tested for sterility and freedom from contamination with other agents, especially viruses as described in chapters 1.1.8 and 1.1.9. Furthermore, materials of bovine or ovine origin should originate from a country with negligible risk for transmissible spongiform encephalopathies (see chapter 1.1.9).

2.2.3. In-process controls

In-process controls are part of the manufacturing process. Cultures should be inspected regularly to ensure that they remain free from gross bacterial contamination, and to monitor the health of the cells and the development or absence of CPE, as appropriate. While the basic requirement for efficacy is the capacity to induce an acceptable neutralising antibody response, during production, target concentrations of antigen required to achieve an acceptable response may be monitored indirectly by assessment of the quantity of infectious virus or antigen mass that is produced. Rapid diagnostic assays such as the ELISA are useful for monitoring BVDV antigen production. Alternatively, the quality of a batch of antigen may be determined by titration of the quantity of infectious virus present, although this may underestimate the quantity of antigen. For inactivated vaccines, infectivity is evaluated before inactivation. For inactivated vaccines the inactivation kinetics should be established so that a suitable safety margin can be determined and incorporated into the routine production processes. At the end of production, *in-vitro* cell culture assays should be undertaken to confirm that inactivation has been complete. These innocuity tests should include a sufficient number of passages and volume of inoculum to ensure that very low levels of infectious virus would be detected if present.

- 453 2.2.4. Final product batch tests
 - i) Sterility

Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9.

457 ii) Identity

Identity tests should demonstrate that no other strain of BDV is present when several strains are propagated in a facility producing multivalent vaccines.

460 iii) Safety

Samples from inactivated vaccines should be tested rigorously for viable virus. Samples of the product should be passaged for a minimum of three passages in sensitive cell cultures to ensure absence of live BDV. This *in-vitro* monitoring can be augmented by injecting two BDV-seronegative sheep with 20 doses of unformulated antigen as part of a standard safety test. Presence of live virus will result in the development of a more convincing serological response than will occur with inactivated virus alone. The sheep sera can also be examined for antibody to other prescribed agents.

Safety tests shall also consist of detecting any abnormal local or systemic adverse reactions to the vaccine by all vaccination route(s). Batch-to-batch safety tests are required unless safety of the product is demonstrated and approved in the registration dossier and production is consistent with that described in chapter 1.1.8. Vaccines must either be demonstrated to be safe in pregnant sheep (i.e. no transmission to the fetus), or should be licensed with a warning not to use them in pregnant animals.

iv) Batch potency

Vaccine potency is best tested in seronegative sheep in which the development and level of antibody is measured. BVD vaccines must be demonstrated to produce adequate immune responses when used in their final formulation according to the manufacturer's published instructions. The minimum quantity of infectious virus or antigen required to produce an acceptable immune response should be determined. An indirect measure of potency is given by the level of virus infectivity prior to inactivation. *In-vitro* assays should be used to monitor individual batches during production. The antigen content following inactivation can be assayed by MAb-capture ELISA and related to the results of established

481 482			<i>in-vivo</i> potency results. It should be demonstrated that the lowest recommended dose of vaccine can prevent transplacental transmission of BDV in pregnant sheep.					
483	2.3.	Requi	Requirements for authorisation/registration/licensing					
484		2.3.1.	Manufacturing process					
485 486 487 488			For registration of a vaccine, all relevant details concerning manufacture of the vaccine and quality control testing should be submitted to the relevant authorities. Unless otherwise specified by the authorities, information should be provided from three consecutive vaccine batches with a volume not less than 1/3 of the typical industrial batch volume.					
489 490 491 492			There is no standard method for the manufacture of a BDV vaccine, but conventional laboratory techniques with stationary, rolled or suspension (micro-carriers) cell cultures may be used. Inactivated vaccines can be prepared by conventional methods, such as binary ethylenimine, formalin or beta-propiolactone inactivation (Park & Bolin, 1987). A variety of adjuvants may be used.					
493		2.3.2.	Safety requirements					
494 495 496 497			<i>In-vivo</i> tests should be undertaken using repeat doses (taking into account the maximum number of doses for primary vaccination and, if appropriate, the first revaccination/booster vaccination) and contain the maximum permitted antigen load and, depending on the formulation of the vaccine, the maximum number of vaccine strains.					
498			i) Target and non-target animal safety					
499 500 501 502			The safety of the final product formulation of inactivated vaccines should be assessed in susceptible young sheep that are free of maternally derived antibodies and in pregnant ewes. They should be checked for any local reactions following administration, and, in pregnant ewes, for any effects on the unborn lamb.					
503			ii) Reversion-to-virulence for attenuated/live vaccines and environmental considerations					
504 505 506 507 508			In the event that a live virus vaccine was developed for BDV, virus seeds that have been passaged at least up to and preferably beyond the passage limit specified for the seed should be inoculated into young lambs to confirm that there is no evidence of disease. If a live attenuated vaccine has been registered for use in pregnant animals, reversion to virulence tests should also include pregnant animals. Live attenuated vaccines should not be transmissible to unvaccinated 'in-contact' animals.					
509			iii) Precautions (hazards)					
510 511 512 513 514 515			BDV is not considered to be a human health hazard. Standard good microbiological practice should be adequate for handling the virus in the laboratory. While the inactivated virus in a vaccine should be identified as harmless for people administering the product, adjuvants included in the vaccine may cause injury to people. Manufacturers should provide adequate warnings that medical advice should be sought in the case of self-injection (including for adjuvants, oil-emulsion vaccine, preservatives, etc.) with warnings included on the product label/leaflet so that the vaccinator is aware of any danger.					
516		2.3.3.	Efficacy requirements					
517 518 519 520 521 522 523 523 524			The potency of the vaccine should be determined by inoculation into seronegative and virus negative lambs, followed by monitoring of the antibody response. Antigen content can be assayed by infectivity titration prior to inactivation and subsequently by ELISA and adjusted as required to a standard level for the particular vaccine. Standardised assay protocols applicable to all vaccines do not exist. Live vaccine batches may be assayed by infectivity titration. Each production batch of vaccine should undergo potency and safety testing as batch release criteria. BVD vaccines must be demonstrated to produce adequate immune responses, as outlined above, when used in their final formulation according to the manufacturer's published instructions.					
525		2.3.4.	Vaccines permitting a DIVA strategy (detection of infection in vaccinated animals)					

2.3.4. Vaccines permitting a DIVA strategy (detection of infection in vaccinated animals)

To date, there are no commercially available vaccines for BDV that support use of a true DIVA 526 strategy. 527

2.3.5. Duration of immunity

Inactivated vaccines are unlikely to provide sustained levels of immunity and it is likely that after an initial course of two or three injections annual booster doses may be required. Insufficient information is available to determine any correlation between vaccinal antibody titres in the dam and fetal protection. As there are likely to be different commercial formulations and these involve a range of adjuvants, there are likely to be different periods of efficacy. Consequently, duration of immunity data must be generated separately for each commercially available product by undertaking challenge tests at the end of the period for which immunity has been claimed.

536 **2.3.6.** Stability

There are no accepted guidelines for the stability of BDV vaccines, but it can be assumed that an inactivated virus vaccine should remain potent for at least 1 year if kept at 4°C and probably longer. Lower temperatures could prolong shelf life but adjuvants in a killed vaccine may preclude this. Bulk antigens that have not been formulated into finished vaccine can be reliably stored frozen at low temperatures, but the antigen quality should be monitored with *in-vitro* assays prior to incorporation into a batch of vaccine.

543

528

529

530

531

532

533

534

535

537

538

539

540

541

542

REFERENCES

544 BARLOW R.M. & PATTERSON D.S.P. (1982). Border disease of sheep: a virus-induced teratogenic disorder. *Adv. Vet. Med.* 545 (*Suppl. J. Vet. Med.*), **36**, 1–87.

BECHER P., AVALOS-RAMIREZ R., ORLICH M., CEDILLO ROSALES S., KONIG, M., SCHWEIZER M., STALDER H., SCHIRRMEIR H & THIEL
 H.-J. (2003). Genetic and antigenic characterisation of novel pestivirus genotypes; Implications for classification. *Virology*,
 311, 96–104.

- 549 BECHER P., ORLICH M. & THIEL H.-J. (1998). Complete genomic sequence of border disease virus a pestivirus from sheep. 550 *J. Virol.*, **72**, 5165–5173.
- 551 BRAUN U., HILBE M., EHRENSPERGER F., SALIS F., ALTHER P., STRASSER M., STALDER H.P. & PETERHANS E. (2002). Border 552 Disease in einem Schafbetrieb. *Schweiz. Arch. Tierheilk.*, **144**, 419–426.

553 BRUN A., LACOSTE F., REYNAUD G., KATO F. & SAINT-MARC B. (1993). Evaluation of the potency of an inactivated vaccine 554 against border disease pestivirus infection in sheep. *In:* Proceedings of the Second Symposium on Pestiviruses, Edwards 555 S., ed. Fondation Marcel Merieux, Annecy, France, 1–3 October 1992, 257–259

556 CARLSSON U. (1991). Border disease in sheep caused by transmission of virus from cattle persistently infected with bovine 557 virus diarrhoea virus. *Vet. Rec.*, **128**, 145–147.

CHAPPUIS G., BRUN A., KATO F., DAUVERGNE M., REYNAUD G. & DURET C. (1986). Etudes serologiques et immunologiques
realisees a la suite de l'isolement d'un pestivirus dans un foyer ovina chez des moutons de L'Aveyron. *In:* Pestiviroses des
Ovins et des Bovins, Espinasse J. & Savey M. eds. Ste Françoise de Buiatrie, Paris, France, **55**, 66.

- DECARO N., MARI V., LUCENTE M., SCIARRETTA R., MORENO A., ARMENISE C., LOSURDO M., CAMERO M., LORUSSO E., CORDIOLI
 P., & BUONAVOGLIA C. (2012). Experimental infection of cattle, sheep and pigs with 'Hobi'-like pestivirus. *Vet. Microbiol.*,
 155, 165–171.
- DEKKER A., WENSVOORT G. & TERPSTRA C. (1995). Six antigenic groups within the genus pestivirus as identified by crossneutralisation assays. *Vet. Microbiol.*, **47**, 317–329.
- 566 ENTRICAN G., DAND A. & NETTLETON P.F. (1994). A double monoclonal-antibody ELISA for detecting pestivirus antigen in 567 the blood of viraemic cattle and sheep. *Vet. Microbiol.*, **43**, 65–74.
- 568 FENTON A., SINCLAIR J.A., ENTRICAN G., HERRING J.A. & NETTLETON P.F. (1991). A monoclonal antibody capture ELISA to 569 detect antibody to border disease virus in sheep sera. *Vet. Microbiol.*, **28**, 327–333.
- 570INTERNATIONALCOMMITTEEONTAXONOMYOFVIRUSES(2016).VirusTaxonomy2015release.571http://www.ictvonline.org/virusTaxonomy.asp

- 572 NETTLETON P.F., GILMOUR J.S., HERRING J.A. & SINCLAIR J.A. (1992). The production and survival of lambs persistently 573 infected with border disease virus. *Comp. Immunol. Microbiol. infect. Dis.*, **15**, 179–188.
- 574 NETTLETON P.F., GILRAY J.A., RUSSO P. & DLISSI E. (1998). Border disease of sheep and goats Vet. Res., 29, 327–340.

575 OGUZOGLU T.C., FLOEGEL-NIESMANN G., FREY H.R. & MOENNIG V. (2001). Differential diagnosis of classical swine fever and 576 border disease: seroepidemiological investigation of a pestivirus infection on a mixed sheep and swine farm. *Dtsch* 577 *Tierarztl. Wochenschr.*, **108**, 210–213.

- 578 PARK B.K. & BOLIN S.R. (1987). Molecular changes of bovine viral diarrhea virus polypeptides treated with binary 579 ethylenimine, beta-propiolactone and formalin. *Res. Rep. Rural Dev. Admin.* (*L*&V) *Korea*, **29**, 99–103.
- PATON D.J., SANDS J.J., LOWINGS J.P., SMITH J.E., IBATA G. & EDWARDS S. (1995). A proposed division of the pestivirus genus
 into subgroups using monoclonal antibodies, supported by cross-neutralization assays and genetic sequencing. *Vet. Res.*,
 26, 92–109.
- 583 RIDPATH J.F. & BOLIN S.R. (1997). Comparison of the complete genomic sequence of the border disease virus, BD31, to 584 other pestiviruses. *Virus Res.*, **50**, 237–243.
- THABTI F., FRONZAROLI L., DLISSI E., GUIBERT J.M., HAMMAMI S., PEPIN M. & RUSSO P. (2002). Experimental model of border
 disease virus infection in lambs: comparative pathogenicity of pestiviruses isolated in France and Tunisia. *Vet. Res.*, 33, 35–45.
- 588 THUR B., HILBE M., STRASSER M. & EHRENSPERGER F. (1997). Immunohistochemical diagnosis of pestivirus infection 589 associated with bovine and ovine abortion and perinatal death. *Am. J. Vet. Res.*, **58**, 1371–1375.
- 590 VALDAZO-GONZALEZ B., ALVAREZ-MARTINEZ M. & SANDVIK T. (2007). Genetic and antigenic typing of border Disease virus 591 isolates in sheep from the Iberian peninsula. *Vet. J.*, **174**, 316–324.
- VAN RIJN P.A., VAN GENNIP H.G.P., LEENCLERSE C.H., BRUSCHKE C.J.M., PATON D.J., MOORMANN R.J.M. & VAN OIRSCHOT J.T.
 (1997). Subdivision of the pestivirus genus based on envelope glycoprotein E2 *Virology*, 237, 337–348.
- VANTSIS J.T., BARLOW R.M., FRASER J. & MOULD D.L. (1976). Experiments in border disease VIII. Propagation and properties
 of a cytopathic virus. *J. Comp. Pathol.*, 86, 111–120.
- 596 VANTSIS J.T., RENNIE J.C., GARDINER A.C., WELLS P.W., BARLOW R.M. & MARTIN W.B. (1980). Immunisation against Border 597 disease. J. Comp. Path., **90**, 349–354.
- 598 VILCEK S. & NETTLETON P.F. (2006). Pestiviruses in wild animals Vet. Microbiol., 116, 1–12.
- VILCEK S., NETTLETON P.F., PATON D.J. & BELAK S. (1997). Molecular characterization of ovine pestiviruses. J. Gen. Virol.,
 78, 725–735.
- 601 WENSVOORT G. & TERPSTRA C. (1988). Bovine viral diarrhoea virus infection in piglets born to sows vaccinated against 602 swine fever with contaminated virus. *Res. Vet. Sci.*, **45**, 143–148.
- 603 WENSVOORT G., TERPSTRA C. & DE KLUYVER E.P. (1989). Characterisation of porcine and some ruminant pestiviruses by 604 cross-neutralisation. *Vet. Microbiol.*, **20**, 291–306.
- 605 WILLOUGHBY K., VALDAZO-GONZALEZ, B., MALEY M., GILRAY J. & NETTLETON P.F. (2006). Development of a real time RT-PCR 606 to detect and type ovine pestiviruses. *J. Virol. Methods*, **132**, 187–194.
- 607
 *

 608
 *

 609
 NB: At the time of publication (2017) there were no WOAH Reference Laboratories

 610
 for border disease (please consult the WOAH Web site:

 611
 https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).

 612
 NB: FIRST ADOPTED IN 1996. MOST RECENT UPDATES ADOPTED IN 2017.

7

SUMMARY

8 Sheep pox and goat pox are <u>contagious</u>, viral diseases of sheep and goats characterised by fever, 9 generalised papules or nodules, vesicles (rarely), internal lesions (particularly in the lungs), and death. Both 10 diseases are caused by strains of capripoxvirus, all of which can infect sheep and goats. Although most of 11 the strains examined cause more severe clinical disease in either sheep or goats, some strains have been 12 isolated that are equally pathogenic in both species.

Sheeppox virus (SPPV) and goatpox virus (GTPV) are the causative agents of sheep pox and goat pox, and with lumpy skin disease virus (<u>LSDV</u>) make up the genus Capripoxvirus in the family Poxviridae. Sheep pox and goat pox are endemic in Africa north of the Equator, the Middle East and Asia, while some parts of Europe have experienced outbreaks recently. <u>See WAHIS (https://wahis.woah.org/#/home) for recent</u> information on distribution at the country level. <u>Countries that reported outbreaks of the disease between</u> 2010 and 2015 include Bulgaria, Chinese Taipei, Israel, Kazakhstan, Kyrgyzstan, Mongolia, Morocco, Greece and Russia, with Greece, Israel and Russia having experienced recurring incidences.

Identification of the agent: Laboratory confirmation of capripoxvirus is most rapid using the polymerase 20 21 chain reaction (PCR) method in combination with a clinical history consistent with generalised capripoxvirus 22 infection. Isolation of the virus is possible as capripoxviruses will grow on tissue culture of ovine, caprine or 23 bovine origin, although field isolates may require up to 14 days to grow or require one or more additional 24 tissue culture passage(s). The virus causes intracytoplasmic inclusions that can be clearly seen using haematoxylin and eosin staining. The antigen can also be detected in tissue culture using specific sera and 25 immunoperoxidase or immunofluorescence techniques. Capripoxvirus antigen and inclusion bodies may be 26 27 seen in stained cryostat or paraffin sections of biopsy or post-mortem lesion material.

An antigen detection enzyme-linked immunosorbent assay (ELISA) using a polyclonal detection serum
 raised against a recombinant immunodominant antigen of capripoxvirus has been developed.

- Serological tests: The virus neutralisation test is the most specific serological test. The indirect immunofluorescence test is less specific due to cross-reactions with antibody to other poxviruses. Western blotting using the reaction between the P32 antigen of capripoxvirus with test sera is both sensitive and specific, but is expensive and difficult to carry out. <u>An enzyme-linked immunosorbent assay (ELISA) has</u> <u>been developed and validated to detect antibodies to capripoxviruses, however it cannot differentiate</u> <u>between SPPV, GTPV and LSDV.</u>
- 36 The use of this antigen, or other appropriate antigens expressed by a suitable vector, in an ELISA offers the 37 prospect of an acceptable and standardised serological test in the future.

Requirements for vaccines: Live and inactivated vaccines have been used for the control of
 capripox<u>viruses</u>. All strains of capripoxvirus so far examined share a major neutralisation site and some will
 cross protect. Inactivated vaccines give, at best, only short-term immunity.

41

A. INTRODUCTION

42 The Capripoxvirus genus, in the family Poxviridae, consists of three species - lumpy skin disease virus (LSDV), which causes disease in cattle only (see Chapter 3.4.12), and sheeppox virus (SPPV) and goatpox virus (GIPPV), which cause 43 sheep pox and goat pox, respectively. Sheep pox and goat pox are characterised by disseminated cutaneous nodules and 44 up to 100% mortality in fully susceptible breeds of sheep and goats. In indigenous animals, generalised disease and 45 mortality are less common, although they are seen where disease has been absent from an area or village for a period of 46 time, when intensive husbandry methods are introduced, or in association with other disease agents, such as peste des 47 petits ruminants virus or foot and mouth disease virus. Sheep pox and goat pox are major constraints to the introduction 48 of exotic breeds of sheep and goats to endemic areas, and to the development of intensive livestock production. 49

Strains of SPPV and GTPV can pass between sheep and goats, although most cause more severe clinical disease in only one their homologous host species. SPPV and GTPV are transboundary diseases that regularly spread into adjacent, nonendemic areas. Sheep pox and goat pox are endemic in Africa north of the Equator and parts of the Middle East and Asia (see WAHIS for most up-to-date information on distribution: https://wahis.woah.org/#/home). Outbreaks have been reported in non-endemic countries of Asia, Europe and the Middle East.

The incubation period of sheep pox and goat pox is between 8 and 13 days following contact between infected and 55 56 susceptible animals. It may be as short as 4 days following experimental infection by intradermal inoculation or mechanical transmission by insects. Some breeds of European sheep, such as Soay, may die of acute infection before the 57 development of skin lesions. In other breeds there is an initial rise in rectal temperature to above 40°C, followed in 2-5 58 days by the development of, at first, macules - small circumscribed areas of hyperaemia, which are most obvious on 59 60 unpigmented skin - and then of papules - hard swellings of between 0.5 and 1 cm in diameter - which may cover the body or be restricted to the groin, axilla and perineum. Papules may be covered by fluid-filled vesicles, but this is rare. Some 61 researchers have distinguished between a vesicular and nodular form of sheep pox and goat pox (Zro et al., 2014b). 62

Within 24 hours of the appearance of generalised papules, affected animals develop rhinitis, conjunctivitis and enlargement of all the superficial lymph nodes, in particular the prescapular lymph nodes. Papules on the eyelids cause blepharitis of varying severity. As the papules on the mucous membranes of the eyes and nose ulcerate, so the discharge becomes mucopurulent, and the mucosae of the mouth, anus, and prepuce or vagina become necrotic. Breathing may become laboured and noisy due to pressure on the upper respiratory tract from the swollen retropharyngeal lymph nodes, due to the developing lung lesions.

If the affected animal does not die in this acute phase of the disease, the papules start to become necrotic from ischaemic necrosis following thrombi formation in the blood vessels at the base of the papule. In the following 5–10 days the papules form scabs, which persist for up to 6 weeks, leaving small scars. The skin lesions are susceptible to fly strike, and secondary pneumonia is common. Anorexia is not usual unless the mouth lesions physically interfere with feeding. Abortion is rare.

On post-mortem examination of the acutely infected animal, the skin lesions are often less obvious than on the live animal. The mucous membranes appear necrotic and all the body lymph nodes are enlarged and oedematous. Papules, which may be ulcerated, can usually be found on the abomasal mucosa, and sometimes on the wall of the rumen and large intestine, on the tongue, hard and soft palate, trachea and oesophagus. Pale areas of approximately 2 cm in diameter may occasionally be seen on the surface of the kidney and liver, and have been reported to be present in the testicles. Numerous hard lesions of up to 5 cm in diameter are commonly observed throughout the lungs, but particularly in the diaphragmatic lobes.

The clinical signs and post-mortem lesions vary considerably with breed of host and strain of capripoxvirus. Indigenous breeds are less susceptible and frequently show only a few lesions, which could be confused with insect bites or contagious pustular dermatitis. However, lambs that have lost their maternally derived immunity, animals that have been kept isolated and animals brought into endemic areas from isolated villages, particularly if they have been subjected to the stress of moving long distances and mixing with other sheep and goats, and their pathogens, can often be seen with generalised and sometimes fatal capripoxvirus infections. Invariably there is high mortality in unprotected imported breeds of sheep and goats following capripoxvirus infection. Capripoxvirus is not infectious to humans.

B. DIAGNOSTIC TECHNIQUES

	Purpose					
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	Prevalence of infection – surveillance	Immune status in individual animals or populations post- vaccination
Identification of the agent ^(a)						
Virus isolation	+	++	+	+++	+	_
Antigen detection	++	++	++	++	++	-
<u>IFAT</u>	<u>+</u>	<u>+</u>	<u>+</u>	<u>++</u>	<u>+</u>	≣
IHC	<u>+</u>	<u>±</u>	<u></u>	<u>++</u>	<u>+</u>	≡
PCR	++	+++	++	+++	++	-
Detection of immune response						
VN <u>T</u>	++	++	++	++	++	++
IFAT	+	+	+	+	+	+
ELISA	<u>++</u>	<u>++</u>	++	<u>++</u>	++	<u>++</u>

Table 1. Test methods available for diagnosis of sheep pox and goat pox and their purpose

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; – = not appropriate for this purpose.

IFAT = indirect fluorescent antibody test; IHC = ; immunohistochemistry; PCR = polymerase chain reaction;

VN<u>T</u> = virus neutralisation; <u>ELISA = enzyme-linked immunosorbent assay</u>.

^(a)A combination of agent identification methods applied on the same clinical sample is recommended.

95 **1. Identification of the agent**

1.1. Specimen collection and submission

97 Material for virus isolation and antigen detection should be collected by biopsy or at post-mortem from skin papules, 98 lung lesions or lymph nodes. Samples for virus isolation and antigen detection enzyme linked immunosorbent assay 99 (ELISA) should be collected within the first week of the occurrence of clinical signs, before the development of 100 neutralising antibodies. Samples for genome detection by polymerase chain reaction (PCR) may be collected before 101 or after the development of neutralising antibody responses. Buffy coat from blood collected into EDTA (ethylene 102 diamine tetra-acetic acid) during the viraemic stage of capripox<u>virus</u> infection (before generalisation of lesions or 103 within 4 days of generalisation), can also be used for virus isolation.

- Samples for histology should include tissue from the surrounding area and should be placed immediately following
 collection into ten times the sample volume of 10% formalin or neutral buffered 10% formal saline. Tissues in formalin
 have no special transportation requirements.
- Blood samples, for virus isolation from the buffy coat, should be collected in tubes containing anticoagulant, placed immediately on ice and processed as soon as possible. In practice, the blood samples may be kept at 4°C for up to 2 days prior to processing, but should not be frozen or kept at ambient temperatures. Tissues and dry scabs for virus isolation, antigen detection and genome detection should preferably be kept at 4°C, on ice or at -20°C. If it is necessary to transport samples over long distances without refrigeration, the medium should contain 10% glycerol; the samples should be of sufficient size that the transport medium does not penetrate the central part of the biopsy, which should be used for virus isolation/detection.

88

89

90

91

92

93

94

96

114 **1.2. Virus isolation**

Lesion material for virus isolation and antigen detection is homogenised. The following is an example of one technique 115 for homogenisation: The tissue is minced using sterile scissors and forceps, and then macerated in a steel ball bearing 116 mixer mill or ground with a sterile pestle in a mortar with sterile sand and an equal volume of sterile phosphate 117 buffered saline (PBS) or serum-free Modified Eagle's Medium (MEM) containing sodium penicillin (1000 international 118 units [IU]/ml), streptomycin sulphate (1 mg/ml), mycostatin (100 IU/ml) or fungizone (2.5 µg/ml) and neomycin 119 120 (200 IU/ml). The homogenised suspension is freeze-thawed three times and then partially clarified by centrifugation 121 using a bench centrifuge at 600 g for 10 minutes. In cases where bacterial contamination of the sample is expected (such as when virus is isolated from skin samples), the supernatant can be filtered through a 0.45 µm pore size filter 122 after the centrifugation step, however, the amount of virus in the supernatant might be reduced. Buffy coats may be 123 prepared from 5-8 ml unclotted blood by centrifugation at 600 g for 15 minutes; the buffy coat is carefully removed 124 into 5 ml of cold double-distilled water using a sterile Pasteur pipette. After 30 seconds, 5 ml of cold double-strength 125 growth medium is added and mixed. The mixture is centrifuged at 600 g for 15 minutes, the supernatant is discarded 126 and the cell pellet is suspended in 5 ml of growth medium, such as Glasgow's modified Eagle's medium (GMEM). 127 After centrifugation at 600 g for a further 15 minutes, the resulting pellet is suspended in 5 ml of fresh GMEM. 128 Alternatively, the buffy coat may be separated from a heparinised sample using a density gradient. 129

130 Capripoxvirus will grow in tissue culture of bovine, ovine or caprine origin, although primary or secondary cultures of lamb testis (LT) or lamb kidney (LK) cells are considered to be the most susceptible. Care needs to be taken to ensure 131 they are not contaminated with viruses such as bovine viral diarrhoea virus, particularly those derived from a wool 132 sheep breed (see chapter 1.1.9). <u>Madin–Darby bovine kidney (MDBK) cells have been shown to be suitable for</u> capripoxvirus isolation (Fay et al., 2020). The following is an example of an isolation technique: either 1 ml of buffy 133 134 coat cell suspension or 1 ml of clarified biopsy preparation supernatant is inoculated on to a 25 cm² tissue culture 135 flask of appropriate cells at 90% confluent LT or LK cells confluence, and the supernatant is allowed to adsorb for 1 136 hour at 37°C. The culture is then washed with warm PBS and covered with 10 ml of a suitable medium, such as 137 138 GMEM, containing antibiotics and 2% fetal calf serum. If available, tissue culture tubes-containing LT or LK cells and a, flying cover-slips, or tissue culture microscope slides, are can also infected. 139

The flasks should be examined daily for 7-14 days for evidence of cytopathic effect (CPE). Contaminated flasks 140 should be discarded. Infected cells develop a characteristic CPE consisting of retraction of the cell membrane from 141 surrounding cells, and eventually rounding of cells and margination of the nuclear chromatin. At first only small areas 142 of CPE can be seen, sometimes as soon as 4 days after infection; over the following 4-6 days these expand to 143 involve the whole cell sheet. If no CPE is apparent by day 7, the culture should be freeze-thawed three times, and 144 clarified supernatant inoculated on to fresh LT or LK-cell cultures. At the first sign of CPE in the flasks, or earlier if a 145 number of infected cover-slips are being used, a cover-slip should be removed, fixed in acetone and stained using 146 147 H&E. Eosinophilic intracytoplasmic inclusion bodies, which are variable in size but up to half the size of the nucleus 148 and surrounded by a clear halo, are indicative of poxvirus infection. Syncytia formation is not a feature of capripoxvirus infection. If the CPE is due to capripoxvirus infection of the cell culture, it can be prevented or delayed by inclusion of 149 specific anti-capripoxvirus serum in the medium; this provides a presumptive identification of the agent. Some strains 150 of capripoxvirus have been adapted to grow on African green monkey kidney (Vero) cells, but these cells are not 151 152 recommended for primary isolation.

153 **1.3. Electron microscopy**

154 The characteristic poxvirus virion can be visualised using a negative-staining preparation technique followed by 155 examination with an electron microscope. There are many different negative-staining protocols, an example is given 156 below:

Material from the original tissue suspension is prepared for transmission electron microscope examination, prior to 157 centrifugation, by floating a 400-mesh hexagon electron microscope grid, with piloform-carbon substrate activated by 158 glow discharge in pentylamine vapour, on to a drop of the suspension placed on parafilm or a wax plate. After 1 159 minute, the grid is transferred to a drop of Tris/EDTA buffer, pH 7.8, for 20 seconds and then to a drop of 1% 160 phosphotungstic acid, pH 7.2, for 10 seconds. The grid is drained using filter paper, air-dried and placed in the 161 electron microscope. The capripoxvirus virion is brick shaped, covered in short tubular elements and measures 162 approximately 290 × 270 nm. A host-cell-derived membrane may surround some of the virions, and as many as 163 possible should be examined to confirm their appearance (Kitching & Smale, 1986). 164

165 The virions of capripoxvirus are indistinguishable from those of orthopoxvirus, but, apart from *Vaccinia* virus, no 166 orthopoxvirus causes lesions in sheep and goats. However, capripoxvirus is distinguishable from the virions of 167 parapoxvirus, that cause contagious pustular dermatitis, as they are smaller, oval in shape, and each is covered in a 168 single continuous tubular element, which appears as striations over the virion.

169 **1.4. Histopathology**

Material for histopathology should be prepared by standard techniques. Following preparation, staining with 170 haematoxylin and eosin (H&E), and mounting of the formalin-fixed biopsy material, a number of sections should be 171 examined by light microscopy. On histological examination, the most striking aspects of acute-stage skin lesions are 172 a massive cellular infiltrate, vasculitis and oedema. Early lesions are characterised by marked perivascular cuffing. 173 Initially infiltration is by macrophages, neutrophils and occasionally eosinophils, and as the lesion progresses, by 174 175 more macrophages, lymphocytes and plasma cells. A characteristic feature of all capripoxvirus infections is the 176 presence of variable numbers of 'sheep pox cells' in the dermis. These sheep pox cells can also occur in other organs where microscopic lesions of sheep and goat pox are present. These cells are large, stellate cells with eosinophilic, 177 poorly defined intracytoplasmic inclusions and vacuolated nuclei. Vasculitis is accompanied by thrombosis and 178 179 infarction, causing oedema and necrosis. Epidermal changes consist of acanthosis, parakeratosis and hyperkeratosis. Changes in other organs are similar, with a predominant cellular infiltration and vasculitis. Lesions in 180 181 the upper respiratory tract are characterised by ulceration.

182 **1.5. Immunological methods**

183

184

185 186

187

188

189 190

198

199

200

201

202 203

208

209

210

211

212 213

214 215

216

217

218

219

1.5.1. Fluorescent antibody tests

Capripoxvirus antigen can also be identified on infected cover-slips or tissue culture slides using fluorescent antibody tests. Cover-slips or slides should be washed and air-dried and fixed in cold acetone for 10 minutes. The indirect test using immune sheep or goat sera is subject to high background colour and nonspecific reactions. However, a direct conjugate can be prepared from sera from convalescent sheep or goats or from rabbits hyperimmunised with purified *Capripoxvirus*. Uninfected tissue culture should be included as a negative control because cross-reactions, due to antibodies to cell culture antigens, can cause problems. The fluorescent antibody tissue section technique has also been used on cryostat-prepared slides.

191 **1.6. Nucleic acid recognition methods**

Amplification methods for detection of the viral DNA genome are specific to the genus *Capripoxvirus* DNA are and both specific and sensitive for detection throughout the course of disease, including before and after the emergence of antibody responses. These methods include conventional PCR, real-time PCR, and most recently loop-mediated isothermal amplification (<u>LAMP</u>). Nucleic acid recognition methods can be used to detect the *Capripoxvirus* genome in biopsy, swab or tissue culture samples.

197 **1.6.1. Conventional PCR methods**

- Several conventional PCR methods have been reported with varying specificity for capripoxviruses in general, SPPV, or GTPV (Heine *et al.*, 1999; Ireland & Binepal, 1998; Zro *et al.*, 2014a). <u>A</u> <u>conventional PCR assay that differentiates GTPV and LSDV from SPPV has been described (Lamien <u>et al.</u>, 2011a). Conventional PCR methods are particularly useful for obtaining sufficient genetic material necessary for species identification by subsequent sequence and phylogenetic analysis (Le Goff *et al.*, 2009).</u>
- 204
 The conventional gel-based PCR method described below is a simple, fast and sensitive method for

 205
 the detection of capripoxvirus genome in EDTA blood, semen or tissue culture samples (Tuppurainen

 206
 et al., 2005).

207 <u>Test procedure</u>

- The extraction method described below can be replaced using commercially available DNA extraction kits.
 - i) <u>Freeze and thaw 200 µl of blood in EDTA, semen or tissue culture supernatant and suspend in</u> <u>100 µl of lysis buffer containing 5 M guanidine thiocyanate, 50 mM potassium chloride, 10 mM</u> <u>Tris/HCl (pH 8); and 0.5 ml Tween 20.</u>
 - ii) <u>Cut skin and other tissue samples into fine pieces using a sterile scalpel blade and forceps.</u> <u>Grind with a pestle in a mortar. Suspend the tissue samples in 800 µl of the above mentioned</u> <u>lysis buffer.</u>
- iii) Add 2 µl of proteinase K (20 mg/ml) to blood samples and 10 µl of proteinase K (20 mg/ml) to tissue samples. Incubate at 56°C for 2 hours or overnight, followed by heating at 100°C for 10 minutes. Add phenol:chloroform:isoamylalcohol (25:24:1 [v/v]) to the samples in a 1:1 ratio. Vortex and incubate at room temperature for 10 minutes. Centrifuge the samples at 16,060 g

000			for 15 minutes at 4° C. Carefully collect the upper aqueous phase (up to 200 µl) and trapefor
220 221			for 15 minutes at 4°C. Carefully collect the upper, aqueous phase (up to 200 µl) and transfer into a clean 2.0 ml tube. Add two volumes of ice cold ethanol (100%) and 1/10 volume of 3 M
222			sodium acetate (pH 5.3). Place the samples at -20°C for 1 hour. Centrifuge again at 16,060 g
223			for 15 minutes at 4°C and discard the supernatant. Wash the pellets with ice cold 70% ethanol
224			(100 μ) and centrifuge at 16,060 g for 1 minute at 4°C. Discard the supernatant and dry the
225			pellets thoroughly. Suspend the pellets in 30 µl of nuclease-free water and store immediately at
226			-20°C (Tuppurainen et al., 2005). Alternatively a column-based extraction kit may be used.
227		iv)	The primers for this PCR assay were developed from the gene encoding the viral attachment
228		,	protein. The size of the expected amplicon is 192 bp (Ireland & Binepal, 1998). The primers
229			have the following gene sequences:
230			Forward primer 5'-TCC-GAG-CTC-TTT-CCT-GAT-TTT-TCT-TAC-TAT-3'
231			Reverse primer 5'-TAT-GGT-ACC-TAA-ATT-ATA-TAC-GTA-AAT-AAC-3'.
232		V)	DNA amplification is carried out in a final volume of 50 µl containing: 5 µl of 10 × PCR buffer,
233		•)	1.5 µl of MgCl ₂ (50 mM), 1 µl of dNTP (10 mM), 1 µl of forward primer, 1 µl of reverse primer,
234			1 µl of DNA template (~10 ng), 0.5 µl of Taq DNA polymerase and 39 µl of nuclease-free water.
235			The volume of DNA template required may vary and the volume of nuclease-free water must
236			<u>be adjusted to the final volume of 50 µl.</u>
237		vi)	Run the samples in a thermal cycler as follows: 2 minutes at 95°C; then 45 seconds at 95°C,
238			50 seconds at 50°C and 1 minute at 72°C (34 cycles); 2 minutes at 72°C and hold at 4°C until
239			analysis.
240		vii)	Mix 10 µl of each sample with loading dye and load onto a 1.5% agarose gel in TAE buffer
241			(Tris/acetate buffer containing EDTA). Load a parallel lane with a 100 bp DNA-marker ladder.
242 243			<u>Electrophoretically separate the products using approximately 8–10 V/cm for 40–60 minutes</u> and visualise with a suitable DNA stain and transilluminator.
240			
244	1.6.2.		I-time PCR methods
245		Seve	eral highly sensitive and specific fluorescent detection-based real-time PCR methods have been
246		deve	loped and validated (Balinsky <i>et al.</i> , 2008; Bowden <i>et al.</i> , 2008; Das <i>et al.</i> , 2012; Stubbs <i>et al.</i> ,
247		deve 2012	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these
247 248		deve 2012 meth	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these loods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct
247		deve 2012 meth capri	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these
247 248 249 250 251		deve 2012 meth capri desc	P). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these hods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct pox <u>virus genotyping species differentiation without</u> the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; <u>Wolff <i>et al.</i>, 2021</u>). real-time PCR method described below is a rapid, sensitive and specific method for the detection
247 248 249 250 251 252		deve 2012 meth capri desc <u>The r</u> of DN	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipox <u>virus</u> genotyping <u>species differentiation without</u> the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; <u>Wolff <i>et al.</i>, 2021</u>).
247 248 249 250 251		deve 2012 meth capri desc <u>The r</u> of DN	P). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these hods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct pox <u>virus</u> genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; <u>Wolff <i>et al.</i>, 2021</u>). real-time PCR method described below is a rapid, sensitive and specific method for the detection
247 248 249 250 251 252 253 254		deve 2012 meth capri desc <u>The I</u> of DN <u>DNA</u> <u>A nu</u>	P). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genetyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-
247 248 249 250 251 252 253 254 255		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nu</u> time	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genetyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate
247 248 249 250 251 252 253 253 254 255 256		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nu</u> time meth	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genetyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate and for the sample type being extracted. WOAH Reference Laboratories can be contacted for
247 248 249 250 251 252 253 254 255		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nu</u> time meth	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genetyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate
247 248 249 250 251 252 253 253 254 255 256		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nu</u> time <u>meth</u> advic	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate to for the sample type being extracted. WOAH Reference Laboratories can be contacted for ce on suitable commercial kits.
247 248 249 250 251 252 253 254 255 256 257 258 259		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nu</u> time <u>meth</u> advic	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate to for the sample type being extracted. WOAH Reference Laboratories can be contacted for ce on suitable commercial kits. time PCR
247 248 249 250 251 252 253 254 255 256 257 258 259 260		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nur</u> time <u>meth</u> advic	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate tod for the sample type being extracted. WOAH Reference Laboratories can be contacted for ce on suitable commercial kits. time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden et al. (2008). and further validated by Stubbs et al. (2012). Cycling conditions and reagent
247 248 249 250 251 252 253 254 255 256 257 258 259		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nur</u> time <u>meth</u> advic	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate to for the sample type being extracted. WOAH Reference Laboratories can be contacted for ce on suitable commercial kits. time PCR
247 248 249 250 251 252 253 254 255 256 257 258 259 260		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nur</u> time <u>meth</u> advic	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these lods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate tod for the sample type being extracted. WOAH Reference Laboratories can be contacted for ce on suitable commercial kits. time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden et al. (2008). and further validated by Stubbs et al. (2012). Cycling conditions and reagent
247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262		deve 2012 meth capri desc <u>The I</u> of DN <u>DNA</u> <u>A nuu</u> time <u>meth</u> advic i)	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these hods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate od for the sample type being extracted. WOAH Reference Laboratories can be contacted for be on suitable commercial kits. time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden <i>et al.</i> (2008). and further validated by Stubbs <i>et al.</i> (2012). Cycling conditions and reagent concentrations can be altered to ensure optimal performance in individual laboratories. Forward and reverse primers should be prepared at concentrations of 20 µM. A minor grove
247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263		deve 2012 meth capri desc <u>The r</u> of Dt <u>DNA</u> <u>A nur</u> time <u>meth</u> advic i)	P) Each test detects a small conserved genetic locus within the capripoxvirus genome, but these hods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipox <u>virus genotyping species differentiation without</u> the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011b; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate and for the sample type being extracted. WOAH Reference Laboratories can be contacted for the consultable commercial kits. time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden <i>et al.</i> (2008), and further validated by Stubbs <i>et al.</i> (2012). Cycling conditions and reagent concentrations can be altered to ensure optimal performance in individual laboratories. Forward and reverse primers should be prepared at concentrations of 20 µM. A minor grove binder (MGB) TaqMan probe should be prepared at a concentration of 10 µM.
247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264		deve 2012 meth capri desc <u>The I</u> of Dt <u>DNA</u> <u>A nui</u> time <u>meth</u> advic i) ii)	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these hods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct poxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011 <u>b</u> ; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate od for the sample type being extracted. WOAH Reference Laboratories can be contacted for the consultable commercial kits. time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden <i>et al.</i> (2008), and further validated by Stubbs <i>et al.</i> (2012). Cycling conditions and reagent concentrations can be altered to ensure optimal performance in individual laboratories. Forward and reverse primers should be prepared at concentrations of 20 µM. A minor grove binder (MGB) TaqMan probe should be prepared at a concentration of 10 µM.
247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264 265		deve 2012 meth capri desc <u>The r</u> of Df <u>DNA</u> <u>A nut</u> time <u>meth</u> advic Real i) ii) ii)	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these hods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. Lextraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate od for the sample type being extracted. WOAH Reference Laboratories can be contacted for the consultable commercial kits. Lime PCR The real-time PCR method outlined below uses the primers and probe described by Bowden et al. (2008), and further validated by Stubbs et al. (2012). Cycling conditions and reagent concentrations can be altered to ensure optimal performance in individual laboratories. Forward and reverse primers should be prepared at a concentration of 10 µM. Forward primer: 5'-AAA-TGA-AAC-CAA-TGG-ATG-GGA-TA-3'
247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264 265 266 265		deve 2012 meth capri desc <u>The r</u> of Df <u>DNA</u> <u>A nut</u> time <u>meth</u> advic i) ii) ii) ii) v) v) v) vi)	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these looks do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye et al., 2013; Lamien et al., 2011b; Wolff et al., 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. Lextraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate od for the sample type being extracted. WOAH Reference Laboratories can be contacted for ze on suitable commercial kits. time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden et al. (2008). and further validated by Stubbs et al. (2012). Cycling conditions and reagent concentrations can be altered to ensure optimal performance in individual laboratories. Forward and reverse primers should be prepared at concentrations of 20 µM. A minor grove binder (MGB) TaqMan probe should be prepared at a concentration of 10 µM. Forward primer: 5'-AAA-ACG-GTA-TAT-GGA-ATA-GAG-TTG-GAA-3' Reverse primer: 5'-AAA-TGA-AAC-CAA-TGG-ATG-GGA-TA-3' Probe: 5'-FAM-TGG-CTC-ATA-GAT-TTC-CT-MGB-3' Mastermix is prepared by combining 10 µl of 2 × real-time PCR mastermix with 0.4 µl of forward primer, 0.4 µl of reverse primer, 0.5 µl of probe and 6.7 µl of RNase free water per reaction.
247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264 265 266 267		deve 2012 meth capri desc <u>The r</u> of Df <u>DNA</u> <u>A nut</u> time <u>meth</u> advic i) ii) ii) ii) v) v) v) vi)	2). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these looks do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct ipoxvirus genotyping species differentiation without the need for gene sequencing have been ribed (Gelaye <i>et al.</i> , 2013; Lamien <i>et al.</i> , 2011b; Wolff <i>et al.</i> , 2021). real-time PCR method described below is a rapid, sensitive and specific method for the detection NA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species. extraction from blood and tissue mber of DNA extraction kits are commercially available for the isolation of template DNA for real-PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate of of the sample type being extracted. WOAH Reference Laboratories can be contacted for ac on suitable commercial kits. -time PCR The real-time PCR method outlined below uses the primers and probe described by Bowden <i>et al.</i> (2008). and further validated by Stubbs <i>et al.</i> (2012). Cycling conditions and reagent concentrations can be altered to ensure optimal performance in individual laboratories. Forward and reverse primers should be prepared at concentration of 10 µM. Forward primer: 5'-AAA-ACG-GTA-TAT-GGA-ATA-GAG-TTG-GAA-3' Reverse primer: 5'-AAA-TGA-AAC-CAA-TGG-ATG-GGA-TA-3' Probe: 5'-FAM-TGG-CTC-ATA-GAT-TTC-CT-MGB-3' Mastermix is prepared by combining 10 µl of 2 × real-time PCR mastermix with 0.4 µl of forward

272 273

271

274

278 279

280

281

282

283

284

295

296

297

298

299

300

301 302

303

304

305 306

307

308 309

310

311

312 313

314 315

316

317 318

319

320

321

322

323

275 276

277

1.6.3. Isothermal genome amplification

negative.

ix)

Molecular tests using loop mediated isothermal amplification (LAMP) to detect capripoxvirus genomes are reported to provide sensitivity and specificity similar to real-time PCR with a simpler method and at lower cost (Das *et al.*, 2012; Murray *et al.*, 2013). Field validation of the Das *et al.* (2012) LAMP method has been further reported by (Omoga *et al.*, 2016) and a combination of this universal capripoxvirus test with two additional LAMP assays was reported to show utility in discriminating between GTPV and SPPV (Zhao *et al.*, 2014).

285 2. Serological tests

286 **2.1. Virus neutralisation**

287 A test serum can either be titrated against a constant titre of capripoxvirus (100 TCID₅₀ [50% tissue culture infective 288 dose]) or a standard capripoxvirus strain can be titrated against a constant dilution of test serum in order to calculate a neutralisation index. Because of the variable sensitivity of tissue culture to capripoxvirus, and the consequent 289 290 difficulty of ensuring the use of 100 TCID₅₀, the neutralisation index is the preferred method, although it does require 291 a larger volume of test sera. The test is described using 96-well flat-bottomed tissue culture grade microtitre plates, but it can be performed equally well in tissue culture tubes with the appropriate changes to the volumes used, although 292 293 it is more difficult to read an end-point in tubes. The use of Vero cells in the virus neutralisation test has been reported to give more consistent results (Kitching & Taylor, 1985). 294

2.1.1. Test procedure

i) Test sera including a negative and a positive control are diluted 1/5 in Eagle's/HEPES (N-2hydroxyethylpiperazine, N-2-ethanesulphonic acid) and inactivated at 56°C for 30 minutes.

viii) 95°C for 10 minutes followed by 45 cycles of 95°C for 15 seconds, 60°C for 60 seconds.

Following completion of the real-time PCR, a cycle threshold (C_T) should be set. Samples with

 C_T values less than 35 are considered positive. Samples with a C_T value greater than 35 but

less than 45 are considered inconclusive and require further investigation. Samples which do

not yield a C_T value, i.e. the amplification curve does not cross the threshold, are considered

Fluorescence detection should be performed at the end of each cycle.

- ii) Next, 50 µl of the first inactivated serum is added to columns 1 and 2, rows A to H of the microtitre plate. The second serum is placed in columns 3 and 4, the third in columns 5 and 6, the positive control serum is placed in columns 7 and 8, the negative control serum is placed in columns 9 and 10, and 50 µl of Eagle's/HEPES without serum is placed in columns 11 and 12 and to all wells of row H.
 - iii) A reference strain of capripoxvirus, usually a vaccine strain known to grow well in tissue culture, with a titre of over log₁₀ 6 TCID₅₀ per ml is diluted in Eagle's/HEPES in bijoux bottles to give a log dilution series of log₁₀ 5.0; 4.0; 3.5; 3.0; 2.5; 2.0; 1.5 TCID₅₀ per ml (equivalent to log₁₀ 3.7; 2.7; 2.2; 1.7; 1.2; 0.7; 0.2 TCID₅₀ per 50 μl).
 - iv) Starting with row G and the most diluted virus preparation, 50 µl of virus is added to each well in that row. This is repeated with each virus dilution, the highest titre virus dilution being placed in row A.
 - v) The plates are covered and incubated for 1 hour at 37°C.
- vi) LT cells are <u>An appropriate cell suspension (such as MDBK cells)</u> is prepared from pregrown monolayers as a suspension of 10⁵ cells/ml in Eagle's medium containing antibiotics and 2% fetal calf serum. Following incubation of the microtitre plates, 100 µl of cell suspension is added to all the wells, except wells H11 and H12, which serve as control wells for the medium. The remaining wells of row H are cell and serum toxicity controls.
 - vii) The microtitre plates are covered and incubated at 37°C for 9 days.
- viii) Using an inverted microscope, the monolayers are examined daily starting at day 4 for evidence of CPE. There should be no CPE in the cells of row H. Using the 0240 KSGP vaccine strain of capripoxvirus, the final reading is taken on day 9, and the titre of virus in each duplicate titration is calculated according to the Kärber method. If left longer, there is invariably a 'breakthrough' of virus in which virus that was at first neutralised appears to disassociate from the antibody.
 - ix) Interpretation of the results: The neutralisation index is the log titre difference between the titre of the virus in the negative serum and in the test serum. An index of ≥1.5 is positive. The test

324 325 326 327	can be made more sensitive if serum from the same animal is examined before and after infection. Because immunity to capripox <u>virus</u> is predominantly cell mediated, a negative result, particularly following vaccination in which the response is necessarily mild, does not imply that the animal from which the serum was taken is not protected.
328	A constant virus/varying serum method has been described using serum dilutions in the range
329	1/5 to 1/500 and fetal calf muscle cells. Because these cells have a lower sensitivity to

capripoxvirus than LT cells, the problem of virus 'breakthrough' is overcome.

331 **2.2.** Indirect fluorescent antibody test

Capripoxvirus-infected tissue culture grown on flying cover-slips or tissue culture microscope slides can be used for the indirect fluorescent antibody test. Uninfected tissue culture control, and positive and negative control sera, should be included in the test. The infected and control cultures are fixed in acetone at -20°C for 10 minutes and stored at 4°C. Dilutions of test sera are made in PBS, starting at 1/5, and positives are identified using an anti-sheep gammaglobulin conjugated with fluorescein isothiocyanate (Davies & Otema, 1978). Cross-reactions can occur with orf, bovine papular stomatitis virus and perhaps other poxviruses.

338 2.3. Western blot analysis

330

339 Western blotting of test sera against capripoxvirus-infected cell lysate provides a sensitive and specific system for 340 the detection of antibody to capripoxvirus structural proteins, although the test is expensive and difficult to carry out 341 (Chand *et al.*, 1994).

342 **2.4. Enzyme-linked immunosorbent assay**

- 343 No validated ELISA is available for the serological diagnosis of SPP or GTP.
- 344Both in-house and commercial enzyme-linked immunosorbent assay (ELISAs) are available, but these tests cannot345discriminate between antibodies to different capripoxviruses (LSD or SPP/GTP).

346 C. REQUIREMENTS FOR VACCINES 347 [THIS SECTION IS UNDER REVIEW IN THE 2024/2025 REVIEW CYCLE]

348 **1. Background**

349 **1.1. Rationale and intended use of the product**

A variety of attenuated live and inactivated capripoxvirus vaccines has been used to provide protection against sheeppox and goatpox. All strains of capripoxvirus of ovine, caprine or bovine origin examined so far share a major neutralising site, so that animals recovered from infection with one strain are resistant to infection with any other strain (Capstick, 1961). Consequently, it is possible to use a single strain of capripoxvirus to protect both sheep and goats against all field strains of virus, regardless of whether their origin was in Asia or Africa (Kitching *et al.*, 1986; Kitching & Taylor, 1985). However, field evidence suggests some strains are quite host-specific and are used only in sheep against SPPV and only in goat against GTPV.

A number of strains of capripoxvirus have had widespread use as live vaccines (Davies & Mbugwa, 1985), for example the Romanian and RM-65 strains used mainly in sheep and the Mysore and Gorgan strains used in goats. The real identity of the commonly used Kenyan sheep and goat pox vaccine virus (KSGP) 0240 was recently shown to be actually LSDV (Tuppurainen *et al.*, 2014). Virus strain identity and attenuation properties must be ascertained and taken into consideration when selecting vaccine strains for use in cattle, sheep and goats. The protective dose depends on the vaccine strain used. Immunity in sheep and goats against capripox<u>virus</u> following vaccination with the 0240 strain lasts over a year and the Romanian strain gave protection for at least 30 months.

Killed vaccines produced from tissue culture contain only the intracellular mature virion form of the virus, and lack the
 less robust but biologically crucial extracellular enveloped virion form. As a result, the vaccine does not stimulate
 immunity against the extracellular enveloped virion, resulting in poor protection. Killed capripox<u>virus</u> vaccines provide,
 at best, only temporary protection.

2. Outline of production and minimum requirements for conventional vaccines 368

General requirements set for the facilities used for the production of vaccines and for the documentation and record keeping 369 370 throughout the whole manufacturing process are described in Chapter 1.1.8 Principles of veterinary vaccine production. The documentation should include the standard operating procedures (SOP) for the method of manufacture and each step 371 for the testing of cells and reagents used in the process, each batches and the final product. 372

373 2.1. Characteristics of the seed

374

380

381

382

383

384

385 386

387 388

392 393

394

395

413

414

2.1.1. **Biological characteristics**

375 A strain of capripoxvirus used for vaccine production must be accompanied by a history describing its origin and tissue culture or animal passage. It must be safe to use in all breeds of sheep and goats 376 for which it is intended, including pregnant and young animals. It must be non-transmissible, remain 377 attenuated after further tissue culture passage, and provide complete protection against challenge 378 with virulent field strains for a minimum of 1 year. A quantity of master seed vaccine virus should be 379 prepared and stored in order to provide a consistent working seed for regular vaccine production.

2.1.2. Quality criteria (sterility, purity, freedom from extraneous agents)

Each master seed must be tested to ensure its identity and shown to be free from adventitious viruses, in particular pestiviruses, such as border disease and bovine viral diarrhoea virus, and free from contamination with bacteria, fungi and/or mycoplasmas. The general procedures for sterility or purity tests are described in chapter 1.1.9. The master seed must also be safe and produce no clinical reaction in all breeds of sheep or goats when given by the recommended route and stimulate complete immunity to capripoxvirus in all breeds of sheep and goats for at least 1 year. The necessary safety and potency tests are described in Section C.2.2.4 Final product batch tests.

2.2. Method of manufacture 389

- The method of manufacture should be documented as the Outline of Production. 390
- 2.2.1. Procedure 391

Vaccine seed should be lyophilised and stored in 2 ml vials at -20°C. It may be stored wet at -20°C, but when wet, is more stable at -70°C or lower. The virus should be cultured in primary or secondary LT or LK cells of wool sheep origin for maximum yield. Vero cells may also be used with suitably adapted strains.

for certain viral vaccines because its mode of action cannot be guaranteed to be totally effective in

inactivating all the live virus. This has not been fully investigated for capripoxvirus.

396 397 398 399 400 401 402 403	Vaccine batches are produced on fresh monolayers of secondary LT or primary LK cells. A vial of seed virus is reconstituted with GMEM or another appropriate medium and inoculated on to an LT or LK monolayer that has been previously washed with warm PBS, and allowed to adsorb for 15 minutes at 37°C before being overlaid with additional GMEM. After 4–6 days, there will be extensive (80–90%) CPE. The culture should be examined for any evidence of nonspecific CPE, medium cloudiness or change in medium pH. The culture is freeze–thawed three times, the suspension removed and centrifuged at 600 g for 20 minutes. A second passage may be required to produce sufficient virus for a production batch. Live vaccine may be produced on roller bottles.
404	The procedure is repeated and the harvests from individually numbered flasks are each mixed
405	separately with an equal volume of sterile and chilled 5% lactalbumin hydrolysate and 10% sucrose,
406	and transferred to individually numbered bottles for storage at -20° C. Prior to storage, 0.2 ml is
407	removed from each bottle for sterility control. An additional 0.2 ml is removed for virus titration; 2 ml
408	pools composed of 0.2 ml samples taken from ten bottles are used. A written record of all the
409	procedures must be kept for all vaccine batches.
410	Inactivated vaccines are produced, usually from unattenuated field strains of capripoxvirus, grown in
411	tissue culture as described above, inactivated with 0.03% formaldehyde, and mixed with an equal
412	volume of alhydrogel as adjuvant. Formaldehyde is no longer considered to be a suitable inactivant

2.2.2. Requirements for substrate and media

- The specification and source of all ingredients used in the manufacturing procedure should be documented and the freedom from extraneous agents: bacteria, fungi, mycoplasma and any other viruses should be tested. The detailed testing procedure is described in the chapter 1.1.9. The use of antibiotics must meet the requirements of the licensing authority.
- 420 2.2.3. In-process controls
 - i) Cells

Cells should be obtained from the testis or kidney of a healthy young lamb from a scrapie-free flock of a wool sheep breed. During cultivation, cells must be observed for any evidence of CPE, and for normal morphology (predominantly fibroblastic). They can usually be passaged successfully up to ten times. When used for vaccine production, uninfected control cultures should be grown in parallel and maintained for at least three additional passages for further observation. They should be checked for the presence of noncytopathic strains of bovine virus diarrhoea or border disease viruses by immunofluorescence or immunoperoxidase techniques. If possible, cells should be prepared and screened prior to vaccine production and stocked in 1–2 ml aliquots containing 2 × 10⁷ cells/ml in sterile 10% DMSO (dimethyl sulphoxide) and 90% FBS (fetal bovine serum) solution stored in liquid nitrogen.

432 ii) Serum

Bovine serum used in the growth or maintenance medium must be free from transmissible spongiform encephalopathies (TSEs) and antibody to capripoxvirus, and tested for contamination with pestivirus or any other viruses, extraneous bacteria, mycoplasma or fungi.

iii) Medium

Medium must be tested free from contamination with pestivirus or any other viruses, extraneous bacteria, mycoplasma or fungi.

439 iv) Virus

Seed virus and final vaccine must be titrated in tissue culture tubes or microtitre plates. Vaccine samples must be examined for the presence of adventitious viruses including cytopathic and noncytopathic strains of pestivirus, and should be mixed with a high-titre capripoxvirus-immune serum that has tested negative for antibody to pestivirus to prevent the vaccine virus itself interfering with the test. The vaccine bulk can be held at -20° C or below until all sterility tests and titrations have been completed, at which time it should be freeze-dried in 1 ml aliquots in vials sufficient for 100 doses. The vaccine harvest diluted with lactalbumin hydrolysate and sucrose should have a minimum titre log₁₀ 4.5 TCID₅₀ per ml after freeze-drying, equivalent to a field dose of log₁₀ 2.5 TCID₅₀. A further titration is carried out on five randomly chosen vials of the freeze-dried preparation to confirm the titre.

- 450 2.2.4. Final product batch tests
- 451 i) Sterility/purity

i) Sterility/purity

Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9.

ii) Safety

The safety studies should be demonstrated by statistically valid vaccination studies using seronegative young sheep and goats of known susceptibility to capripox virus. The procedure described is suitable for vaccine strains such as 0240 that are equally immunogenic in both sheep and goats. The choice of target animal should be adapted for strains with a more restricted host preference.

460 iii) Potency

461Potency tests must be undertaken if the minimum immunising dose of the virus strain is not known.462This is usually carried out by comparing the titre of a virulent challenge virus on the flanks of463vaccinated and control animals. Following vaccination, the flanks of at least three animals and three464controls are shaved of wool or hair. Log₁₀ dilutions of the challenge virus are prepared in sterile PBS465and six dilutions are inoculated intradermally (0.1 ml per inoculum) along the length of the flank; four

466 467 468 469 470 471 472 473	2.3.	Requi	replicates of each dilution are inoculated down the flank. An oedematous swelling will develop at possibly all 24 inoculation sites on the control animals, although preferably there will be little or no reaction at the four sites of the most dilute inocula. The vaccinated animals should develop an initial hypersensitivity reaction at sites of inoculation within 24 hours, which should quickly subside. Small areas of necrosis may develop at the inoculation site of the most concentrated challenge virus. The macule/papule is measured at between 8 and 10 days post-challenge. The titre of the challenge virus is calculated for the vaccinated and control animals; a difference of log ₁₀ titre > 2.5 is taken as evidence of protection.
		2.3.1.	Safety requirements
475		2.3.1.	
476			i) Target and non-target animal safety
477 478 479			The vaccine must be safe to use in all breeds of sheep and goats for which it is intended, including young and pregnant animals. It must also be non-transmissible, remain attenuated after further tissue culture passage.
480			Safety tests should be carried out on the final product of each batch as described in Section C.2.2.4.
481 482 483			The safety of the vaccine in non-target animals must have been demonstrated using mice and guinea-pigs as described in Section C.2.2.4. There should be no evidence of pathology caused by the vaccine.
484			ii) Reversion-to-virulence for attenuated/live vaccines
485			The selected final vaccine should not revert to virulence during a further passages in target animals.
486			iii) Environmental consideration
487 488 489 490			Attenuated vaccine should not be able to perpetuate autonomously in cattle, sheep or goat populations. Vaccines using the 0240 strain should not be used in <i>Bos taurus</i> breeds. Strains of capripoxvirus are not a hazard to human health. There are no precautions other than those described above for sterility and freedom from adventitious agents.
491		2.3.2.	Efficacy requirements
492			i) For animal production
493 494			The efficacy of the vaccine must be demonstrated in vaccination challenge experiment under laboratory conditions. As described in Section C.2.2.4.
495 496 497			Once the potency of the particular strain being used for vaccine production has been determined in terms of minimum dose required to provide immunity, it is not necessary to repeat this on the final product of each batch, provided the titre of virus present has been ascertained.
498			ii) For control and eradication
499			Vaccination is the only effective way to control the sheep pox and goat pox outbreaks in endemic
500 501			countries. Unfortunately, currently no marker vaccines allowing the differentiation of infected from vaccinated animals are available.
502 503			Immunity to virulent field virus following vaccination of sheep or goats with the 0240 strain lasts over 1 year, and protection against generalised infection following intradermal challenge lasts at least
504 505			3 years and is effective lifelong. The duration of immunity produced by other vaccine strains should be ascertained in both sheep and goats by undertaking controlled trials in an environment in which
506			there is no possibility of field strains of capripoxvirus confusing the results. The inactivated vaccines
507 508			provide immunity for less than 1 year, and for the reasons given at the beginning of this section, may not give immunity to the form of capripoxvirus usually associated with natural transmission.
509		2.3.3.	Stability
510 511 512			All vaccines are initially given a shelf-life of 24 months before expiry. Real-time stability studies are then conducted to confirm the appropriateness of the expiry date. Multiple batches of the vaccine should be re-titrated periodically throughout the shelf-life to determine the vaccine variability.

- 513Properly freeze-dried preparations of capripox vaccine, particularly those that include a protectant,514such as sucrose and lactalbumin hydrolysate, are stable for over 25 years when stored at -20°C and515for 2-4 years when stored at 4°C. There is evidence that they are stable at higher temperatures, but516no long-term controlled experiments have been reported. The inactivated vaccines must be stored at5174°C, and their shelf- life is usually given as 1 year.
- 518 No preservatives other than a protectant, such as sucrose and lactalbumin hydrolysate, are required 519 for the freeze-dried preparation.

520 3. Vaccines based on biotechnology

521 3.1. Vaccines available and their advantages

522 Currently, no recombinant vaccines for capripoxviruses are commercially available. However, a new generation of 523 capripox vaccines is being developed that uses the capripoxvirus genome as a vector for the genes of other ruminant 524 pathogens such as peste des petits ruminants (PPR) virus (Berhe *et al.*, 2003; Tuppurainen *et al.*, 2014).

525 **3.2.** Special requirements for biotechological vaccines, if any

- 526 Not applicable.
- 527

REFERENCES

- 528 BALINSKY C.A, DELHON G, SMOLIGA G, PRARAT M, FRENCH R.A, GEARY S.J, ROCK D.L & RODRIGUEZ L.L. (2008). Rapid 529 preclinical detection of sheep pox virus by a real-time PCR assay. *J. Clin. Microbiol.*, **46**, 438–442.
- BERHE G., MINET C., LE GOFF C., BARRETT T., NGANGNOU A., GRILLET C., LIBEAU G., FLEMING M., BLACK D.N. & DIALLO A.
 (2003). Development of a dual recombinant vaccine to protect small ruminants against peste-des-petits-ruminants virus and capripoxvirus infections. J. Virol., 77, 1571–1577.
- 533 BOWDEN T.R, BABIUK S.L, PARKYN G.R., COPPS J.S. & BOYLE D.B. (2008). Capripox virus tissue tropism and shedding: A 534 quantitative study in experimentally infected sheep and goats. *Virology*, **371**, 380–393.
- 535 CAPSTICK P.B. (1961). Annual Report. Kenya Veterinary Department, Kenya, 45–47.
- 536 CHAND P., KITCHING R.P. & BLACK D.N. (1994). Western blot analysis of virus-specific antibody responses to capripoxvirus 537 and contagious pustular dermatitis infections in sheep. *Epidemiol. Infect.*, **113**, 377–385.
- 538 DAS A., BABIUK S. & MCINTOSH M.T. (2012). Development of a loop-mediated isothermal amplification assay for rapid 539 detection of capripoxviruses. *J. Clin. Microbiol.*, **50**, 1613–1620.
- 540 DAVIES F.G. & MBUGWA G. (1985). The alterations in pathogenicity and immunogenicity of a Kenya sheep and goat pox 541 virus on serial passage in bovine foetal muscle cell cultures. *J. Comp. Pathol.*, **95**, 565–576.
- 542 DAVIES F.G. & OTEMA C. (1978). The antibody response in sheep infected with a Kenyan sheep and goat pox virus. *J.* 543 *Comp. Pathol.*, **88**, 205–210.
- 544 FAY P.C., COOK C.G., WIJESIRIWARDANA N., TORE G., COMTET L., CARPENTIER A., SHIH B., FREIMANIS G., HAGA I.R. & BEARD
 545 P.M. (2020). Madin–Darby bovine kidney (MDBK) cells are a suitable cell line for the propagation and study of the bovine
 546 poxvirus lumpy skin disease virus. J. Virol. Methods, 285, 113943. doi: 10.1016/j.jviromet.2020.113943.
- 547 GELAYE E., LAMIEN C.E., SILBER R., TUPPURAINEN E.S., GRABHERR R. & DIALLO A.(2013). Development of a cost-effective 548 method for capripoxvirus genotyping using snapback primer and dsDNA intercalating dye. *PLoS One*, **8** (10): e75971.
- 549 HEINE H.G., STEVENS M.P., FOORD A.J. & BOYLE D.B. (1999). A capripoxvirus detection PCR and antibody ELISA based on 550 the major antigen P32, the homolog of the vaccinia virus H3L gene. *J. Immunol. Methods*, **227**, 187–196.

IRELAND D.C. & BINEPAL Y.S. (1998). Improved detection of capripoxvirus in biopsy samples by PCR. J. Virol. Methods, 74, 1–7.

- 553 KITCHING R.P., HAMMOND J.M. & TAYLOR W.P. (1986). A single vaccine for the control of capripox infection in sheep and 554 goats. *Res. Vet. Sci.*, **42**, 53–60.
- 555 KITCHING R.P. & SMALE C. (1986). Comparison of the external dimensions of capripoxvirus isolates. *Res. Vet. Sci.*, **41**, 425– 556 427.
- 557 KITCHING R.P. & TAYLOR W.P. (1985). Clinical and antigenic relationship between isolates of sheep and goat pox viruses. 558 *Trop. Anim. Health Prod.*, **17**, 64–74.

LAMIEN C.E., GOFF C.L., SILBER R., WALLACE D.B., GULYAZ V., TUPPURAINEN E., MADANI H., CAUFOUR P., ADAM T., EL HARRAK
 M., LUCKINS A.G., ALBINA E. & DIALLO A. (2011a). Use of the Capripoxvirus homologue of Vaccinia virus 30kDa RNA
 polymerase subunit (RPO30) gene as a novel diagnostic and genotyping target: development of a classical PCR method
 to differentiate Goat poxvirus from Sheep poxvirus. *Vet. Microbiol.*, **149**, 30–39. doi: 10.1016/j.vetmic.2010.09.038.

- LAMIEN C.E., LELENTA M., GOGER W., SILBER R., TUPPURAINEN E., MATIJEVIC M., LUCKINS A.G. & DIALLO A. (2011<u>b</u>). Real time
 PCR method for simultaneous detection, quantitation and differentiation of capripoxviruses. *J. Virol. Methods*, **171**, 134–
 140.
- Le GOFF C., LAMIEN C.E., FAKHFAFH E., CHADEYRAS A., ABU-ADULUGBAD E., LIBEAU G., TUPPURAINEN E., WALLACE D., ADAM T.,
 SILBER R., GULYAZ V., MADANI H., CAUFOUR P., HAMAMMI S., DIALLO A. & ALBINA E. (2009). Capripoxvirus G-protein-coupled
 chemokine receptor, a host-range gene suitable for virus-animal origin discrimination. *J. Gen. Virol.*, **90**, 67–77.
- 569 MURRAY L., EDWARDS L., TUPPURAINEN E.S., BACHANEK-BANKOWSKA K., OURA C.A., MIOULET V. & KING D.P. (2013). Detection 570 of capripoxvirus DNA using a novel loop-mediated isothermal amplification assay. *BMC Vet. Res.*, **9**, 90.
- 571 OMOGA D.C.A., MACHARIA M., MAGIRI E., KINYUA J., KASIITI J. & HOLTON T. (2016) Molecular based detection, validation of a 572 LAMP assay and phylogenetic analysis of capripoxvirus in Kenya. *J. Adv. Biol. Biotech.*, **7**, 1–12.
- 573 STUBBS S., OURA C.A., HENSTOCK M., BOWDEN T.R., KING D.P. & TUPPURAINEN E.S. (2012). Validation of a high-throughput 574 real-time polymerase chain reaction assay for the detection of capripoxviral DNA. *J. Virol. Methods*, **179**, 419–422.
- TUPPURAINEN E.S.M., PEARSON C.R., BACHANEK-BANKOWSKA K., KNOWLES N.J., AMAREEN S., FROST L., HENSTOCK M.R.,
 LAMIEN C.E., DIALLO A. & MERTENS P.P.C. (2014). Characterization of sheep pox virus vaccine for cattle against lumpy skin
 disease virus. *Antiviral Res.*, **109**, 1–6.
- 578 <u>TUPPURAINEN E.S.M., VENTER E.H. & COETZER J.A.W. (2005). The detection of lumpy skin disease virus in samples of</u> 579 <u>experimentally infected cattle using different diagnostic techniques. *Onderstepoort J. Vet. Res.*, **72**, 153–164.</u>
- 580 <u>WOLFF J., & BEER M. & Hoffmann B. (2021). Probe-Based Real-Time qPCR Assays for a Reliable Differentiation of Capripox</u>
 581 <u>Virus Species. *Microorganisms*, 9, 765. https://doi.org/10.3390/microorganisms9040765</u>
- ZHAO Z., FAN B., WU G., YAN X., LI Y., ZHOU X., YUE H., DAI X., ZHU H., TIAN B., LI J. & ZHANG Q. (2014) Development of loop mediated isothermal amplification assay for specific and rapid detection of differential goat Pox virus and Sheep Pox virus.
 BMC Microbiol., **14**, 1–10.
- ZRO K., AZELMAT S., BENDOURO Y., KUHN J.H., EL FAHIME E. & ENNAJI M.M. (2014a). PCR-based assay to detect sheeppox
 virus in ocular, nasal, and rectal swabs from infected Moroccan sheep. *J. Virol. Methods*, **204**, 38–43.
- 587 ZRO K., ZAKHAM F., MELLOUL M., EL FAHIME E. & MUSTAPHA M. (2014b). A sheeppox outbreak in Morocco: isolation and 588 identification of virus responsible for the new clinical form of disease. *BMC Vet Res.*, **10**, 31.

589 * *
 590 * * *
 591 592 NB: There are WOAH Reference Laboratories for sheep pox and goat pox (please consult the WOAH Web site: https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).
 593 593 Please contact the WOAH Reference Laboratories for any further information on diagnostic tests, reagents and vaccines for sheep pox and goat pox
 595 NB: FIRST ADOPTED IN 1989. MOST RECENT UPDATES ADOPTED IN 2017.

Annex 16. Item 5.1. – Chapter 3.9.1. African swine fever (infection with African swine fever virus)
MEETING OF THE BIOLOGICAL STANDARDS COMMISSION
Paris, 4–8 September 2023
SECTION 3.9.
SUIDAE
CHAPTER 3.9.1.

AFRICAN SWINE FEVER (INFECTION WITH AFRICAN SWINE FEVER VIRUS)

SUMMARY

African swine fever (ASF) is an infectious disease of domestic and wild pigs of all breeds and ages, caused by ASF virus (ASFV). The clinical syndromes vary from peracute, acute, subacute to chronic, depending on the virulence of the virus. Acute disease is characterised by high fever, haemorrhages in the reticuloendothelial system, and a high mortality rate. Soft ticks of the Ornithodoros genus, especially O. moubata and O. erraticus, have been shown to be both reservoirs and transmission vectors of ASFV. The virus is present in tick salivary glands and passed to new hosts (domestic or wild suids) when feeding. It can be transmitted sexually between ticks, transovarially to the eggs, or transtadially throughout the tick's life.

18 ASFV is the only member of the Asfarviridae family, genus Asfivirus.

10

- 19 Laboratory diagnostic procedures for ASF fall into two groups: detection of the virus and serology. The 20 selection of the tests to be carried out depends on the disease situation and laboratory diagnostic capacity 21 in the area or country.
- 22Identification of the agent: Laboratory diagnosis must be directed towards isolation of the virus by23inoculation of pig leukocyte or bone marrow cultures, the detection of antigen in smears or cryostat sections24of tissues by fluorescent antibody test and/or the detection of genomic DNA by the polymerase chain reaction25(PCR) or real-time PCR. The PCRs are excellent, highly sensitive, specific and rapid techniques for ASFV26detection and are very useful under a wide range of circumstances. They are especially useful if the tissues27are unsuitable for virus isolation and antigen detection. In doubtful cases, the material is passaged in28leukocyte cell cultures and the procedures described above are repeated.
- Serological tests: Pigs that survive natural infection usually develop antibodies against ASFV from 7–10 days post-infection and these antibodies persist for long periods of time. Where the disease is endemic, or where a primary outbreak is caused by a strain of low or moderate virulence, the investigation of new outbreaks should include the detection of specific antibodies in serum or extracts of the tissues submitted. A variety of methods such as the enzyme-linked immunosorbent assay (ELISA), the indirect fluorescent antibody test (IFAT), the indirect immunoperoxidase test (IPT), and the immunoblotting test (IBT) is available for antibody detection.

36 37

38

Requirements for vaccines: At present, there is no vaccine for ASF. <u>Commercially produced modified live</u> virus vaccines are available and licenced in some countries.

A. INTRODUCTION

The current distribution of African swine fever (ASF) extends across more than 50 countries in three continents (Africa, Asia and Europe). Several incursions of ASF out of Africa were reported between the 1960s and 1970s. In 2007, ASF was introduced into Georgia, from where it spread to neighbouring countries including the Russian Federation. From there ASF spread to eastern European countries extending westwards and reaching the European Union in 2014. Further westward and southern spread in Europe has occurred since that time. In all these countries, both hosts – domestic pig and wild boar – were affected by the disease. In August 2018, the People's Republic of China reported its first outbreak of ASF and further spread in Asia has occurred.

46 ASF virus (ASFV) is a complex large, enveloped DNA virus with icosahedral morphology. It is currently classified as the 47 only member of the Asfaviridae family, genus Asfivirus (Dixon et al., 2005). More than 60 structural proteins have been 48 identified in intracellular virus particles (200 nm) (Alejo et al., 2018). More than a hundred infection-associated proteins have been identified in infected porcine macrophages, and at least 50 of them react with sera from infected or recovered 49 pigs (Sánchez-Vizcaíno & Arias, 2012). The ASFV double-stranded linear DNA genome comprises between 170 and 50 51 193 kilobases (kb) and contains between 150 and 167 open reading frames with a conserved central region of about 125 kb and variable ends. These variable regions encode five multigene families that contribute to the variability of the virus 52 genome. The complete genomes of several ASFV strains have been sequenced (Bishop et al., 2015; Chapman et al., 53 2011; de Villiers et al., 2010; Portugal et al., 2015). Different strains of ASFV vary in their ability to cause disease, but at 54 55 present there is only one recognised serotype of the virus detectable by antibody tests.

56 The molecular epidemiology of the disease is investigated by sequencing of the 3' terminal end of the B646L open reading frame encoding the p72 protein major capsid protein, which differentiates up to 24 distinct genotypes (Achenbach et al., 57 2017; Boshoff et al., 2007; Quembo et al. 2018). To distinguish subgroups among closely related ASFV, sequence analysis 58 59 of the tandem repeat sequences (TRS), located in the central variable region (CVR) within the B602L gene (Gallardo et al., 2009; Lubisi et al., 2005; Nix et al., 2006) and in the intergenic region between the I73R and I329L genes, at the right 60 end of the genome (Gallardo et al., 2014), is undertaken. Several other gene regions such as the E183L encoding p54 61 protein, the CP204L encoding p30 protein, and the protein encoded by the EP402R gene (CD2v), have been proved as 62 useful tools to analyse ASFVs from different locations and hence track virus spread. 63

ASF viruses produce a range of syndromes varying from peracute, acute to chronic disease and subclinical infections. Pigs are the only domestic animal species that is naturally infected by ASFV. European wild boar and feral pigs are also susceptible to the disease, exhibiting clinical signs and mortality rates similar to those observed in domestic pigs. In contrast African wild pigs such as warthogs (*Phacochoerus aethiopicus*), bush pigs (*Potamochoerus porcus*) and giant forest hogs (*Hylochoerus meinertzhageni*) are resistant to the disease and show few or no clinical signs. These species of wild pig act as reservoir hosts of ASFV in Africa (Costard *et al.*, 2013; Sánchez-Vizcaíno *et al.*, 2015).

70 The incubation period is usually 4-19 days. The more virulent strains produce peracute or acute haemorrhagic disease characterised by high fever, loss of appetite, haemorrhages in the skin and internal organs, and death in 4-10 days, 71 sometimes even before the first clinical signs are observed. Case fatality rates may be as high as 100%. Less virulent 72 73 strains produce mild clinical signs - slight fever, reduced appetite and depression - which can be readily confused with many other conditions in pigs and may not lead to suspicion of ASF. Moderately virulent strains are recognised that induce 74 75 variable disease forms, ranging from acute to subacute. Low virulence, non-haemadsorbing strains can produce subclinical non-haemorrhadic infection and seroconversion, but some animals may develop discrete lesions in the lungs or on the 76 skin in areas over bony protrusions and other areas subject to trauma. Animals that have recovered from either acute, 77 78 subacute or chronic infections may potentially become persistently infected, acting as virus carriers. The biological basis for the persistence of ASFV is still not well understood, nor it is clear what role persistence plays in the epidemiology of 79 80 the disease.

ASF cannot be differentiated from classical swine fever (CSF) by either clinical or post-mortem examination, and both diseases should be considered in the differential diagnosis of any acute febrile haemorrhagic syndrome of pigs. Bacterial septicaemias may also be confused with ASF and CSF. Laboratory tests are essential to distinguish between these diseases.

In countries free from ASF but suspecting its presence, the laboratory diagnosis must be directed towards isolation of the virus by the inoculation of pig leukocyte or bone marrow cultures, the detection of genomic DNA by polymerase chain reaction (PCR) or the detection of antigen in smears or cryostat sections of tissues by direct fluorescent antibody test (FAT). Currently the PCR is the most sensitive technique and can detect ASFV DNA from a very early stage of infection in tissues, ethylene diamine tetra-acetic acid (EDTA)-blood and serum samples. The PCR is particularly useful if samples 90 submitted are unsuitable for virus isolation and antigen detection because they have undergone putrefaction. Pigs that 91 have recovered from acute, subacute or chronic infections usually exhibit a viraemia for several weeks making the PCR 92 test a very useful tool for the detection of ASFV DNA in pigs infected with low or moderately virulent strains. Virus isolation 93 by the inoculation of pig leukocyte or bone marrow cultures and identification by haemadsorption tests (HAD) are 94 recommended as a confirmatory test when ASF is positive by other methods, particularly in the event of a primary outbreak 95 or a case of ASF.

96 As no vaccine is available, the presence of ASEV antibodies is indicative of previous infection and, as antibodies are 97 produced from the first week of infection and persist for long periods, they are a good marker for the diagnosis of the 98 disease, particularly in subacute and chronic forms.

99 Vaccines should be prepared in accordance with Chapter 1.1.8 Principles of veterinary vaccine production. ASF modified live virus (MLVs) vaccines are based on the live virus that have been naturally attenuated or attenuated by targeted genetic 100 recombination through cell cultures (Gladue & Borca, 2022). MLV production is based on a seed-lot system consistent with 101 the European Pharmacopoeia (11th edition) and that has been validated with respect to virus identity, sterility, purity, 102 potency, safety, non-transmissibility, stability and immunogenicity. ASF MLV first generation vaccines - defined as those 103 104 for which peer-reviewed publications are in the public domain - should meet or exceed the minimum standards as 105 described below. Paramount demonstration of acceptable safety and efficacy against the epidemiologically relevant ASFV 106 field strain(s) where the vaccine is intended for use are required. At the present time, acceptable efficacy should be shown 107 against the B646L (p72) genotype II pandemic virus lineage currently circulating widely in domestic pigs and wild boar.

ASF MLV first generation vaccines allowing the differentiation of infected animals from vaccinated animals (DIVA) by
 suitable methods (e.g. serology-based tests) are preferred. Demonstration of MLV safety and efficacy in breeding-age
 boars, gilts and pregnant sows, and onset and duration of protective immunity, are also preferred but are not required to
 meet the minimum standard.

ASF epidemiology is complex with different epidemiological patterns of infection occurring in Africa and Europe. ASF occurs through transmission cycles involving domestic pigs, wild boar, wild African suids, and soft ticks (Sánchez-Vizcaíno *et al.*, 2015). In regions where *Ornithodoros* soft-bodied ticks are present, the detection of ASFV in these reservoirs of infection contributes to a better understanding of the epidemiology of the disease. This is of major importance in establishing effective control and eradication programmes (Costard *et al.*, 2013).

117 ASF is not a zoonotic disease and does not affect public health (Sánchez-Vizcaíno *et al.*, 2009).

118 ASFV should be handled with an appropriate level of bio-containment, determined by risk analysis in accordance with 119 Chapter 1.1.4 *Biosafety and biosecurity: Standard for managing biological risk in the veterinary laboratory and animal* 120 *facilities.*

121

122

C. REQUIREMENTS FOR VACCINES

. . .

123 At present there is no commercially available vaccine for ASF.

124 <u>**1. Background**</u>

125 <u>The ASF p72 genotype II strain (ASFV Georgia 2007/1 lineage) (NCBI, 2020) is recognised to be the current highest global</u>
 126 <u>threat for domestic pig production worldwide (Penrith *et al.*, 2022).
</u>

127 Guidelines for the production of veterinary vaccines are given in Chapter 1.1.8 *Principles of Veterinary Vaccine Production*.
 128 Varying additional requirements relating to quality (including purity and potency), safety, and efficacy will apply in particular
 129 countries or regions for manufacturers to comply with local regulatory requirements.

Wherever live, virulent ASFV or ASF MLVs are stored, handled and disposed, the appropriate biosecurity level, procedures
 and practices should be used. The ASF MLV vaccine production facility should meet the requirements for containment
 outlined in Chapter 1.1.4 Biosafety and biosecurity: Standard for managing biological risk in the veterinary laboratory and
 animal facilities.

An optimal ASF MLV first generation vaccine for the target host should have the following general characteristics (minimum
 standards):

- Safe: demonstrate absence of fever and clinical signs of acute or chronic ASF in vaccinated and in-contact animals, minimal and ideally no vaccine virus transmission, and absence of an increase in virulence (genetic and phenotypic stability);
- Efficacious: protects against mortality, reduces acute disease (fever accompanied by the appearance of clinical signs caused by ASF) and reduces vertical (boar semen and placental) and horizontal disease transmission;
- Quality purity: free from wild-type ASFV and extraneous microorganisms that could adversely affect the safety, potency or efficacy of the product;
- Quality potent: the log₁₀ virus titre maintained throughout the vaccine shelf life that guarantees the efficacy demonstrated by the established minimum immunising (protective) dose.
- Identity: based on the capacity to protect against the ASFV B646L (p72) genotype II pandemic strain or other p72 genotypes of recognised epidemiologic importance.
- 147 <u>Vaccine production should be carried out using a validated, controlled and consistent manufacturing process.</u>
- ASF MLV first generation vaccines must be safe (i.e. an acceptable safety profile) for non-target species and the
 environment in general.
- Ideally, ASF MLV first generation vaccines that meet the minimum standards should also fulfil the following additional
 general characteristics: i) prevents acute and persistent (carrier state) disease; ii) prevents horizontal and vertical disease
 transmission; iii) induces rapid protective immunity (e.g. < 2 weeks); and iv) confers stable, life-long immunity.

Furthermore, ASF MLV second and future generation vaccines should meet the minimum safety and efficacy standards
 as ASF MLV first generation vaccines, and ideally provide additional product profile benefits, including but not limited to: i)
 contain a negative marker allowing the differentiation of infected from vaccinated animals (DIVA) by reliable discriminatory
 tests such as serology-based tests; and ii) confer broad range of protection against other p72 genotype field strains of
 varying virulence (low, moderate, and high).

The majority of ASF global vaccine research groups and companies are currently focused on ASF MLV first generation
 vaccine candidates that are safe and efficacious against ASF viruses belonging to the ASFV p72 genotype II pandemic
 strain (ASFV Georgia 2007/1 lineage) (NCBI, 2020).

- 161 <u>Currently, two gene deleted MLV recombinant vaccines (ASFV-G-ΔI177L and ASFV-G-ΔMGF) have been licenced for</u>
 162 <u>field use in Vietnam following supervised field testing to evaluate the safety and effectiveness of several vaccine batches.</u>
- 163 <u>There are numerous, promising ASF MLV vaccine candidates targeting the p72 genotype II pandemic strain under</u> 164 <u>development, including:</u>
- A naturally attenuated field strain (Lv17/WB/Rei1) (Barasona *et al.*, 2019) being developed as an oral bait vaccine for wild boars;
- 167 <u>A laboratory thermo-attenuated field strain (ASFV-989) (Bourry et al., 2022);</u>
- <u>Single gene-deleted, recombinant viruses (e.g. SY18ΔI226R, ASFV-G-ΔA137R) (Gladue *et al.*, 2021; Zhang *et al.*, 2021);
 <u>2021)</u>;
 </u>
- Double gene-deleted, recombinant viruses (e.g. ASFV-G-Δ9GL/ΔUK; ASFV-SY18-ΔCD2v/UK; Arm-ΔCD2v-ΔA238L)
 (O'Donnell *et al.*, 2016; Pérez-Núñez *et al.*, 2022; Teklue *et al.*, 2020);
- Multiple gene-deleted, recombinant viruses (ASFV-G-ΔI177L/ΔLVR; ASFV-G-ΔMGF; BA71ΔCD2; HLJ/18-7GD; ASFVGZΔI177LΔCD2vΔMGF) (Borca *et al.*, 2021; Chen *et al.*, 2020; Liu *et al.*, 2023; Monteagudo *et al.*, 2017; O'Donnell *et al.*, 2015).
- 175 Information regarding many of these MLV vaccine candidates can be found in a recent review publication (Brake, 2022).
- 176 Different DIVA strategies using serological methods (e.g. ELISA) or genome detection methods (e.g. differential real-time
- 177 PCR) are not widely available for these ASF MLV first generation vaccine candidates. Therefore, there is still room for 178 improvement with respect to marker vaccines and their companion diagnostic tests.
- Inactivated (non-replicating) whole virus vaccines are not presently available and may be difficult to develop to meet
 minimum efficacy standards. Recombinant vectored, subunit vaccine candidates that can be produced in scalable vaccine
 platform expression systems and mRNA-based ASF vaccines are being evaluated in ongoing laboratory research, testing
- 182 and evaluation in experimental challenge models. The publicly available Center of Excellence for African Swine Fever

183 <u>Genomics (ASFV Genomics, 2022⁴³) that provides the structural protein predictions for all 193 ASFV proteins may help
 184 accelerate ASF first and second generation vaccine research and development.
</u>

185 <u>Fit-for-purpose vaccine use scenarios matched to the intended use in a domestic pig specific type of production system</u>
 186 <u>may require different vaccine product profiles or may influence the focus of essential versus ideal vaccine requirements.</u>
 187 <u>As with any MLV vaccine, all ASF MLV vaccines should be used according to the label instructions, under the strict control</u>
 188 <u>of the country's Regulatory Authority.</u>

 The minimum standards given here and in chapter 1.1.8 are intended to be general in nature and may be supplemented by national, regional, and veterinary international medicinal product harmonised requirements. Minimum data requirements for an authorisation in exceptional circumstances should be considered where applicable.

192 **2.** Outline of production and minimum requirements for vaccines

193 **2.1. Characteristics of the seed**

194

2.1.1. Biological characteristics of the master seed

195		MLVs are produced from ASFV field strains derived from naturally attenuated field isolates or using
196		DNA homologous (genetically targeted) recombination techniques in cell cultures to delete one or
197		more ASFV genes or gene families. These molecular techniques typically involve replacement of the
198		targeted ASFV gene(s) with one or more positive, marker fluorescent (e.g. BFP, eGFP, mCherry) or
199		enzyme-based (e.g. β-glucuronidase) ASFV promoter-reporter gene systems that allow the use of
200		imaging microscopy or flow cytometry to visualise, select, and clone gene-deleted, recombinant, ASF
201		MLVs. MLV production is carried out in cell cultures based on a seed-lot system.
000		Master and viruses (MSVa) for MIVa should be calested and produced based on their access of
202		Master seed viruses (MSVs) for MLVs should be selected and produced based on their ease of
203		growth in cell culture, virus yield (log ₁₀ infectious titre) and genetic stability over multiple cell
204		passages. Preferably, a continuous well-characterised cell line (e.g. ZMAC-4; PIPEC; IPKM) (Borca
205		et al., 2021; Masujin et al., 2021; Portugal et al., 2020) is used to produce a master cell bank (MCB)
206		on which the MSV and MSV-derived working seed virus (WSV) can be produced. The exact source
207		of the underlying ASFV isolate, the whole genome sequence, and the passage history must be
208		recorded.
209	<u>2.1.2.</u>	Quality criteria (sterility, purity, freedom from extraneous agents)
210		Only MSVs that have been established as sterile, pure (free of wild-type parental virus and free of
211		extraneous agents as described in Chapter 1.1.9 Tests for sterility and freedom from contamination
212		of biological materials intended for veterinary use, and those listed by the appropriate licensing
213		authorities) and immunogenic, should be used as the vaccine virus (WSV and vaccine batch
214		production). Live vaccines must be shown not to cause disease or other adverse effects in target
215		animals in accordance with chapter 1.1.8, Section 7.1 Safety tests (for live attenuated MSVs), that
216		includes target animal safety tests, increase in virulence tests, assessing the risk to the environment)
217		and if possible, no transmission to other animals.
218		Identity of the MSV must be confirmed using appropriate methods (e.g. through the use of vaccine
219		strain-specific whole genome detection methods such as next generation sequencing).
210		oran operation of generic detection methode oden de nort generation orquenengy.
220		Demonstration of MSV stability over several cell passages is necessary, typically through at least five
221		passages (e.g. MSV+5). For those MLV vaccines for which attenuation is linked to specific
222		characteristics (gene deletion, gene mutations, etc.), genetic stability of attenuation throughout the
223		production process should be confirmed using suitable methods. Suitable techniques to demonstrate
224		genetic stability may include but are not limited to: genome sequencing, biochemical, proteomic,
225		genotypic (e.g. detection of genetic markers) and phenotypic strain characterisation. If final product
226		vields (infectious titres) are relatively low, genetic stability at a minimum of MSV+10 should be
227		demonstrated to allow more flexibility in the outline of production. For example, if MSV+8 is the
228		maximum passage for use in final product manufacturing, demonstration of genetic stability to at least
229		MSV+10 is warranted.

43 http://asfvgenomics.com. Accessed 4/4/2023.

230		<u>2.1.3.</u>	Validation as a vaccine strain
231			The vaccine derived from the MSV must be shown to be satisfactory with respect to safety and
232			efficacy.
233			Even if pigs are not known for susceptibility to transmissible spongiform encephalopathy (TSE)
234			agents, consideration should also be given to minimising the risk of TSE transmission by ensuring
235 236			that animal origin materials from TSE-relevant species, if no alternatives exist for vaccine virus propagation, comply with the measures on minimising the risk of transmission of TSE.
237 238			Ideally, the vaccine virus in the final product should generally not differ by more than five passages from the master seed lot.
239			ASF vaccine should be presented in a suitable pharmaceutical form (e.g. lyophilisate or liquid form).
240	<u>2.2.</u>	Metho	d of manufacture
241		<u>2.2.1.</u>	Procedure
242			The MLV virus is used to infect swine primary cell cultures obtained from specific-pathogen free pigs.
243 244			the requirements for which are defined in specific monographs (Chapter 2.3.3 <i>Minimum requirements for the organisation and management of a vaccine manufacturing facility</i> , Section 2.4.2). Compared
244 245			with primary cell cultures, use of a continuous cell line generally allows for more consistency, higher
246			serial volumes in manufacturing and aligns better with a seed lot system. Thus, preferably a master
247			cell bank based established, continuous cell line shown to support genetically stable ASFV replication
248			and acceptable titres over several passages should be used.
249 250			<u>Cell cultures shall comply with the requirements for cell cultures for production of veterinary vaccines</u> in chapter 1.1.8. Regardless of the production method, the substrate should be harvested under
251			aseptic conditions and may be subjected to appropriate methods to release cell-associated virus (e.g.
252			freeze-thaw cycles, detergent lysis). The harvest can be further processed by filtration and other
253			purification methods. A stabiliser or other excipients may be added as appropriate. The vaccine is
254			
			homogenised to ensure a uniform batch/serial.
255		<u>2.2.2.</u>	Requirements for ingredients
255 256			Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8.
		<u>2.2.2.</u> <u>2.2.3.</u>	Requirements for ingredients
256			Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8.
256 257 258			Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk
256 257 258 259		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests.
256 257 258 259 260		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests
256 257 258 259 260 261		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility
256 257 258 259 260 261 262 263 264		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity
256 257 258 259 260 261 262 263		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9.
256 257 258 259 260 261 262 263 264 265		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity Appropriate methods such as specific genome detection methods (e.g. specific differential real-time
256 257 258 259 260 261 262 263 264 265 266 266 267 268		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity Appropriate methods such as specific genome detection methods (e.g. specific differential real-time PCR) should be used for confirmation of the identity of the vaccine virus. iii) <u>Purity</u> Appropriate methods should be used to ensure that the final product batch does not contain any
256 257 258 259 260 261 262 263 264 265 266 265		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity Appropriate methods such as specific genome detection methods (e.g. specific differential real-time PCR) should be used for confirmation of the identity of the vaccine virus. iii) Purity
256 257 258 259 260 261 262 263 264 265 266 265 266 267 268 269 270		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity Appropriate methods such as specific genome detection methods (e.g. specific differential real-time PCR) should be used for confirmation of the identity of the vaccine virus. iii) Purity Appropriate methods should be used to ensure that the final product batch does not contain any residual wild-type ASFV. iv) Safety
256 257 258 259 260 261 262 263 264 265 266 265 266 267 268 269 270 271		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity Appropriate methods such as specific genome detection methods (e.g. specific differential real-time PCR) should be used for confirmation of the identity of the vaccine virus. iii) Purity Appropriate methods should be used to ensure that the final product batch does not contain any residual wild-type ASEV. iv) Safety Batch safety testing is to be carried out unless consistent safety of the product is demonstrated and
256 257 258 259 260 261 262 263 264 265 266 265 266 267 268 269 270		<u>2.2.3.</u>	Requirements for ingredients All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8. In-process controls In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests. Final product batch tests i) Sterility Tests for sterility and freedom from contamination of biological materials intended for veterinary use may be found in chapter 1.1.9. ii) Identity Appropriate methods such as specific genome detection methods (e.g. specific differential real-time PCR) should be used for confirmation of the identity of the vaccine virus. iii) Purity Appropriate methods should be used to ensure that the final product batch does not contain any residual wild-type ASFV. iv) Safety

274	v) <u>Batch/serial potency</u>
275	Virus titration is a reliable indicator of vaccine potency once a relationship has been established
276	between the vaccine minimum immunising dose (MID) (minimum protective dose) and titre of the
277	modified live vaccine in vitro. In the absence of a demonstrated correlation between the virus titre
278	and protection, an efficacy test will be necessary (Section C.2.3.3 Efficacy requirements, below).
279	vi) <u>Residual humidity/residual moisture</u>
280	The test should be carried out consistent with VICH 44 GL26 (Biologicals: Testing of Residual
281	Moisture, 2003 ⁴⁵). Required for MLV vaccines presented as lyophilisates for suspension for injection.
282	2.3. Requirements for authorisation/registration/licensing
283	2.3.1. Manufacturing process
284	For regulatory approval of a vaccine, all relevant details concerning history of the pre-MSV,
285 286	preparation of MSV, manufacture of the vaccine and quality control testing (Sections C.2.1 Characteristics of the seed and C.2.2 Method of manufacture) should be submitted to the authorities.
287	Information shall be provided from three consecutive vaccine batches originating from the same MSV
288	and representative of routine production, with a volume not less than 1/10, and more preferably with
289 290	a volume not less than 1/3 of the typical industrial batch volume. The in-process controls are part of the manufacturing process.
291	2.3.2. Safety requirements
292 293	For the purpose of gaining regulatory approval, the following safety tests should be performed satisfactorily.
294	As a minimum standard, vaccines should be tested for any pathogenic effects on healthy domestic
295 296	pigs of the target age intended for use. Additional demonstration of MLV safety in breeding age gilts and pregnant sows is preferred but not required as a minimum standard.
297	i) <u>Safety in young animals</u>
298	Carry out the test by each recommended route of administration using, in each case, piglets a
299	minimum of 6-weeks old and not older than 10-weeks old.
300	The test is conducted using no fewer than eight healthy piglets, and preferably no fewer than ten
301	healthy piglets.
302	Use vaccine virus at the least attenuated passage level that will be present in a batch of the vaccine.
303	Administer to each piglet a quantity of the vaccine virus equivalent to not less than ten times the
304	maximum virus titre (e.g. 50% haemadsorption dose [HAD ₅₀], 50% tissue culture infective dose
305	[TCID ₅₀], guantitative PCR, etc.) (maximum release dose) likely to be contained in one dose of the
306	vaccine. To obtain individual and group mean baseline temperatures, the body temperature of each
307	vaccinated piglet is measured on at least the 3 consecutive days preceding administration of the
308	vaccine.
309	To confirm the presence or absence of fever accompanied by acute and chronic disease, observe
310	the piglets 4 hours after vaccination and then at least once daily for at least 45 days, preferably 60
311	days post-vaccination. Carry out the daily observations for signs of acute and chronic disease using
312	a quantitative clinical scoring system adding the values for multiple clinical signs (e.g. Gallardo et al.,
313	2015a). These clinical signs should include fever, anorexia, recumbency, skin haemorrhage or
314 315	cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive findings).
316	At a minimum of 45 days post-vaccination, humanely euthanise all vaccinated piglets. Conduct gross
317	pathology on spleen, lung, tonsil, and kidney tissue samples and at least three different lymph nodes

 ⁴⁴ VICH: International Cooperation on Harmonisation of Technical Requirements for Registration of Veterinary Medical Products

 45
 https://www.ema.europa.eu/en/documents/scientific-guideline/vich-gl26-biologicals-testing-residual-moisture-step-7_en.pdf

318	(which should include lymph node closest to site of inoculation, gastrohepatic and submandibular
319	nodes).
320	The vaccine complies with the test if:
320	The vaccine completes with the test in.
321	• No piglet shows abnormal (local or systemic) reactions, reaches the pre-determined humane
322	endpoint defined in the clinical scoring system or dies from causes attributable to the vaccine;
323	• The average body temperature increase for all vaccinated piglets (group mean) for the
323	observation period does not exceed 1.5°C above baseline; and no individual piglet shows a
325	temperature rise above baseline greater than 2.5°C for a period exceeding 3 days.
326	 <u>No vaccinated pigs show notable signs of disease by gross pathology</u>
327	ii) <u>Safety test in pregnant sows and test for transplacental transmission</u>
000	
328	There is currently an absence of published information on ASFV pathogenesis in breeding-age gilts
329	and in pregnant sows associated with ASFV transplacental infection and fetus abortion/stillbirth. If a
330	label claim is pursued for use in breeding age gilts and sows, then a safety study in line with VICH
331	<u>GL44 (Guidelines on Target Animal Safety for Veterinary Live and Inactivated Vaccines, Section 2.2.</u>
332	Reproductive Safety Test, 2009 ⁴⁶) should be completed.
333	iii) Horizontal transmission
000	
334	The test is conducted using no fewer than 12 healthy piglets, a minimum of 6-weeks old and not older
335	than 10-weeks old and of the same origin, that do not have antibodies against ASFV, and blood
336	samples are negative on real-time PCR. All piglets are housed together from day 0 and the number
337	of vaccinated animals is the same as the number of naïve, contact animals. Co-mingle equal numbers
338	of vaccinated and naïve, contact piglets in the same pen or room.
339	Use vaccine virus at the least attenuated passage level that will be present between the master seed
340	lot and a batch of the vaccine. Administer by each recommended route of administration to no fewer
341	than six piglets a quantity of the vaccine virus equivalent to not less than the maximum virus titre
342	(maximum release dose) likely to be contained in 1 dose of the vaccine.
343	To obtain individual and group mean baseline temperatures, the body temperature of each naïve,
344	contact piglet is measured on at least the 3 consecutive days preceding co-mingling with vaccinated
345	piglets. The body temperature of each naïve, contact piglet is then measured daily for at least 45
346	<u>days, preferably 60 days.</u>
347	To confirm the presence or absence of fever accompanied by disease, observe the naïve, contact
348	piglets daily for at least 45 days, preferably 60 days. Carry out the daily observations for signs of
349	acute and chronic clinical disease using a quantitative clinical scoring system adding the values for
350	multiple clinical signs (e.g. Gallardo et al., 2015a). These clinical signs should include fever, anorexia,
351	recumbency, skin haemorrhage or cyanosis, joint swelling and necrotic lesions around the joints,
352	respiratory distress and digestive findings.
002	
252	
353	In addition, blood should be taken from the naïve contact piglets at least twice a week for the first 21
354	days post-vaccination and then on a weekly basis. From the blood samples, determine infectious
355	virus titres by quantitative virus isolation (HAD50/ml or TCID50/ml) and using a real-time PCR test. If
356	the vaccine virus is non-haemadsorbing or does not cause cytopathic effects, a real-time PCR test
357	only may be used.
358	Collect blood (serum) samples from the naïve contact pigs at least at day 21 and day 28 days and
359	carry out an appropriate test to detect vaccine virus antibodies. At a minimum of 45 days, humanely
360	euthanise all naïve, contact piglets. Conduct gross pathology on spleen, lung, tonsil, and kidney
361	tissue samples and at least three different lymph nodes. Determine virus titres in all collected samples
362	by quantitative virus isolation (HAD ₅₀ /mg or TCID ₅₀ /mg) and real-time(RT)-PCR (see Section B.1.
363	Identification of the agent). If the vaccine virus is non-haemadsorbing or does not cause cytopathic
364	effects, a real-time PCR test or other appropriate method (e.g. titration using IPT or FAT detection)
365	may be used.

^{46 &}lt;u>https://www.ema.europa.eu/en/documents/scientific-guideline/vich-gl44-target-animal-safety-veterinary-live-inactived-vaccines-step-</u> 7_en.pdf_

366	The vaccine complies with the test if:
367 368	 <u>No vaccinated or naïve contact piglet shows abnormal (local or systemic) reactions, reaches the predetermined humane endpoint defined in the clinical scoring system or dies from causes</u>
369	attributable to the vaccine;
370	• The average body temperature increase for all naïve, contact piglets (group mean) for the
371 372	observation period does not exceed 1.5°C: above baseline; and no individual piglet shows a temperature rise above baseline greater than 2.5°C for a period exceeding 3 days;
373 374	 <u>No naïve, contact piglet shows notable signs of disease by gross pathology and no virus is</u> detected in their blood or tissue samples
375	<u>No naïve contact pigs test positive for antibodies to the vaccine virus.</u>
376	iv) <u>Post-vaccination kinetics of viral replication (MLV blood and tissue dissemination) study</u>
377	Prior to the reversion to virulence study (Section C2.3.2.v. below), a minimum of one study should be
378	performed to determine the post-vaccination kinetics of virus replication in the blood (viremia), tissues
379	and viral shedding.
380	The test consists of the administration of the vaccine virus from the master seed lot to no fewer than
381	eight healthy piglets, and preferably ten healthy piglets, a minimum of 6-weeks old and not older than
382 383	<u>10-weeks old and of the same origin, that do not have antibodies against ASFV, and blood samples</u> are negative on real-time PCR.
384	Administer to each piglet, using the recommended route of administration most likely to result in
385	spread (such as the intramuscular route or intranasal route), a quantity of the master seed vaccine
386	virus equivalent to not less than the maximum virus titre (maximum release dose) likely to be
387	contained in 1 dose of the vaccine.
388 389	Record daily body temperatures and observe inoculated animals daily for clinical disease for at least 45 days, preferably 60 days.
390	Carry out the daily observations for signs of acute and chronic clinical disease using a quantitative
391	clinical scoring system adding the values for multiple clinical signs (e.g. Gallardo et al. (2015a). These
392 393	clinical signs should include fever, anorexia, recumbency, skin haemorrhage or cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive findings.
394	Collect blood samples from all the piglets at least two times per week from 3 days post-vaccination
395	for the first 2 weeks, then weekly for the duration of the test. Determine vaccine virus titres by
396 397	<u>quantitative virus isolation (HAD₅₀/ml or TCID₅₀/ml) and using a real-time PCR test. If the vaccine</u> virus is non-haemadsorbing or does not cause cytopathic effects, a real-time PCR test only may be
398	used.
399	Determine which blood timepoint(s) should be used in the design of the reversion to virulence study
400	(Section C2.3.2.v. below).Collect oral, nasal and faecal swab samples (preferably devoid of blood to
401	minimise assay interference) at least two times per week from 3-days post-vaccination for the first
402	2 weeks, then weekly for the duration of the test. Test the swabs for the presence of vaccine virus. Determine virus titres in all collected samples by quantitative virus isolation (HAD ₅₀ /ml or TCID ₅₀ /ml)
403 404	and using a real-time PCR test. If the vaccine virus is non-haemadsorbing or does not cause
405	cytopathic effects, a real-time PCR test or other appropriate method (e.g. titration using IPT or FAT
406	detection) may be used.
407	Euthanise at least two piglets on days 7, 14, 21, and preferably on day 28 (±2 days at each timepoint)
408	and collect spleen, lung, tonsil, kidney tissue samples and at least three different lymph nodes (which
409 410	<u>should include lymph node closest to site of inoculation, gastrohepatic and submandibular nodes).</u> Determine virus titres in all collected samples by quantitative virus isolation (HAD₅₀/mg or TCID₅₀/mg)
411	and using real-time PCR test. If the vaccine virus is non-haemadsorbing or does not cause cytopathic
412	effects, a real-time PCR test or other appropriate method (e.g. titration using IPT or FAT detection)
413	<u>may be used.</u>
414	Determine which tissue(s) and timepoint(s) should be used to aid in the design of the reversion to
415	virulence study (Section C.2.3.2.v), for example, specific tissues at specific timepoints which show
416	the highest titres should be considered for selection and use in the reversion to virulence study.

417	v) <u>Reversion to virulence</u>
418	The test should be carried out consistent with VICH GL41 (Examination of live veterinary vaccines in
419	target animals for absence of reversion to virulence, 2008 ⁴⁷).
420	The test for increase in virulence consists of the administration of the vaccine master seed virus to
421	healthy piglets of an age (e.g. between 6-weeks and 10-weeks old) suitable for recovery of the strain
422	and of the same origin, that do not have antibodies against ASFV, and blood samples that are
423	negative on real-time PCR. This protocol is typically repeated five times.
424	<u>First pass (p1)</u>
425	Administer to no fewer than two piglets, and preferably no fewer than four piglets using the intended
426	route of administration for the final product, a quantity of the master seed vaccine virus equivalent to
427	not less than the maximum virus titre (maximum release dose) likely to be contained in 1 dose of the
428	vaccine. Observe inoculated animals daily for the appearance of at least two and preferably at least
429	three clinical signs and record daily body temperatures.
430	Based on results from at least one completed vaccine shed and spread (virus blood and tissue
431	dissemination study, Section C.2.3.2.iv above) collect an appropriate quantity of blood from each
432	piglet on the predetermined single timepoint (day 5–13). Determine virus titres in individual blood
433	samples by quantitative virus isolation (HAD ₅₀ /ml or TCID ₅₀ /ml) and by real-time PCR. If the vaccine
434 435	virus is non-haemadsorbing or does not cause cytopathic effects, a real-time PCR test or other appropriate method (e.g. titration using IPT or FAT detection) may be used. Identify the individual
435	blood sample(s) with the highest infectious titre and reserve for the subsequent <i>in-vivo</i> passage
430	(second pass, p2).
437	<u>(Second pass, pz).</u>
438	Based on results from at least one completed vaccine virus blood and tissue distribution
439	dissemination study, Section C.2.3.2.iv above) euthanise piglets on the predetermined timepoint (i.e.
440	day 7, 14, 21, or 28). Determine infectious virus titres in individual tissue samples by quantitative
441	virus isolation (HAD50/ml or TCID50/ml). If the vaccine virus is non-haemadsorbing or does not cause
442	cytopathic effects, a real-time PCR test or other appropriate method (e.g. titration using IPT or FAT
443	detection) may be used. Identify individual tissue sample(s) with the highest infectious titre. Pool the
444	tissues from different organs from all animals with the highest titres and prepare at least a 10%
445	suspension in PBS, pH 7.2 kept at 4°C or at -70°C for longer storage. Test each blood and tissue
446	pool used for inoculation by PCR to confirm the absence of potential viral agent contaminants (i.e.
447 448	<u>CSFV, FMDV, PRRS, PCV2). Blood and pooled tissue (p1) are used to inoculate 2 ml of positive</u> material using the intended route of administration for the final product to each of least two and ideally
448	at least four further pigs of the same age and origin.
449	<u>a treastrour luttrer pigs of the same age and origin.</u>
450	<u>Second pass (p2)</u>
451	If no virus is found (p1), repeat the administration by the intended route once again with the same
452	pooled material (blood and pooled tissue, p1) in another ten healthy piglets of the same age and
453	<u>origin.</u>
454	If no virus is found at this point, end the process here. If, however, virus is found, carry out a second
455	series of passages by administering 2 ml of positive material using the intended route of
456	administration for the final product to each of no fewer than two piglets, and preferably no fewer than
457	four piglets of the same age and origin. Observe inoculated animals daily for the appearance of at
458	least two and preferably at least three clinical signs and record daily body temperatures.
459	<u>Third and fourth pass (p3 and p4)</u>
460	If no virus (p2), repeat the intramuscular administration once again with the same pooled material
461	(blood and pooled tissue, p2) in another eight healthy piglets of the same age and origin.
462	If no virus is found at this point, end the process here. If, however, virus is found, carry out this
463	passage operation no fewer than two additional times (p3 and p4) (to each of no fewer than two
464	piglets, and preferably no fewer than four piglets of the same age and origin) and verifying the
465	presence of the virus at each passage in blood and tissues. Observe inoculated animals daily for the
	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, _,, _

^{47 &}lt;u>https://www.ema.europa.eu/en/documents/scientific-guideline/vich-gl41-target-animal-safety-examination-live-veterinary-vaccines-target-animals-absence-reversion_en.pdf.</u>

466 467	<u>appearance of at least two and preferably at least three clinical signs and record daily body</u> temperatures.
468	<u>Fifth pass (p5)</u>
469	Administer 2 ml of the blood and pooled tissue (4) to each of at least eight healthy piglets of the same
470	age and origin. Observe inoculated animals daily for at least 28 days post-inoculation for the
471	appearance of at least two and preferably at least three clinical signs, and daily body temperature.
472	The vaccine virus complies with the test if:
473 474	 <u>No piglet shows abnormal local or systemic reaction, reaches the pre-determined humane end</u> point defined in the clinical scoring system or dies from causes attributable to the vaccine; and
475	 There is no indication of increasing virulence (as monitored by daily body temperature
476 477	accompanied by clinical sign observations) of the maximally passaged virus compared with the master seed virus.
478	At a minimum, a safe MLV vaccine shall demonstrate ALL the following features (minimal standards):
479	Absence of fever (defined as average body temperature increase for all vaccinated piglets (group)
480 481	mean) for the observation period does not exceed 1.5°C above baseline; and no individual piglet shows a temperature rise above baseline greater than 2.5°C for a period exceeding 3 days);
482	Absence of chronic and acute clinical signs and gross pathology over the entire test period or
483 484	minimal chronic clinical signs (defined as mild swollen joints with a low clinical score that resolve within 1 week).
485	 Minimal (defined as no naïve, contact piglet shows notable signs of disease by clinical signs and
486	gross pathology and no or a low percentage of contact piglets test both real-time PCR positive
487	and seropositive) or no vaccine virus transmission (defined as no naïve, contact piglet shows
488	notable signs of disease by clinical signs and gross pathology and no contact piglets test both
489	real-time PCR positive and seropositive) over the entire test period;
490	<u>Absence of an increase in virulence (genetic and phenotypic stability) (complies with the reversion</u>
491	to virulence test).
492	In addition, the vaccines in their commercial presentation before being authorised for general use
493	should be tested for safety in the field (see chapter 1.1.8 Section 7.2.3). Additional field safety
494	evaluation studies may include but are not limited to: environmental persistence (e.g. determination
495	of virus recovery from bedding or other surfaces), assessment of immunosuppression, and negative
496	impacts on performance.
497 <u>2.3.3.</u>	Efficacy requirements
498	i) <u>Protective dose</u>
499	Vaccine efficacy is estimated in immunised animals directly, by evaluating their resistance to live
500	virus challenge. The test consists of a vaccination/challenge trial in piglets a minimum of 6-weeks old
501	and not more than 10-weeks old, free of antibodies to ASFV, and negative blood samples by real-
502 502	time PCR. The test is conducted using no fewer than 15 and preferably no fewer than 24 vaccinated pigs, and no fewer than five non-vaccinated control piglets.
503	pigs, and no lewer than nive non-vaccinated control piglets.
504	The test is conducted to determine the minimal immunising dose (MID) (also referred to as the
505	minimal protective dose [MPD] or protective fraction); using at least three groups of no fewer than
506	five and preferably not fewer than eight vaccinated piglets per group, and one additional group of no
507	fewer than five non-vaccinated piglets of the same age and origin as controls. Use vaccine containing
508	virus at the highest passage level that will be present in a batch of vaccine.
509	Each group of piglets, except the control group, is immunised with a different vaccine virus content
510	in the same vaccine volume. In at least one vaccinated group, piglets are immunised with a vaccine
511	dose containing not more than the minimum virus titre (minimum release dose) likely to be contained
512	in one dose of the vaccine as stated on the label.
513	Twenty-eight days (±2 days) after the single injection of vaccine (or if using two injections of the
514	vaccine then 28 days [±2 days] following the second injection), challenge all the piglets by the
515	intramuscular route. If previous studies have demonstrated acceptable efficacy using IM challenge,

516	then a different challenge route (e.g. direct contact, oral or oronasal) may be used. Challenged,
517	vaccinated piglets may be housed in one or more separate pens in the same room or in different
518	rooms. Challenged, naïve controls can be housed in one or more rooms that are separate from
519	challenged, vaccinated piglets.
520	Carry out the test using an ASFV representative strain of the epidemiologically relevant field strain(s)
521	where the vaccine is intended for use (e.g. ASFV B646L [p72] genotype II pandemic strain and other
522	
	p72 virulent genotype of recognised epidemiologic importance). For gene deleted, recombinant MLV
523	viruses, if neither challenge virus type is available, then carry out the test with the parental, virulent
524	virus used to generate the MLV recombinant virus. Use a 10e3–10e4 HAD ₅₀ (or TCID ₅₀ for non-HAD
525	viruses) challenge dose sufficient to cause death or meet the humane endpoint in 100% of the
526	nonvaccinated piglets in less than 21 days. Higher or lower challenge doses can be considered if
527	appropriately justified.
528	The rectal temperature of each vaccinated piglet is measured on at least the 3 days preceding
529	<u>administration of the challenge virus, at the time of challenge, 4 hours after challenge, and then daily</u>
530	for at least 28 days, preferably 35 days. Observe the piglets at least daily for at least 28 days,
531	preferably 35 days. Carry out the daily observations for signs of acute and chronic clinical disease
532	using a quantitative clinical scoring system adding the values for multiple clinical signs (e.g. Gallardo
533	et al., 2015). These clinical signs should include fever, anorexia, recumbency, skin haemorrhage or
534	cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive
535	findings.
536	Collect blood samples from the vaccinated challenged piglets at least two times per week from 3 days
537	post-challenge for at least 28, preferably 35 days. From the blood samples, determine infectious virus
538	titres by quantitative virus isolation (HAD50/ml or TCID50/ml) and using a real-time PCR test. If the
539	vaccine virus is non-haemadsorbing or does not cause cytopathic effects, a real-time PCR test only
540	may be used.
340	<u>may be useu.</u>
541	At the end of the test period, humanely euthanise all vaccinated challenged piglets. Conduct gross
542	pathology on spleen, lung, tonsil, and kidney tissue samples and at least three different lymph nodes.
543	(which should include lymph node closest to site of inoculation, gastrohepatic and submandibular
544	nodes). Determine virus titres in all collected samples by quantitative virus isolation (HAD ₅₀ /mg or
545	TCID ₅₀ /mg) and real-time PCR (see Section B.1. Identification of the agent). If the vaccine virus is
546	non-haemadsorbing or does not cause cytopathic effects, a real-time PCR test or other appropriate
547	method (e.g. titration using IPT or FAT detection) may be used.
548	The test is invalid if fewer than 100% of control piglets die or reach a humane endpoint.
549	The vaccine (or a specific vaccine virus dose if conducting a vaccine dose titration study) complies
550	with the test if:
551	• No vaccinated challenged piglet shows abnormal (local or systemic) reactions, reaches the
552	humane endpoint or dies from causes attributable to ASF;
553	• The average body temperature increase for all vaccinated challenged piglets (group mean) for
555	the observation period does not exceed 2.0°C above baseline; and no individual piglet shows a
555	temperature rise above baseline greater than 2.0°C;
556	• The vaccinated challenged piglets display a reduction or absence of typical acute clinical signs of
557	disease and gross pathology and a reduction or absence of challenge virus levels in blood and
558	tissues.
559	ii) Assessment for horizontal transmission (challenge virus shed and spread study)
560	The ASF basic reproduction number, R0, can be defined as the average number of secondary ASF
561	disease cases caused by a single ASFV infectious pig during its entire infectious period in a fully
562	susceptible population (Hayes et al., 2021). In general, if the ASFV effective reproduction number
563	<u>Re=R0 × (S/N) (S= susceptible pigs; N= total number of pigs in a given population) is greater than</u>
564	1.0, disease is predicted to spread. Ideally, ASF vaccination should reduce Re to less than 1.0 by
565	reducing the number of susceptible, naïve, contact pigs exposed to vaccinated, infected pigs.
EGG	To evaluate ASF vaccine impact on ASF disease transmission, the test consists of a
566	
567	vaccination/challenge trial in piglets a minimum of 6-weeks old and not older than 10-weeks old, free
568	of antibodies to ASFV, and negative blood samples by real-time PCR.

569	The test is conducted using no fewer than 15 healthy piglets at a ratio comprising twice the number
570	of vaccinated piglets to naïve piglets (e.g. ten vaccinated and five naïve). Use vaccine containing
571	virus at the highest passage level that will be present in a batch of the vaccine.
572	The quantity of vaccine virus administered to each pig is equivalent to be not more than the minimum
573	virus titre (minimum dose) likely to be contained in one dose of the vaccine as stated on the label.
574	Following immunisation, vaccinated and naïve piglets should continue to be co-mingled.
575	Twenty-eight days [±2 days] after the single injection of vaccine (or if using two injections of the
576	vaccine then 28 days [± 2 days] following the second injection), temporarily separate [into different
577	pen(s) or room(s)] all vaccinated piglets from naïve piglets. Challenge all vaccinated piglets by the
578	intramuscular or other previously verified route. Carry out the challenge using an ASFV
579 580	representative strain of the epidemiologically relevant field strain(s) where the vaccine is intended for
580 581	use (e.g. ASFV B646L [p72] genotype II pandemic strain and other p72 virulent genotype of recognised epidemiological importance). For gene deleted, recombinant MLV viruses, if neither
582	challenge virus type is available, then carry out the test with the parental, virulent virus used to
583	generate the MLV recombinant virus. Use a 10e3–10e4 HAD ₅₀ (or TCID ₅₀ for non-HAD viruses
584	challenge dose sufficient to cause death or met the humane endpoint in 100% of the nonvaccinated
585	piglets in less than 21 days. Higher or lower challenge doses can be considered if appropriately
586	justified.
587	Approximately 18-24 hours later, re-introduce naïve piglets to vaccinated, challenged piglets and
588	allow for direct nose to nose contact exposure with vaccinated, challenged piglets. Allow for
589	continuous contact exposure by co-mingling both groups through the end of the study. If more than
590	one pen or room is used for co-housing, following reintroduction initially maintain a ratio of 2:1 of
591	challenged, vaccinated piglets to contact exposed, naïve piglets.
592	The rectal temperature of each contact piglet is measured on at least the 3 days preceding
593	administration of the challenge virus to vaccinated pigs, immediately prior to direct contact exposure,
594	4 hours post-contact exposure, and then daily for at least 28, preferably 35 days. Observe all contact
595	exposed piglets at least daily for at least 28 days, and preferably for at least 35 days.
596	Carry out the daily observations in each contact piglet for signs of acute and chronic clinical disease
597	using a quantitative clinical scoring system adding the values for multiple clinical signs (e.g. Gallardo
598	et al., 2015a). These clinical signs should include fever, anorexia, recumbency, skin haemorrhage or
599	cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive
600	findings.
601	In addition, blood should be taken from the naïve contact piglets at least twice a week from 3 days
602	post-contact exposure for the duration of the test period. From the blood samples, determine
603	infectious challenge virus titres by quantitative virus isolation (HAD ₅₀ /ml or TCID ₅₀ /ml) and using a
604	real-time PCR test. If the vaccine virus is non-haemadsorbing or does not cause cytopathic effects,
605	a real-time PCR test only may be used.
606	Collect blood (serum) samples from the naïve contact pigs at least at day 21 and day 28 (±2 days),
607	and at the end of the test period, and carry out an appropriate test to detect vaccine virus antibodies.
608	<u>Collect oral, nasal and faecal swab samples (preferably devoid of blood to minimise assay</u>
609	interference) from all contact-exposed naïve piglets at least two times per week from 3-days post-
610	contact exposure for the first 2 weeks, then weekly for the duration of the test and test swabs for the
611	presence of challenge virus. Determine virus titres in all collected samples by quantitative virus
612	isolation (HAD50/ml or TCID50/ml) and using a real-time PCR test. If the vaccine virus is non-
613	haemadsorbing or does not cause cytopathic effects, a real-time PCR test or other appropriate
614	method (e.g. titration using IPT or FAT detection) may be used.
615	At the end of the test period, humanely euthanise all contact piglets. Conduct gross pathology on
616	spleen, lung, tonsil, and kidney tissue samples and at least three different lymph nodes. (which should
617	include lymph node closest to site of inoculation, gastrohepatic and submandibular nodes).
618	Determine virus titres in all collected samples by quantitative virus isolation (HAD ₅₀ /mg or TCID ₅₀ /mg)
619	and real-time PCR (see Section B.1. Identification of the agent). If the vaccine virus is non-
620 621	haemadsorbing or does not cause cytopathic effects, a real-time PCR test or other appropriate method (e.g. titration using IPT or FAT detection) may be used.
021	meniou (e.g. unanon using in i or fai uerecnon) may be useu.

622 623		<u>The test is invalid if the vaccine fails to comply with the compliance criteria described for the protected</u> dose test in vaccinated pigs (Section C.2.3.3.i above).
624		The vaccine complies with the test for a reduction in horizontal disease transmission if:
625 626		 <u>No naïve, contact exposed piglet shows abnormal (local or systemic) reactions, reaches the</u> defined humane endpoint or dies from causes attributable to ASF;
627 628		 <u>No naïve, contact exposed piglet displays fever accompanied by typical signs of disease,</u> <u>including gross pathology.</u>
629		Naïve contact pigs show a reduction or absence of challenge virus levels in blood and tissues.
630 631		 None of or a reduced number of naïve contact exposed pigs test positive for antibodies to the challenge virus.
632 633		<u>At a minimum, an efficacious MLV vaccine shall demonstrate ALL the following features (minimal standards):</u>
634		<u>Protects against mortality;</u>
635 636		• <u>Reduces acute disease (fever accompanied by a reduction of typical clinical and pathological signs of acute disease)</u>
637 638 639		 <u>Reduces horizontal disease transmission (no naïve, contact exposed piglet shows abnormal [local or systemic] reactions, reaches the humane endpoint or dies from causes attributable to ASF, and displays fever accompanied by typical acute disease signs caused by ASF)</u>
640		<u>Reduces levels of viral shedding and viraemia.</u>
641 642 643 644		In addition, the vaccines in their commercial presentation before being authorised for general use should be tested for efficacy in the field (see chapter 1.1.8 Section 7.2.3). Additional field efficacy evaluation studies may include but are not limited to: onset of immunity, duration of immunity, and impact on disease transmission.
645	2.3.4.	Duration of immunity
646 647 648		Although not included in the guidance for ASF MLV first generation vaccines, manufacturers are encouraged as part of the authorisation procedure, to demonstrate the duration of immunity of a given vaccine by evaluation of potency at the end of the claimed period of protection.
649	2.3.5.	Stability
650 651 652 653 654		Stability of the vaccine should be demonstrated over the shelf life recommended for the product. Although not included in the standards for first generation MLV ASF vaccines, manufacturers are encouraged, as part of the authorisation procedure, to generate data supporting the period of validity of a lyophilised or other pharmaceutical form of the ASF vaccine as part of the authorisation procedure.
655		REFERENCES
656	Achenbach J.E., Ga	llardo C., Nieto-Pelegrín E., Rivera-Arroyo B., Degefa-Negi T., Arias M., Jenberie S., Mulisa D.D.,

ACHENBACH J.E., GALLARDO C., NIETO-PELEGRÍN E., RIVERA-ARROYO B., DEGEFA-NEGI T., ARIAS M., JENBERIE S., MULISA D.D.,
GIZAW D., GELAYE E., CHIBSSA T.R., BELAYE A., LOITSCH A., FORSA M., YAMI M., DIALLO A., SOLER A., LAMIEN C.E. & SÁNCHEZVIZCAÍNO J.M. (2017). Identification of a New Genotype of African Swine Fever Virus in Domestic Pigs from Ethiopia. *Transbound. Emerg. Dis.*, **64**, 1393–1404.

AGÜERO M., FERNÁNDEZ J., ROMERO L., SANCHEZ C., ARIAS M. & SÁNCHEZ-VIZCAÍNO J.M. (2003). Highly sensitive PCR assay for the routine diagnosis of African swine fever virus in clinical samples, *J. Clin. Microbiol.*, **41**, 4431–4434.

AGÜERO M., FERNÁNDEZ J., ROMERO L., ZAMORA M.J., SANCHEZ C., BELÁK S., ARIAS M. & SÁNCHEZ-VIZCAÍNO J.M. (2004). A
 highly sensitive and specific gel-based multiplex RT-PCR assay for the simultaneous and differential diagnosis of African
 swine fever and Classical swine fever. *Vet. Res.*, **35**, 1–13.

ALEJO A., MATAMOROS T., GUERRA M. & ANDRÉS G. (2018). A Proteomic Atlas of the African Swine Fever Virus Particle. J.
 Virol., 92, pii: e01293-18. doi: 10.1128/JVI.01293-18.

- BARASONA J.A., GALLARDO C., CADENAS-FERNÁNDEZ E., JURADO C., RIVERA B., RODRÍGUEZ-BERTOS A., ARIAS M. & SÁNCHEZ VIZCAÍNO J.M. (2019). First oral vaccination of Eurasian wild boar against African swine fever virus genotype II. *Front. Vet.* Sci., 6, 137. https://doi: 10.3389/fvets.2019.00137.
- BASTO A.P., PORTUGAL R.S., NIX R.J., CARTAXEIRO C., BOINAS F., DIXON L.K., LEITAO A. & MARTINS C. (2006). Development of
 a nested PCR and its internal control for the detection of African swine fever virus (ASFV) in *Ornithodoros erraticus. Arch. Virol.*, **151**, 819–826
- BISHOP R.P., FLEISCHAUER C., DE VILLIERS E.P., OKOTH E.A., ARIAS M., GALLARDO C. & UPTON C. (2015). Comparative analysis
 of the complete genome sequences of Kenyan African swine fever virus isolates within p72 genotypes IX and X. *Virus Genes*, **50**, 303–309.
- BORCA M.V., RAI A., RAMIREZ-MEDINA E., SILVA E., VELAZQUEZ-SALINAS L., VUONO E., PRUITT S., ESPINOZA N. & GLADUE D.P.
 (2021). A Cell Culture-Adapted Vaccine Virus against the Current African Swine Fever Virus Pandemic Strain. J. Virol., 95,
 e0012321. https://doi: 10.1128/JVI.00123-21.
- 679 BOSHOFF C.I., BASTOS A.D., GERBER L.J. & VOSLOO W. (2007). Genetic characterisation of African swine fever viruses from 680 outbreaks in southern Africa (1973–1999). *Vet. Microbiol.*, **121**, 45–55.
- BOURRY O., HUTET E., LE DIMNA M., LUCAS P., BLANCHARD Y., CHASTAGNER A., PABOEUF F. & LE POTIER M.F. (2022).Oronasal
 or Intramuscular Immunization with a Thermo-Attenuated ASFV Strain Provides Full Clinical Protection against Georgia
 2007/1 Challenge. *Viruses*, 14, 2777. doi: 10.3390/v14122777
- 684 <u>BRAKE D.A. (2022). African Swine Fever Modified Live Vaccine Candidates: Transitioning from Discovery to Product</u> 685 <u>Development through Harmonized Standards and Guidelines. *Viruses*, 14, 2619. doi: 10.3390/v14122619</u>
- 686 <u>CADENAS-FERNÁNDEZ E., SÁNCHEZ-VIZCAÍNO J.M., KOSOWSKA A., RIVERA B., MAYORAL-ALEGRE F., RODRÍGUEZ-BERTOS A., YAO</u>
 687 J., BRAY J., LOKHANDWALA S., MWANGI W. & BARASONA J.A. (2020). Adenovirus-vectored African Swine Fever Virus Antigens
 688 <u>Cocktail Is Not Protective against Virulent Arm07 Isolate in Eurasian Wild Boar. Pathogens</u>, 9, 171. doi:
 689 <u>10.3390/pathogens9030171.</u>
- 690 CHAPMAN D.A., DARBY A.C., DA SILVA M., UPTON C., RADFORD A.D. & DIXON L.K. (2011). Genomic analysis of highly virulent 691 Georgia 2007/1 isolate of African swine fever virus. *Emerg. Infect. Dis*, **17**, 599–605.
- 692 <u>CHEN W., ZHAO D., HE X., LIU R., WANG Z., ZHANG X., LI F., SHAN D., CHEN H., ZHANG J., WANG L., WEN Z., WANG X., GUAN Y.,</u>
 693 <u>LIU J. & BU Z. (2020). A seven-gene-deleted African swine fever virus is safe and effective as a live attenuated vaccine in</u>
 694 <u>pigs. Sci. China Life Sci., 63</u>, 623. https://doi: 10.1007/s11427-020-1657-9.
- COSTARD S., MUR L., LUBROTH J., SANCHEZ-VIZCAINO J.M. & PFEIFFER D.U. (2013). Epidemiology of African swine fever virus.
 Virus Res., 173, 191–197.
- DE LEÓN P., BUSTOS M.J. & CARRASCOSA A.L. (2013). Laboratory methods to study African swine fever virus. *Virus Res.*,
 173, 168–179.
- 699 DE VILLIER E.P., GALLARDO C., ARIAS M., DA SILVA M., UPTON C., MARTIN R. & BISHOP R.P. (2010). Phylogenomic analysis of 700 11 complete African swine fever virus genome sequences. *Virology*, **400**, 128–136.
- DIXON L.K., ESCRIBANO J.M., MARTINS C., ROCK D.L., SALAS M.L. & WILKINSON P.J. (2005). *In:* Virus Taxonomy, VIIIth Report
 of the ICTV, Fauquet C.M., Mayo M.A., Maniloff J, Desselberger U.& Ball L.A., eds. Elsevier/Academic Press, London, UK,
 135–143.
- ESCRIBANO J.M., PASTOR M.J. & SANCHEZ VIZCAINO J.M. (1989). Antibodies to bovine serum albumin in swine sera: implications for false positive reactions in the sero diagnosis of African swine fever. *Am. J. Vet. Res.*, **50**, 1118–1122.
- FERNÁNDEZ-PACHECO P., NIETO R., SIMÓN A., GARCÍA CASTEY T.A., MARTÍN E., ARIAS M. & GALLARDO C. (2016). Comparative
 evaluation of the performance of six ELISA tests for the detection of antibodies against African swine fever virus (ASFV).
 EPIZONE 10th Annual Meeting, Madrid, Spain, 27–29 September 2016, p. https://www.epizone-eu.net/en/Home/Downloads.htm
- 710 FERNÁNDEZ-PINERO J., GALLARDO C., ELIZALDE M., RASMUSSEN T.B., STAHL K., LOEFFEN W., BLOME S., BATTEN C., CROOKE H.,
- LE POTIER M.F., UTTENTHAL Å., LEBLANC N., ALBINA E., KOWALCZYK A., MARKOWSKA-DANIEL I., TIGNON M., DE MIA G.M.,
 GIAMMARIOLI M., ARIAS M. & HOFFMANN B. (2010). EPIZONE ring trial on ASFV real-time PCR. Annual Meeting of National
 African swine fever Laboratories, 18 May 2010, Pulawy, Poland.

FERNÁNDEZ-PINERO J., GALLARDO C., ELIZALDE M., ROBLES A., GÓMEZ C., BISHOP R., HEATH L., COUACY-HYMANN E., FASINA
 F.O., PELAYO V., SOLER A & ARIAS M. (2013). Molecular diagnosis of African Swine Fever by a new real-time PCR using
 universal probe library. *Transbound. Emerg. Dis.*, **60**, 48–58.

GALLARDO C., FERNÁNDEZ-PINERO J., PELAYO V., GAZAEV I., MARKOWSKA-DANIEL I., PRIDOTKAS G., NIETO R., FERNÁNDEZ PACHECO P., BOKHAN S., NEVOLKO O., DROZHZHE Z., PÉREZ C., SOLER A., KOLVASOV D. & ARIAS M. (2014). Genetic variation
 among African swine fever genotype II viruses, eastern and central Europe. *Emerg. Infect. Dis.*, 20, 1544–1547.

GALLARDO C., MWAENGO D.M., MACHARIA J.M., ARIAS M., TARACHA E.A., SOLER A., OKOTH E., MARTÍN E., KASITI J. & BISHOP
 RP. (2009). Enhanced discrimination of African swine fever virus isolates through nucleotide sequencing of the p54, p72, and pB602L (CVR) genes. *Virus Genes*, **38** 85–95.

GALLARDO C., NIETO R., SOLER A., PELAYO V., FERNÁNDEZ-PINERO J., MARKOWSKA-DANIEL I., PRIDOTKAS G., NURMOJA I.,
 GRANTA R., SIMÓN A., PÉREZ C., MARTÍN E., FERNÁNDEZ-PACHECO P. & ARIAS M. (2015b). Assessment of African swine fever
 diagnostic techniques as a response to the epidemic outbreaks in Eastern European Union countries: How to improve
 surveillance and control programs. *J. Clin. Microbiol.*, **53**, 2555–2565.

GALLARDO C., SOLER A., NIETO R., CARRASCOSA A.L., DE MIA G.M., BISHOP R.P., MARTINS C., FASINA F.O., COUACY-HYMMAN
 E., HEATH L., PELAYO V., MARTIN E., SIMON A., MARTIN R., OKURUT A.R., LEKOLOL I., OKOTH E. & ARIAS M. (2013). Comparative
 evaluation of novel African swine fever virus (ASF) antibody detection techniques derived from specific ASF viral genotypes
 with the OIE internationally prescribed serological tests. *Vet. Microbiol.*, **162**, 32–43.

- GALLARDO C., SOLER A., NIETO R., SÁNCHEZ M.A.S., MARTINS E., PELAYO V., CARRASCOSA A., REVILLA Y., SIMON A., BRIONES
 V., SÁNCHEZ-VIZCAÍNO J.M. & ARIAS M. (2015a). Experimental Transmission of African Swine Fever (ASF) Low Virulent
 Isolate NH/P68 by Surviving Pigs. *Transbound. Emerg. Dis.*, 62, 612–622.
- GLADUE D.P. & BORCA M.V. (2022). Recombinant ASF Live Attenuated Virus Strains as Experimental Vaccine Candidates.
 Viruses, 14, 878. http://doi:10.3390/v14050878.

GLADUE D.P., RAMIREZ-MEDINA E., VUONO E., SILVA E., RAI A., PRUITT S., ESPINOZA N., VELAZQUEZ-SALINAS L. & BORCA M.V.
 (2021). Deletion of the A137R Gene from the Pandemic Strain of African Swine Fever Virus Attenuates the Strain and
 Offers Protection against the Virulent Pandemic Virus. J. Virol., 95 (21):e0113921. doi: 10.1128/JVI.01139-21. Epub 2021
 Aug 18. PMID: 34406865; PMCID: PMC8513468.

HAINES F.J., HOFMANN M.A., KING D.P., DREW T.W. & CROOKE H.R. (2013). Development and validation of a multiplex, realtime RT PCR assay for the simultaneous detection of classical and African swine fever viruses. *PLoS One*, **8** (7).

- HAYES B.H., ANDRAUD M., SALAZAR L.G., ROSE N. & VERGNE T. (2021). Mechanistic modelling of African swine fever: A
 systematic review. *Prev. Vet. Med.*, **191**, 105358. doi: 10.1016/j.prevetmed.2021.105358.
- KING K., CHAPMAN D., ARGILAGUET J.M., FISHBOURNE E., HUTET E., CARIOLET R., HUTCHINGS G., OURA C.A., NETHERTON C.,
 MOFFAT K., TAYLOR G., LE POTIER M.-F., DIXON L.K. & TAKAMATSU H.-H. (2011). Protection of European domestic pigs from
 virulent African isolates of African swine fever virus by experimental immunisation. Vaccine, 29, 4593–4600. https://doi:
 10.1016/j.vaccine.2011.04.052
- KING D.P., REID S.M., HUTCHINGS G.H., GRIERSON S.S., WILKINSON P.J., DIXON L.K., BASTOS A.D.S. & DREW T.W. (2003).
 Development of a TaqMan® PCR assay with internal amplification control for the detection of African swine fever virus. *J. Virol. Methods*, **107**, 53–61.
- LIU Y., XIE Z., LI Y., SONG Y., DI D., LIU J., GONG L., CHEN Z., WU J., YE Z., LIU J., YU W., LV L., ZHONG Q., TIAN C., SONG Q.,
 WANG H. & CHEN H. (2023). Evaluation of an I177L gene-based five-gene-deleted African swine fever virus as a live
 attenuated vaccine in pigs. *Emerg. Microbes Infect.*, **12**, 2148560. doi: 10.1080/22221751.2022.2148560. PMID:
 36378022; PMCID: PMC9769145.
- LUBISI B.A., BASTOS A.D., DWARKA R.M. & VOSLOO W. (2005). Molecular epidemiology of African swine fever in East Africa.
 Arch Virol., **150**, 2439–2452.

 MASUJIN K., KITAMURA K., KAMEYAMA K.I., OKADERA K., NISHI T., TAKENOUCHI T., KITANI H. & KOKUHO T. (2021). An immortalized porcine macrophage cell line competent for the isolation of African swine fever virus. *Sci. Rep.*, **11**, 4759.
 doi.org/10.1038/s41598-021-84237-2.

MONTEAGUDO P.L., LACASTA A., LOPEZ L., BOSCH J., COLLADO J., PINA-PEDRERO S., CORREA-FIZ F., ACCENSI F., NAVAS M.J.,
 VIDAL E., BUSTOS M.J., RODRÍGUEZ J.M., GALLEI A., NIKOLIN V., SALAS M.L. & RODRÍGUEZ F. (2017). BA71deltaCD2: A new

- recombinant live attenuated African swine fever virus with cross-protective capabilities. J. Virol., 91, e1058–e1117.
 https://doi: 10.1128/JVI.01058-17.
- NCBI (NATIONAL CENTER FOR BIOTECHNOLOGY INFORMATION) (2020). https://www.ncbi.nlm.nih.gov/nuccore/FR682468.
 Accessed 4/4/2023.
- NIX R.J., GALLARDO C., HUTCHINGS G., BLANCO E. & DIXON L.K. (2006). Molecular epidemiology of African swine fever virus studied by analysis of four variable genome regions. *Arch. Virol.*, **151**, 2475–244.

 O'DONNELL V., HOLINKA L.G., GLADUE .DP., SANFORD B., KRUG P.W., LU X., ARZT J., REESE B., CARRILLO C., RISATTI G.R. & BORCA M.V. (2015). African swine fever virus Georgia isolate harboring deletions of MGF360 and MGF505 genes is attenuated in swine and confers protection against challenge with virulent parental virus. *J. Virol.*, **89**, 6048. https://doi: 10.1128/JVI.00554-15.

- O'DONNELL V., RISATTI G.R., HOLINKA L.G., KRUG P.W., CARLSON J., VELAZQUEZ-SALINAS L., AZZINARO P.A., GLADUE D.P. &
 BORCA M.V. (2016). Simultaneous Deletion of the 9GL and UK Genes from the African Swine Fever Virus Georgia 2007
 Isolate Offers In-creased Safety and Protection against Homologous Challenge. J. Virol., 91, e01760-16. https://doi:
 10.1128/JVI.01760-16.
- PASTOR M.J., ARIAS M. & ESCRIBANO J.M. (1990). Comparison of two antigens for use in an enzyme-linked immunosorbent assay to detect African swine fever antibody. *Am. J. Vet. Res.*, **51**, 1540–1543.
- PASTOR M.J., LAVIADA M.D., SANCHEZ VIZCAINO J.M. & ESCRIBANO J.M. (1989). Detection of African swine fever virus antibodies by immunoblotting assay. *Can. J. Vet. Res.*, **53**, 105–107.
- PENRITH M.L., VAN HEERDEN J., HEATH L., ABWORO E.O. & BASTOS A.D.S. (2022). Review of the Pig-Adapted African Swine
 Fever Viruses in and Outside Africa. *Pathogens*, **11**, 1190. doi: 10.3390/pathogens11101190.
- PÉREZ-NÚÑEZ D., SUNWOO S.-Y., GARCÍA-BELMONTE R., KIM C., VIGARA-ASTILLERO G., RIERA E., KIM D.-M., JEONG J., TARK D.,
 KO Y.-S., YOU Y.-K. & REVILLA Y. (2022). Recombinant African Swine Fever Virus Arm/07/CBM/c2 Lacking CD2v and A238L
 Is Attenuated and Protects Pigs against Virulent Korean Paju Strain. Vaccines, 10, 1992.
- 785 https://doi.org/10.3390/vaccines10121992.
- PORTUGAL R., COELHO J., HÖPER D., LITTLE N.S., SMITHSON C., UPTON C., MARTINS C., LEITÃO A. & KEIL G.M. (2015). Related
 strains of African swine fever virus with different virulence: genome comparison and analysis. *J. Gen. Virol.*, **96** (Pt 2), 408–419.
- PORTUGAL R., GOATLEY L.C., HUSMANN R., ZUCKERMANN F.A. & DIXON L.K. (2020). A porcine macrophage cell line that supports high levels of replication of OURT88/3, an attenuated strain of African swine fever virus. *Emerg. Microbes Infect.*, 9, 1245–1253. doi.org/10.1080/22221751.2020.1772675.
- QUEMBO C.J., JORI F., VOSLOO W. & HEATH L. (2018). Genetic characterization of African swine fever virus isolates from soft
 ticks at the wildlife/domestic interface in Mozambique and identification of a novel genotype. *Transbound. Emerg. Dis.*, 65,
 420–431.
- SÁNCHEZ-VIZCAÍNO J.M. (1987). African swine fever diagnosis. *In:* African Swine Fever, Becker Y., ed. Martinus Nijhoff,
 Boston, USA, 63–71.
- SÁNCHEZ-VIZCAÍNO J.M. & ARIAS M. (2012). African swine fever. *In*: Diseases of Swine, tenth Edition, Straw B., D'Allaire S.,
 Mengeling W., Taylor D., eds. Iowa State University, USA, pp.396–404.
- SÁNCHEZ-VIZCAÍNO J.M., MARTINEZ-LÓPEZ B., MARTINEZ-AVILÉS M., MARTINS C., BOINAS B., VIAL L., MICHAUD V., JORI F., ETTER
 E., ALBINA E. & ROGER F. (2009). Scientific Review on African swine fever. CFP/EFSA/AHAW/2007/02, pp: 1–141.
- 801 SÁNCHEZ-VIZCAÍNO J.M., MUR L., GOMEZ-VILLAMANDOS J.C. & CARRASCO L. (2015). An update on the epidemiology and 802 pathology of African swine fever. *J. Comp. Pathol.*, **152**, 9–21.
- <u>TEKLUE T., WANG T., LUO Y., HU R., SUN Y.& QIU H.J. (2020). Generation and Evaluation of an African Swine Fever Virus</u>
 <u>Mutant with Deletion of the CD2v and UK Genes. Vaccines</u>, 8, 763.https://doi:10.3390/vaccines8040763.

TIGNON M., GALLARDO C., ISCARO C., HUTET E., VAN DER STEDE Y., KOLBASOV D., DE MIA G.M., LE POTIER M.F., BISHOP R.P.,
 ARIAS M. & KOENEN F. (2011). Development and inter-laboratory validation study of an improved new real-time PCR assay
 with internal control for detection and laboratory diagnosis of African swine fever virus. *J. Virol. Methods*, **178**, 161–167.

WANG N., ZHAO D., WANG J., ZHANG Y., WANG M., GAO Y., LI F., WANG J., BU Z., RAO Z. & WANG X. (2019). Architecture of
 African swine fever virus and implications for viral assembly. *Science*, **366**, 640–644. doi: 10.1126/science.aaz1439. Epub
 2019 Oct 17

811 812 813	ZHANG Y., KE J., ZHANG J., YANG J., YUE H., ZHOU X., QI Y., ZHU R., MIAO F. & LI Q. (2021). African Swine Fever Virus Bearing an I226R Gene Deletion Elicits Robust Immunity in Pigs to African Swine Fever. J. Virol., 95 , e0119921. https://doi: 10.1128/JVI.01199-21.
814	*
815	* *
816	NB: There are WOAH Reference Laboratories for African swine fever
817	(please consult the WOAH Web site:
818	https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).
819	Please contact the WOAH Reference Laboratories for any further information on
820	diagnostic tests and reagents for African swine fever
821	NB: First adopted in 1990. Most recent updates adopted in 2021.

Draft Standards for African Swine Fever (ASF) Modified Live Virus (MLV) Vaccines for Domestic and Wild Pigs

I. Background

Under a project funded by a Collaborative Agreement between the WOAH and USDA-ARS, and in collaboration with CRDF Global, a consultant, Dr David Brake of BioQuest Associates, LLC, was engaged to develop guidelines on the development and manufacture of safe and efficacious ASF vaccines.

Draft guidelines for ASF modified live virus (MLV) vaccine standards were developed using the source information from WOAH *Terrestrial Manual*, principles and standards described in applicable and current guidelines for veterinary MLVs (published by EMA CVMP, USDA CVB-PEL, VICH and WOAH), peer-reviewed publications on ASF MLV lead vaccine candidates, through sessions at Global ASF Research Alliance (GARA) meetings, as well as general ASF vaccine and laboratory-specific surveys and one-on-one exchanges with laboratory subject matter experts (SMEs). Four technical workshops were also organised to solicit input from SMEs with follow-up surveys and a workshop organised with key opinion leaders from the regulatory sector.

II. Summary of key discussions areas

	Safety		Efficacy	Qu	ality (purity/potency) and other
1.	Breed and gender	1.	Breed and gender	1.	Master seed virus purity –
2.	Age/weight range	2.	Age/weight range		screening for presence of wild- type virus
3.	Group size and housing	3.	Group size and housing	2.	Gene markers in MLV
4.	4. Route of immunisation		Dose		recombinant vaccine
5.	Dose studies	5.	Challenge route	2	candidates
6.	Clinical observations:	6.	Challenge strain and dose	3.	Estimated stability
	frequency, duration, rectal temp, disease/clinical scoring	7.	Challenge timepoint		
7.	Analytical readouts: viremia shedding	8.	Clinical observations: frequency, duration, rectal temp, survival, clinical scoring		
8.	Short <i>vs</i> long term	9.	Analytical readouts: viremia,		
9.	Post-mortem readouts: pathology, tissue persistence	0.	shedding, challenge virus transmission		
10.	Transmission studies	10.	Protective dose (MID vs PD)		
11.	Reversion to virulence	11.	Duration of immunity		
12.	Recombination	12.	Cross (heterologous) protection		
13.	Pregnant animals	13.	DIVA		
14.	Wild boars	14.	Wild boars		
15.	Definitions – minimum standards for fever, clinical signs, residual virulence, viremia, shedding	15.	Definition – minimum standards for fever, clinical signs, "prevents" vs "reduces"		

III. Summary of Points of Consensus

<u>A. General</u>

- Technical requirements should be in standards; whereas some vaccine parameters are more considered national policy based and should be excluded. Examples to omit from draft standards included: i) strict DIVA requirements, ii) how and where to use MLV vaccines +/- stamping out, iii) GMO statements on MLV vaccines.
- International standards should focus on <u>vaccine development criteria</u> to allow a minimum level of regulatory consistency amongst MLV vaccine candidates; standards can also inform on vaccine discovery (labbased) future studies.
- Generally, standards should not be highly restrictive and sufficiently generic for regulatory authorities to use; however, for some parameters more specific definitions should be used when it all possible (i.e. based on current knowledge/publications) but for other parameters less specific in other cases (i.e. to reflect knowledge gaps).
- 4. Consensus building process through identification of uniform safety and efficacy animal models, then developing key safety and efficacy definitions based on minimum acceptability statements.
- 5. First generation published standards should not contain efficacy requirement associated with cross-/heterologous protection.
- 6. Standards should be based on published data and reflect a sensible/achievable level of safety and efficacy. *"Goldilocks Principle"* – neither too hot nor too cold; just the right amount.
- CSF vaccine standards in *Terrestrial Manual* and EMA monograph used as reference for drafting the ASF MLV vaccine standard guidelines; consider adding statements specific to ASF disease pathogenesis where applicable.
- 8. Include standards for wild boar oral vaccines that should be independent from domestic pig vaccine standards.
- 9. Vaccine purity NGS is problematic due to current lack of standards and sensitivity.

B. Laboratory safety specific – minimum standards

- MLV transmission more important than shedding, thus vaccine safety should include measurement of MLV transmission to naïve pigs, particularly in regions where several wild type or unauthorised MLV vaccine strains/genotypes may be co-circulating; however, little published information is currently available on MLV transmission.
- 2. There is a general lack of correlation between: viremia and residual virulence, viremia and ability to shed, viremia and ability to transmit; thus, viremia may not be a highly informative parameter to evaluate vaccine safety; viremia quantitation may not be important in vaccine safety definition and caution should be exercised in setting viremia threshold cut-off.
- 3. Useful to measure both virus isolation and RT-PCR (blood and swabs), however hard to set safety quantitative thresholds.
- 4. In reversion to virulence, not essential to conduct next generation/deep sequencing on ASF MLV virus full genome obtained after the last *in-vivo* passage, in part due to NGS complexity, data interpretation, absence of SOP standard, etc; however, consideration could be given to limiting sequence analysis to genome regions containing gene deletion(s).

Parameter	Minimum safety standard	Consensus comment		
Breed and gender	Nonprescriptive	Breed intended for use		
Age/weight range	6–10-week old	Practicality (procurement); IACUC and animal welfare requirements		
Group sizes	Prescriptive range	Minimum ("at least") and "preferably"		
Penning/housing	Prescriptive mainly for transmission studies	Flexibility to meet IACUC and animal welfare requirements		
Rectal temperature/fever	Important to measure as standalone parameter Fever needs to be accompanied by 2– 3 specific clinical signs			
Clinical sign observations	Carry out the daily observations for clinical disease using a numerical clinical scoring system (e.g. King <i>et al.</i> , 2011)			
Analytical readouts	Not a safety compliant criterion, however important to measure viremia and nasal and faecal swab shedding infectious titres	Prescriptive timepoints for sampling Nonprescriptive for specific threshold values (HAD ₅₀), RT-PCR Ct		
Pregnant sows and breeding gilts	Not specifically drafted	Very little published data on this subject Optional Consistent with VICH guidelines for pregnant animals		
Horizontal transmission studies	Inclusion in vaccine compliant test	Very little published data on this subject Difficult to fully evaluate in lab setting		
Reversion to virulence	Align with VICH GL41	To recommend prescriptive timepoints and sample types for subsequent passages (e.g. P2–P5), future standards could be based on a MLV lead vaccine candidate summary comparative table for viremia and tissue distribution		
Recombination	Not specifically called out due to technical complexity	Agnostic. Regulatory driven case by case based on risk assessment		
Wild boars		Use domestic pigs for majority of studies until final stage for pivotal safety and efficacy		

Safety-related definitions:

- 1. Absence of fever (defined as average body temperature increase for all vaccinated piglets (group mean) for the observation period does not exceed 1.5°C above baseline; and no individual piglet shows a temperature rise above baseline greater than 2.5°C for a period exceeding 3 days).
- Minimum horizontal transmission (defined as no naive, contact piglet shows notable signs of disease by ASF related clinical signs, gross pathology and a low percentage of contact piglets testing both RT-PCR positive and seropositive).
- 3. Absence of an increase in virulence (genetic and phenotypic stability) (defined as complies with the reversion to virulence test.

C. Laboratory Efficacy specific - minimum standards

1. For efficacy, "prevent mortality" and for all other readouts for "reduction in"

Parameter	Minimum efficacy standard	Consensus comment
Breed and gender	Nonprescriptive	Breed intended for use
Age/weight range	6–10-week old	Practicality (procurement); IACUC and animal welfare requirements
Group sizes	Prescriptive range	Minimum ("at least") and "preferably"
Penning/housing	Only prescriptive for horizontal transmission studies	Flexibility to meet IACUC and animal welfare requirements
Challenge route	Challenge all the piglets by the intramuscular route. If previous studies have demonstrated acceptable efficacy using IM challenge, then a different challenge route (e.g. direct contact, oral or oronasal) may be used	
Challenge dose	Consensus only reached for a relatively broad HAD ₅₀ /TCID ₅₀ range 10e2–10e7	
Challenge strain	ASFV representative strain of the epidemiologically relevant field strain(s) where the vaccine is intended for use	ASFV B646L [p72] genotype II pandemic strain identified as highest importance, as well as other p72 virulent genotype of recognised epidemiological importance
Challenge timepoint	28 days following (last) vaccination	Based on majority of publications
Rectal temperature/fever	Important to measure as standalone parameter Fever needs to be accompanied by 2– 3 specific clinical signs	
Clinical sign observations	Carry out the daily observations for clinical disease using a numerical clinical scoring system (e.g. King <i>et al.</i> , 2011)	
Analytical readouts	Not an efficacy compliant criterion, however important to measure viremia, and nasal and faecal swab shedding infectious titres	Prescriptive timepoints for sampling Nonprescriptive for specific threshold values (HAD ₅₀), RT-PCR Ct as titre meaning uncertain
Challenge virus transmission	Part of efficacy definition	
Protective dose	Minimum effective (protective) dose	PD50 or PD80 was also discussed but generally lacked endorsement
Duration of immunity	Not specifically drafted	Insufficient data to clearly define meaning of, thus consensus not to include (SMEs were agonistic) Regulatory key opinion leaders (KOLs) suggested including but referenced manufacturers decision

Efficacy related definitions:

- 1. Protects against mortality.
- 2. Reduces acute disease (defined as fever accompanied by a reduction of typical acute disease signs caused by ASF).
- 3. Reduces horizontal disease transmission (defined as no naive, contact exposed piglets show abnormal [local or systemic] reactions, reaches the humane endpoint or dies from causes attributable to ASF, or

display fever accompanied by typical acute disease signs caused by an ASFV virulent strain representative of the epidemiologically relevant field strain(s) where the vaccine is intended for use.

IV. Summary of points of dissention

A. General

- Minimum age should be <6 weeks-old as: i) ideal target product profile is to vaccinate piglets as young as possible, and ii) regulatory guidance states that vaccine safety should be conducted in the most susceptible age; younger pigs are generally more susceptible vs older pigs.
- 2. Minimum observation period for vaccine safety 21, 28, 35 or 42 days?
- 3. Prescriptive or nonprescriptive timepoints and target samples (blood, tissues) be used for reversion to virulence study.
- 4. Challenge dose how narrow or prescriptive.

B. Laboratory safety specific - minimum standards

Parameter	Minimum safety standard	Dissention comment
Vaccination route	Route intended for the final product	Vast majority of publications on MLV lead vaccine candidates have used IM route, however these conflicts with regulatory standard which needs to be consistent with Ph. 5.2.6
Rectal temperature/fever	The average body temperature increase for all vaccinated piglets (group mean) for the observation period does not exceed 1.5°C above baseline; and no individual piglet shows a temperature rise above baseline greater than 2.5°C for a period exceeding 3 days	What should the specific cut-off values for individual piglets be? 1.5, 2 or 2.5°C ? For how long (days?)(consecutive days?)
Clinical sign observations	Should numerical threshold be used?	What are the most important 2–3 clinical signs to measure? (e.g. inappetence, behaviour, respiratory [laboured breathing or coughing], or digestive [vomiting, diarrhoea]).
Horizontal transmission studies	Inclusion in vaccine compliant test	What should study length be? Should evaluation of ASFV seroconversion and/or presence of ASFV in tissues in contact piglets be part of the vaccine compliant transmission definition?
Reversion to virulence	Prior to the reversion to virulence study (C.v. below), a minimum of one study should be performed to determine the post-vaccination kinetics of virus replication in the blood (viremia), tissues and viral shedding	Should blood and tissue timepoints with highest titre be selected for subsequent passage?

C. Laboratory efficacy specific - minimum standards

Parameter	Minimum efficacy standard	Dissention comment
Challenge dose	Current recommended standard for challenge dose range (10e3–10e4) needs further discussion	General consensus requires further discussion to try and tighten dose range
Rectal temperature/fever	The average body temperature increase for all vaccinated piglets (group mean) for the observation Period does not exceed 2.0°C above baseline; and no individual piglet shows a temperature rise above baseline greater than 2.0°C;	SMEs did not settle on agreeable cut-off values for individual and group piglets What should the specific cut-off values for individual piglets be? 1.5, 2 or 2.5°C? For how long (days?)(consecutive days?)
Clinical sign observations	Carry out the daily observations for clinical disease using a numerical clinical scoring system (e.g. King <i>et</i> <i>al.</i> , 2011) Maximum threshold for safety not	What are the most important 2–3 clinical signs to measure? (e.g. inappetence, behaviour, respiratory [laboured breathing or coughing], or digestive [vomiting, diarrhoea]). Further SME discussion to agree on most
	specifically defined	important 2–3 clinical signs to measure/score
Challenge virus transmission	Final definition of complaint with vaccine w/r/t ASF seroconversion and presence of ASFV in tissues was not resolved	What should be included? Evaluation of ASFV seroconversion and the presence of ASFV in tissues collected from naive, contact exposed piglets?

V. Concluding remarks and recommendations

At the time of drafting there has been few ASF MLV vaccines approved by any regulatory body, thus there was relatively insufficient information to draft international standards and guidelines that were highly prescriptive. There were two major areas (route of administration for assessing vaccine safety and route of administration for assessing vaccine efficacy) in which SME consensus was inconsistent with current international and/or national regulatory guidelines. *Thus, the draft standard guidelines for Section C of Chapter 3.9.1 for vaccine route of administration were written to be consistent with current regulatory guidelines.*

There was sufficient consensus on final draft Section C for consideration by the WOAH Biological Standards Commission. However, there are four major areas which could benefit from future input:

- 1. ASFV challenge dose (range);
- 2. Cut-off values for fever definition for both vaccine safety and vaccine efficacy;
- 3. Most important 2–3 clinical sign observations to measure and whether or not to use a standard numerical scoring for each clinical sign.
- 4. Use of prescriptive timepoints and sample types for subsequent passages for reversion to virulence study;

It is recommended that:

- 1. A semi-annual review is conducted to identify any new peer-reviewed publications and new technical information on existing and any new ASF MLV candidates
- 2. A comprehensive review of peer-reviewed literature conducted on current ASF MLV licensed vaccines and top 12 candidates to generate vaccine safety and efficacy comparative summary tables of methods and results
- 3. Future distribution and request for feedback for draft Section C of Chapter 3.9.1 *African swine fever* (*infection with African swine fever*).

SUMMARY

Description of the disease: Campylobacter jejuni (C. jejuni) and Campylobacter coli (C. coli) can colonise
 the intestinal tract of most mammals and birds and are the most frequently isolated Campylobacter species
 in humans with gastroenteritis. Although poultry is the main reservoir of Campylobacter, transmission to
 humans is only partly through handling and consumption of poultry meat; other transmission routes are also
 considered to be important. This chapter focuses on C. jejuni and C. coli in primary livestock production with
 regard to food safety.

8

15 Campylobacter jejuni and C. coli do not normally cause clinical disease in adult animals except for sporadic 16 cases of abortion in ruminants-and very rare cases of hepatitis in ostriches. The faecal contamination of 17 meat (especially poultry meat) during processing is considered to be an important source of human food-18 borne disease. In humans, extraintestinal infections, including bacteraemia, can occur and some sequelae 19 of infection, such as polyneuropathies, though rare, can be serious.

20 Identification of the agent: In mammals and birds, detection of intestinal colonisation is based on the 21 isolation of the organism from faeces, rectal swabs or caecal contents, or the use of polymerase chain reaction (PCR). Campylobacter jejuni and C. coli are thermophilic, Gram-negative, highly motile bacteria 22 that, for optimal growth, require microaerobic environment and incubation temperatures of 37–42°C. Agar 23 media containing selective antibiotics are required to isolate these bacteria from faecal/intestinal samples. 24 25 Alternatively, their high motility can be exploited using filtration techniques for isolation. Enrichment 26 techniques to detect intestinal colonisation are not routinely used. Preliminary confirmation of isolates can 27 be made by examining the morphology and motility using a light microscope. The organisms in the log growth phase are short and S-shaped in appearance, while coccoid forms predominate in older cultures. Under 28 phase-contrast microscopy the organisms have a characteristic rapid corkscrew-like motility. Phenotypic 29 30 identification is based on reactions under different growth conditions. Biochemical and molecular tests, 31 including PCR and MALDI-TOF (matrix assisted laser desorption ionisation-time of flight) mass 32 spectrometry can be used to identify Campylobacter strains at species level. PCR assays can also be used for the direct detection of C. jejuni and C. coli. 33

34 **Serological tests:** serological assays are not routinely in use for the detection of colonisation by C. jejuni 35 and C. coli.

36 **Requirements for vaccines:** There are no effective vaccines available for the prevention of enteric 37 Campylobacter infections in birds or mammals.

A. INTRODUCTION

39 1. Disease

38

40 Campylobacter jejuni and C. coli are generally considered commensals of livestock, domestic pet animals and birds. Large High numbers of Campylobacter have been isolated from young livestock with enteritis, including piglets, lambs and calves, 41 42 but the organisms are also found in healthy animals. One specific C. jejuni clone has been associated with abortion in 43 sheep (Tang et al., 2017). Outbreaks of avian hepatitis have been reported, but although C. jejuni is associated with the disease, it is not the causative agent (Jennings et al., 2011). Recently, a new Campylobacter was isolated as the causative 44 agent of spotty liver disease in layers (Crawshaw et al., 2015). Campylobacter jejuni and C. coli are of interest mainly from 45 46 the point of view food safety. Campylobacter is the main cause of human bacterial intestinal disease identified in many industrialised countries (Havelaar et al., 2013; Scallan et al., 2011 CDC, 2022; EFSA, 2021), and C. jejuni and C. coli 47 together account for more than 90% of all human campylobacteriosis cases. Over 80% of cases are caused by C. jejuni 48 and about 10% of cases are caused by C. coli. In humans, C. jejuni/C. coli infection is associated with acute enteritis and 49 50 abdominal pain lasting for 7 days or more. Although such infections are generally self-limiting, complications can arise and may include bacteraemia, Guillain-Barré syndrome, reactive arthritis, and abortion (WHO, 2013). Attribution-Studies have 51 shown that the majority of campylobacteriosis cases in humans can be attributed to poultry and a smaller fraction to cattle 52 (Mughini-Gras et al., 2012) is the main reservoir of Campylobacter and responsible for between 50 and 80% of the human 53 54 infections. In the European Union (EU), an estimated 30-20-40% of the human infections are associated with handling and 55 consumption of poultry meat while up to 80% of the strains infecting humans have their origin in the poultry reservoir (EFSA, 2010). ; but A considerable proportion of the poultry-derived strains has a non-poultry meat transmission route. 56 e.g. via environmental contamination surface water (EFSA, 2010b, Mulder et al., 2020). Contact with pets and livestock, 57 the consumption of contaminated water or raw milk and travelling in high prevalence areas are also considered risks factors 58 in human disease (Domingues et al., 2012; Mughini-Gras et al., 2021). The control of Campylobacter in the food chain has 59 now become a major target of agencies responsible for food safety world-wide. 60

61 Laboratory manipulations should be performed with appropriate biosafety and containment procedures as determined by 62 biorisk analysis (see Chapter 1.1.4 *Biosafety and biosecurity: standard for managing biological risk in the veterinary* 63 *diagnostic laboratory and animal facilities*).

64 2. Taxonomy

There are currently 34-43 Campylobacter species recognised (July 2023), but with the improved diagnostic techniques and 65 genomic analysis, this number is expected to increase over time (ef-List of prokaryotic names with standing in 66 nomenclature: (https://lpsn.dsmz.de/genus/campylobacter_http://www.bacterio.net/index.html). Members of the genus 67 68 Campylobacter are typically Gram-negative, non-spore-forming, S-shaped or spiral shaped bacteria (0.2-0.8-0.5 µm wide 69 and 0.5-5-8 um long), with single polar flagella at one or both ends, conferring a characteristic corkscrew-like motility. These bacteria Campylobacter requires microaerobic conditions, but some strains also grow aerobically or anaerobically. 70 They neither ferment nor oxidise carbohydrates. Some species, particularly C. jejuni, C. coli and C. lari, are thermophilic, 71 72 growing optimally at 42°C. They can colonise mucosal surfaces, usually the intestinal tract, of most mammalian and avian species tested. The species C. jejuni includes two subspecies (C. jejuni subsp. jejuni and C. jejuni subsp. doylei) that can 73 be discriminated on the basis of several phenotypic tests, but this subspeciation has no added value for epidemiological 74 75 or intervention purposes (nitrate reduction, selenite reduction, sodium fluoride, and safranine) and growth at 42°C (subsp. doylei does not grow at 42°C) (Garrity, 2005). Subspecies jejuni is much more frequently isolated then subspecies doylei. 76

77

B. DIAGNOSTIC TECHNIQUES



Table 1. Test methods available for the diagnosis of Campylobacter jejuni and C. coli and their purpose

			Purp	oose ^(a)		
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	<u>Prevalence</u> of infection – <u>surveillance</u>	Immune status in individual animals or populations post- vaccination
	Agent identification ^(b)					
Isolation	+++	-	≡ ***	+++	+++	_
MALDI-TOF	+++	-	+++	+++	+++	-

			Purp	DOSe ^(a)		
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	<u>Prevalence</u> of infection – surveillance	Immune status in individual animals or populations post- vaccination
Antigen detection	++	_	<u>=</u> ++	_	++ <u>+</u>	_
16S rRNA sequencing	++	_	++	++	++	_
Real-time PCR	++ <u>+</u>	_	≡ * *	++	++ <u>+</u>	_
	Detection	of immune respo	nse: n/a for <i>Ca</i>	mpylobacter j	ejuni and C. d	coli

82

83

84

85

86

87

96

97

112

80 81 Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; – = not appropriate for this purpose.

MALDI-TOF = matrix assisted laser desorption ionisation-time of flight; PCR = polymerase chain reaction.

^(a)Regarding the control of the agent: Campylobacter jejuni and C. coli are endemic globally and very rarely cause disease. These species are of interest from the point of view of food safety. There is no eradication programme. For broiler flocks there are worldwide efforts to try to prevent colonisation with C. jejuni and C. coli to prevent contamination of the carcasses during slaughter. Therefore, only the columns 'population freedom' (= broiler flock) and prevalence of infection surveillance are filled in where "infection" should be read as "colonisation".

^(b)A combination of agent identification methods applied on the same clinical sample is recommended.

88 **1.** Isolation and identification of the agent

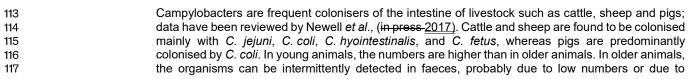
Two ISO (International Organization for Standardization) procedures for detection of *Campylobacter* exist. <u>ISO 10272</u> describes a horizontal method for detection and enumeration of thermotolerant *Campylobacter* spp. (ISO 10272) in food and animal feeding stuffs with 2 parts: (part 1 detection method <u>(ISO 10272-1:2017)</u> and part 2 colony count technique (<u>ISO 10272-2:2017</u>). Both parts of the ISO are under revision and will be published in 2017. The revised standard will include methods for the isolation of *Campylobacter* from live animals, and a procedure for ISO 17995 concerns water quality, with detection and enumeration of thermotolerant *Campylobacter* spp. from water (ISO, 2005 last reviewed in 2014).

1.1. Collection of specimens

1.1.1. Poultry at the farm

98	Poultry is frequently colonised with C. jejuni (65–95%), less often with C. coli and rarely with other
99	Campylobacter species (Newell & Wagenaar, 2000-Wagenaar et al., 2023). Colonisation rates in
100	broiler chickens are age-related. Most flocks are negative until 2 weeks of age. Once Campylobacter
101	colonisation occurs in a broiler flock, transmission, via exposure to faecal contamination, is extremely
102	rapid and up to 100% of birds within a flock can become colonised within a few days. Samples from
103	live birds, destined for the food chain, should therefore be taken as close to slaughter as possible
104	(Newell & Wagenaar, 2000-Wagenaar et al., 2023). The majority of birds shed large numbers of
105	organisms (>10 ⁶ colony-forming units/g faeces). Campylobacters can be isolated from fresh
106	faeces/caecal droppings or cloacal swabs. For reliable detection of Campylobacter by culture, freshly
107	voided faeces (preferably without traces of urine) should be collected. Such samples must be
108	prevented from drying out before culturing. When swabs are used, a transport medium such as
109	Cary Blair, Amies, or Stuart must be used. Sampling strategy in primary poultry has been reviewed
110	(Vidal et al., 2013) and is normally based on boot-swab samples, faecal/caecal droppings or cloacal
111	swabs.

1.1.2. Cattle, sheep and pigs at the farm



intermittent shedding. Fresh samples have to be taken (rectal samples if possible) and **they should be prevented from drying out**. When swabs are used, a transport medium (like Cary Blair, Amies, or Stuart) must be used.

1.1.3. At slaughter

- In poultry, the caeca<u>l contents</u> are usually used for the detection of *Campylobacter*. They-<u>Caeca</u> can be cut with sterile scissors from the remaining part of the intestines and submitted intact to the laboratory in a suitable container.
- Samples from cattle, sheep and pigs are collected from the intestines by aseptically opening the gut wall or by taking guarded rectal swabs.

At all stages from collecting the samples until they are processed in the laboratory, utmost attention should be given to make sure that campylobacters do not die. Follow the instructions for transportation and shipment carefully.

1.2. Transportation and treatment of specimens

1.2.1. Transport

Campylobacters are sensitive to environmental conditions, including dehydration, atmospheric oxygen, sunlight and elevated temperature. Transport to the laboratory and subsequent processing should therefore be as rapid as possible preferably the same day..., but It is recommended to process the samples within 72 hours, but if not possible, storage of samples is accepted up to 96 hours (Tast-Lahti et al., 2022) within at least 3 days. The samples must be protected from light, extreme temperatures and desiccation.

No recommendation on the ideal temperature for transportation can be made, but it is clear that freezing or high temperatures can reduce viability. <u>If possible, samples should be maintained at a temperature of 4°C (±2°C)</u>. High temperatures (>20°C), low temperatures (<0°C) and fluctuations in temperature must be avoided. When the time between sampling and processing is longer than 48 hours, storage at 4°C (±2°C) is advised.

1.2.2. Transport media

Swabs: When samples are collected on boot-swabs or rectal swabs, the use of commercially available transport tubes, containing a medium, such as Cary Blair or Amies, is recommended. This medium may be plain agar or charcoal-based. The function of the medium is not for growth of *Campylobacter* spp., but to protect the swab contents from drying and the toxic effects of oxygen.

When only small amounts of faecal/caecal samples can be collected and transport tubes are not available, shipment of the specimen in transport medium is recommended. Several transport media have been described: <u>Amies</u>, Cary-Blair, modified Cary-Blair, modified Stuart medium, Campy₁ thioglycolate medium, alkaline peptone water and semisolid motility test medium. Good recovery results have been reported using Cary Blair (Luechtefeld *et al.*, 1981; Sjogren *et al.*, 1987).

1.2.3. Maintenance of samples

On arrival at the laboratory, samples should be processed as soon as possible, preferably on the day of arrival. It is recommended to process the samples within 72 hours, but if not possible, storage of samples is accepted up to 96 hours, whereby *C. coli* is more sensitive for long storage times than *C. jejuni* but no longer than 3 days after collecting the samples <u>(Tast-Lahti *et al.*, 2022)</u>. To avoid temperature variation, samples should only be refrigerated when they cannot be processed on the same day, otherwise they should be kept at room temperature <u>when processed the same day</u>. When samples are submitted or kept in the laboratory at 4°C, they should be allowed to equilibrate to room temperature before processing to avoid temperature shock.

1.3. Isolation of *Campylobacter*

For the isolation of *Campylobacter* from faecal/caecal or intestinal samples, no pre-treatment is needed; samples can be plated on selective medium or the filtration method on non-selective agar can be used. In the case of caecal samples, caeca are aseptically opened by cutting the end with a sterile scissors and squeezing out the material to be processed. Enrichment is recommended can be considered to enhance the culture sensitivity of potentially environmentally stressed organisms or in the case of low levels of organisms in faeces <u>(ISO, 2017)</u>, for example from cattle, sheep or pigs. However, enrichment of faecal samples is usually subject to overgrowth by competing bacteria and is not carried out routinely. <u>There is no need to use enrichment</u> media to isolate *Campylobacter* from poultry caeca.

1.3.1. Selective media for isolation

- Many media can be used in the recovery of Campylobacter spp. The selective medium modified charcoal, cefoperazone, desoxycholate agar (mCCDA), is the most commonly recommended medium and is prescribed in the ISO standard, although alternative media may be used (ISO, 2017). A detailed description on Campylobacter detection by culture and the variety of existing media is given by Corry et al. (1995; 2003). The selective media can be divided into two main groups: blood-based media and charcoal-based media. Blood components and charcoal serve to remove toxic oxygen derivatives. Most media are commercially available. The selectivity of the media is determined by the antibiotics used. Cefalosporins (generally cefoperazone) are used, sometimes in combination with other antibiotics (e.g. vancomycin, trimethoprim). Cycloheximide (actidione) and more recently Amphotericin B or cycloheximide are used to inhibit yeasts and molds (Martin et al., 2002). The main difference between the media is the degree of inhibition of contaminating flora. All the selective agents allow the growth of both C. jejuni and C. coli. There is no medium available that allows growth of C. jejuni and inhibits C. coli or vice versa. To some extent, other Campylobacter species (e.g. C. lari, C. upsaliensis, C. helveticus, C. fetus and C. hyointestinalis) will grow on most media, especially at the less selective temperature of 37°C.
 - Examples of selective blood-containing solid media:
 - i) Preston agar

- ii) Skirrow agar
- iií) Butzler agar
 - iv) Campy-cefex

Examples of charcoal-based solid media:

- i) mCCDA (modified charcoal cefoperazone deoxycholate agar), slightly modified version of the originally described CCDA) (Bolton *et al.*, 1984; 1988)
 - ii) Karmali agar or CSM (charcoal-selective medium) (Karmali et al., 1986)
 - iii) CAT agar (cefoperazone, amphotericin and teicoplanin), facilitating growth of *C. upsaliensis* (Aspinall *et al.*, 1993).

<u>1.3.2 Enrichment</u>

The ISO standard describes the isolation of *Campylobacter* from samples with low numbers of *Campylobacter* and high numbers of background flora by using Preston enrichment medium (ISO, 2017). This can be considered for samples from pigs, cattle and sheep. Samples are added to Preston broth with a 1 in 10 dilution (e.g. 10 g faecal sample with 90 ml broth) and incubated under microaerobic conditions for 24 hours at 41.5°C.

After enrichment, campylobacters can be isolated on selective media as described before with plating one loop (10 µl) to solid media.

1.3.3. Passive filtration

Passive filtration, a method developed by Steele & McDermott (1984) obviates the need for selective media; thus it is very useful for the isolation of antimicrobial-sensitive *Campylobacter* species. As the method does not use expensive selective media, it may be used in laboratories with fewer resources. For passive filtration, faeces are mixed with PBS (approximately 1/10 dilution) to produce a suspension. Approximately <u>10–15 drops</u><u>100 µl</u> of this suspension are then carefully layered on to a <u>0.45 or 0.65 µm sterile cellulose acetate</u> filter, which has been previously placed on top of a non-selective blood agar plate. Care must be taken not to allow the inoculum to spill over the edge of the filter. The bacteria are allowed to migrate through the filter for 30–45 minutes at 37°C or room temperature (microaerobic conditions are not required) and the filter is then removed. The plate is incubated microaerobically at <u>37°C or</u> 42°C.

i) Atmosphere 218 219 Microaerobic atmospheres of 5–10% oxygen, 5–10% carbon dioxide are required for optimal growth 220 (Corry et al., 2003; Vandamme, 2000). Appropriate atmospheric conditions may be produced by a variety of methods. In some laboratories, (repeated) gas jar evacuations followed by atmosphere 221 222 replacement with bottled gasses are used. Gas generator kits are available from commercial sources. 223 Variable atmosphere incubators are more suitable if large numbers of cultures are undertaken. 224 ii) Temperature 225 Media may be incubated at 37°C or 42°C, but it is common practice to incubate at 42°C to minimise growth of contaminants and to select for optimal growth of C. jejuni and C. coli. The fungistatic agents 226 eycloheximide or amphothericin B or cycloheximide are added in order to prevent growth of yeasts 227 228 and mould at 37°C (Bolton et al., 1988). In some laboratories, incubation takes place at 41.5°C to harmonise with Salmonella and Escherichia coli O157 isolation protocols (ISO, 2006). 229 230 iii) Time Campylobacter jejuni and C. coli usually show growth on solid media within 24-48 hours at 37-42°C. 231 As the additional number of positive samples obtained by prolonged incubation is very low, 48 hours 232 of incubation is recommended for routine diagnosis (Bolton et al., 1988). 233 Confirmation 1.4. 234 A pure culture is required for confirmatory tests, but a preliminary confirmation can be obtained by direct microscopic 235 236 examination of suspect colony material. 237 1.4.1. Identification on solid medium On Skirrow or other blood-containing agars, characteristic Campylobacter colonies are slightly pink, 238 round, convex, smooth and shiny, with a regular edge. On charcoal-based media such as mCCDA, 239 the characteristic colonies are greyish, flat and moistened, with a tendency to spread, and may have 240 241 a metal sheen. 242 1.4.2. Microscopic examination of morphology and motility 243 Material from a suspect colony is suspended in saline and evaluated, preferably by a phase-contrast 244 microscope, for characteristic, spiral or curved slender rods with a corkscrew-like motility. Older 245 cultures show less motile coccoïd forms. 1.4.3. Detection of oxidase 246 Take material from a suspect colony and place it on to a filter paper moistened with oxidase reagent. 947 248 The appearance of a violet or deep blue colour within 10 seconds is a positive reaction. If a commercially available oxidase test kit is used, follow the manufacturer's instructions. 249 1.4.4. Aerobic growth at 25°C 250 251 Inoculate the pure culture on to a non-selective blood agar plate and incubate at 25°C in an aerobic 252 atmosphere for 48 hours. Latex agglutination tests 253 1.4.5. Latex agglutination tests for confirmation of pure cultures of C. jejuni and C. coli (often also including 254 C. lari) are commercially available. 255 1.5. Biochemical identification of Campylobacter to the species level 256 Among the Campylobacter spp. growing at 42°C, the most frequently encountered species from samples of animal 257 origin are C. jejuni and C. coli. However, low frequencies of other species, including Helicobacter species, have been 258 259

217

1.3.4

Incubation

origin are *C. jejuni* and *C. coli*. However, low frequencies of other species, including *Helicobacter* species, have been
 described. Generally, *C. jejuni* can be differentiated from other *Campylobacter* species on the basis of the hydrolysis
 of hippurate as this is the only hippurate-positive species isolated from veterinary or food samples. The presence of
 hippurate-negative *C. jejuni* strains has been reported (Steinhauserova *et al.*, 2001). Table 2 gives some basic
 classical phenotypic characteristics of the most important thermophilic *Campylobacter* species (ISO, 2006-2017).

- 263 More extensive speciation schemes have been described in the literature (On, 1996; Vandamme, 2000). Speciation 264 results should be confirmed using defined positive and negative controls.
- 265 <u>Biochemical speciation may be supplemented or replaced with MALDI-TOF mass spectrometry. MALDI-TOF can be</u> 266 <u>used to identify *Campylobacter* isolates rapidly and efficiently at the genus and species level (Bessede *et al.*, 2011).</u>

Table 2. Basic phenotypic characteristics of selected thermophilic Campylobacter species

Characteristics	C. jejuni	C. coli	C. lari
Hydrolysis of hippurate	+*	_	-
Hydrolysis of indoxyl acetate	+	+	_

Key: + = positive; - = negative; *not all strains.

The confirmatory tests for the presence of thermophilic campylobacters and the interpretation (ISO, <u>2006-2017</u>) are given in Table 3. Confirm results of confirmation tests using positive and negative controls.

267

Confirmatory test	Result for thermophilic Campylobacter
Morphology	Small curved bacilli
Motility	Characteristic (highly motile and cork-screw like)
Oxidase	+
Aerobic growth at 25°C	-

271 272 273

274 275

276

277 278

279

280 281

282 283

1.5.1. Detection of hippurate hydrolysis

Suspend a loopful of growth from a suspect colony in 400 μ l of a 1% sodium hippurate solution (care should be taken not to incorporate agar). Incubate <u>aerobically</u> at 37°C for 2 hours, then slowly add 200 μ l 3.5% ninhydrin solution to the side of the tube to form an overlay. Re-incubate at 37°C for 40 <u>15–30</u> minutes, and read the reaction. Positive reaction: dark purple/blue. Negative reaction: clear or grey. If commercially available hippurate hydrolysis test disks are used, follow the manufacturer's instructions. <u>The hippurate hydrolysis test is not very robust and the test is often replaced by molecular tests (see below).</u>

1.5.2. Detection of indoxyl acetate hydrolysis

Place a suspect colony on an indoxyl acetate disk and add a drop of sterile distilled water. If indoxyl acetate is hydrolysed a colour change to dark blue occurs within 5–10 minutes. No colour change indicates hydrolysis has not taken place. If commercially available indoxyl acetate hydrolysis test disks are used, follow the manufacturer's instructions.

Biochemical speciation may be supplemented or replaced with molecular methods or MALDI-TOF mass 284 285 spectrometry. MALDI TOF can be used to identify Campylobacter isolates rapidly and efficiently at the genus and species level (Bessede et al., 2011). A variety of DNA probes and polymerase chain reaction (PCR)-based 286 identification assays has been described for the identification of Campylobacter species (On, 1996; Vandamme, 287 288 2000). On & Jordan (2003) evaluated the specificity of 11 PCR-based assays for C. jejuni and C. coli identification. A fast method to differentiate C. jejuni and C. coli strains is a duplex real-time PCR, targeting gene mapA for C. jejuni 289 290 identification and gene CeuE for C. coli identification (Best et al., 2003). Another real-time PCR method commonly used to identify and differentiate between C. jejuni, coli and lari is described by Mayr et al. (2010). A gel-based method 291 that is commonly used differentiates between C. jejuni, C. coli, C. lari and C. upsaliensis (Wang et al., 2002). 292 Campylobacter isolates can also be molecular identified at species level with 16S rRNA sequencing (Gorkiewicz et 293 al., 2003). 294

²⁷⁰

295 **1.6. Molecular detection and identification of Campylobacter**

296 Multiple PCR-based methods for the detection of Campylobacter in animal faecal samples and enriched meat 297 samples have been extensively described in the literature (Bang et al., 2001 Lund et al., 2003; Olsen et al., 1995). Lund et al. describe a real-time PCR method to detect Campylobacter spp. in chicken faecal samples using magnetic 298 bead DNA isolation followed by a real-time PCR targeting the 16S rRNA gene (Lund et al., 2003; 2004). For food 299 samples, a combined method is described of Bolton broth enrichment and multiplex real-time PCR targeting gene 300 301 mapA for C. jejuni, gene ceuE for C. coli and a ATP-binding protein for both C. jejuni and C. coli (Lanzl et al., 2022). 302 Many molecular tests are available to identify Campylobacter species, but there is not a specific recommended one. Campylobacter isolates can be identified at species level with 16S rRNA sequencing (Gorkiewicz et al., 2003). 303 Inclusion of positive and negative reference strains and process controls to detect inhibition of the PCR reaction by 304 305 the sample matrix are required for all molecular Campylobacter detection methods.

306A variety of DNA probes and PCR-based identification assays has been described for the identification of307Campylobacter species (Ferrari et al., 2023; Jribi et al., 2017). On & Jordan (2003) evaluated the specificity of30811 PCR-based assays for C. jejuni and C. coli identification. A fast method to differentiate C. jejuni and C. coli strains309is a duplex real-time PCR, targeting gene mapA for C. jejuni identification and gene ceuE for C. coli identification310(Best et al., 2003). Another real-time PCR method commonly used to identify and differentiate between C. jejuni, C.311coli and C. lari is described by Mayr et al. (2010). Campylobacter isolates can also be identified at species level with31216S rRNA sequencing (Gorkiewicz et al., 2003).

313 **1.7. Antigen-capture-based tests**

Enzyme immunoassays are available for the detection of *Campylobacter* in human and animal stool samples (<u>Ricke</u> *et al.*, 2019). Some are of the lateral flow format. <u>While antigen tests are convenient to use, in an evaluation study</u> where human stool samples were tested with four commercial *Campylobacter* antigen tests, it was shown that no stool antigen test offered the necessary combination of high sensitivity, high specificity, and moderate to high positive predictive value needed in a standalone diagnostic test (Fitzgerald *et al.*, 2016). By using antigen-capture-based tests, the sensitivity and specificity should be critically evaluated through an in-house validation.

320 **2.** Serological tests

321 There are no serological assays in routine use for the detection of colonisation of C. jejuni or /C. coli in livestock.

322 C. REQUIREMENTS FOR VACCINES AND DIAGNOSTIC BIOLOGICALS

323 There are no vaccines specifically developed for *C. jejuni* or *C. coli* in animals or birds.

324

REFERENCES

- ASPINALL S.T., WAREING D.R.A., HAYWARD P.G. & HUTCHINSON D.N. (1993). Selective medium for thermophilic campylobacters including *Campylobacter upsaliensis. J. Clin. Pathol.*, **46**, 829–831.
- BANG D.D., PEDERSEN K. & MADSEN M. (2001). Development of a PCR assay suitable for *Campylobacter* spp. mass
 screening programs in broiler production. *J. Rapid Methods Autom. Microbiol.*, **9**, 97–113.

BESSEDE E., SOLECKI O., SIFRE E., LABADI L. & MEGRAUD F. (2011). Identification of Campylobacter species and related organisms by matrix assisted laser desorption ionization–time of flight (MALDI-TOF) mass spectrometry. *Clin. Microbiol. Infect.*, **17**, 1735–1739.

- BEST E.L., POWEL E.J., SWIFT C., KATHLEEN A.G. & FROST J.A. (2003). Applicability of a rapid duplex real-time PCR assay for speciation of *Campylobacter jejuni* and *Campylobacter coli* directly from culture plates. *FEMS Microbiol.*, **229**, 237–241.
- BOLTON F.J., HUTCHINSON D.N. & COATES D. (1984). Blood-free selective medium for isolation of *Campylobacter jejuni* from faeces. *J. Clin. Microbiol.*, **19**, 169–171.
- BOLTON F.J., HUTCHINSON D.N. & PARKER G. (1988). Reassessment of selective agars and filtration techniques for isolation of *Campylobacter* species from faeces. *Eur. J. Clin. Microbiol. Infect. Dis.*, **7**, 155–160.

- 338
 CDC (CENTRES FOR DISEASE CONTROL AND PROVENTION [OF THE USA]) (2022) FoodNet Homepage, CDC.

 339
 https://www.cdc.gov/foodnet/index.html. Accessed 4 July 2023
- CORRY J.E.L., ATABAY H.I., FORSYTHE S.J. & MANSFIELD L.P. (2003). Culture media for the isolation of campylobacters,
 helicobacter and arcobacters. *In:* Handbook of Culture Media for Food Microbiology, Second Edition, Corry J.E.L., Curtis
 G.D.W. & Baird R.M. eds. Elsevier, Amsterdam, The Netherlands, 271–315.
- CORRY J.E.L., POST D.E., COLIN P. & LAISNEY M.J. (1995). Culture media for the isolation of campylobacters. *Int. J. Food Microbiol.*, **26**, 43–76.
- 345 CRAWSHAW T.R., CHANTER J.I., YOUNG S.C., CAWTHRAW S., WHATMORE A.M., KOYLASS M.S., VIDAL A.B., SALGUERO F.J.-&
 346 IRVINE R.M. (2015). Isolation of a novel thermophilic *Campylobacter* from cases of spotty liver disease in laying hens and
 347 experimental reproduction of infection and microscopic pathology. *Vet. Microbiol.*, **179**, 315–321.
- DOMINGUES A.R., PIRES S.M., HALASA T. & HALD T. (2012). Source attribution of human campylobacteriosis using a metaanalysis of case-control studies of sporadic infections. *Epidemiol. Infect.*, **140**, 970–981.
- EFSA (EUROPEAN FOOD SAFETY AUTHORITY) (2010). Scientific Opinion on Quantification of the risk posed by broiler meat to human campylobacteriosis in the EU. EFSA Panel on Biological Hazards (BIOHAZ). *EFSA J.*, **8**, 1437. [89 pp.].
- 352 <u>EFSA (EUROPEAN FOOD SAFETY AUTHORITY) (2021). The European Union one health 2020 Zoonoses report. EFSA J., 19.</u>
 353 <u>https://doi.org/10.2903/J.EFSA.2021.6971</u>
- FERRARI S., ÁSTVALDSSON A., JERNBERG T., STINGL K., MESSELHÄUßER U. & SKARIN H. (2023). Validation of PCR methods for confirmation and species identification of thermotolerant *Campylobacter* as part of EN ISO 10272 – Microbiology of the food chain – Horizontal method for detection and enumeration of *Campylobacter* spp. *Int. J. Food Microbiol.*, 388, 110064.
 doi: 10.1016/j.ijfoodmicro.2022.110064.
- FITZGERALD C., PATRICK M., GONZALEZ A., AKIN J., POLAGE C.R., WYMORE K., GILLIM-ROSS L., XAVIER K., SADLOWSKI J.,
 MONAHAN J., HURD S., DAHLBERG S., JERRIS R., WATSON R., SANTOVENIA M., MITCHELL D., HARRISON C., TOBIN-D'ANGELO M.,
 DEMARTINO M., PENTELLA M., RAZEQ J., LEONARD C., JUNG C., ACHONG-BOWE R., EVANS Y., JAIN D., JUNI B., LEANO F.,
 ROBINSON T., SMITH K., GITTELMAN R.M., GARRIGAN C. & NACHAMKIN I. (2016). Multicenter Evaluation of Clinical Diagnostic
 Methods for Detection and Isolation of Campylobacter spp. from Stool. J. Clin. Microbiol., 54, 1209–1215. doi:
 10.1128/JCM.01925-15.
- GARRITY G.M. (Editor in Chief) (2005). Bergey's Manual of Systematic Bacteriology, Second Edition. Springer Verlag, New
 York, USA.
- GORKIEWICZ G., FEIERL G., SCHOBER C., DIEBER F., KÖFER J., ZECHNER R. & ZECHNER E.L. (2003). Species-specific identification of Campylobacters by partial 16S rRNA gene sequencing. *J. Clin. Microbiol.*, **41**, 2537–2546.
- 368 HAVELAAR A.H., IVARSSON S., LÖFDAHL M. & NAUTA M.J. (2013). Estimating the true incidence of campylobacteriosis and 369 salmonellosis in the European Union, 2009. *Epidemiol. Infect.*, **141**, 293–302.
- INTERNATIONAL ORGANIZATION FOR STANDARDIZATION (ISO) (2005). ISO 17995:2005. Water quality Detection and
 enumeration of thermophilic *Campylobacter species*. International Organisation for Standardisation (ISO), ISO Central
 Secretariat, 1 rue de Varembé, Case Postale 56, CH 1211, Geneva 20, Switzerland.
- INTERNATIONAL ORGANIZATION FOR STANDARDIZATION (ISO) (<u>2017</u>-2006). ISO 10272-1: <u>2017</u>-2006-AND ISO/TS 10272-2: <u>2017</u>
 Microbiology of food and animal feeding stuffs Horizontal method for the detection and enumeration of
 Campylobacter spp. Part 1: Detection method; Part 2: Colony count technique. International Organisation for
 Standardisation (ISO), ISO Central Secretariat, 1 rue de Varembé, Case Postale 56, CH 1211, Geneva 20, Switzerland.
- JENNINGS J.L., SAIT L.C., PERRETT C.A., FOSTER C., WILLIAMS L.K., HUMPHREY T.J. & COGAN T.A. (2011). Campylobacter jejuni
 is associated with, but not sufficient to cause vibrionic hepatitis in chickens. Vet. Microbiol., 149, 193–199.
- JRIBI H., SELLAMI H., MARIAM S., SMAOUI S., GHORBEL A., HACHICHA S., BENEJAT L., MESSADI-AKROUT F., MÉGRAUD F. & GDOURA
 R. (2017). Isolation and Identification of *Campylobacter* spp. from Poultry and Poultry By-Products in Tunisia by
 Conventional Culture Method and Multiplex Real-Time PCR. J. Food Prot., 80, 1623–1627.
- KARMALI M.A., SIMOR A.E., ROSCOE M., FLEMING P.C., SMITH S.S. & LANE J. (1986). Evalutation of a blood-free, charcoalbased, selective medium for the isolation of *Campylobacter* organisms from feces. *J. Clin. Microbiol.*, **23**, 456–459.

- LANZL M.I., ZWIETERING M.H., ABEE A. & DEN BESTEN H.M.W. (2022). Combining enrichment with multiplex real-time PCR
 leads to faster detection and identification of *Campylobacter* spp. in food compared to ISO 10272–1:2017. *Food Microbiol.*,
 108, 104117.
- LUECHTEFELD N.W., WANG W.L., BLASER M.J. & RELLER L.B. (1981). Evaluation of transport and storage techniques for
 isolation of *Campylobacter fetus* subsp. *jejuni* from turkey cecal specimens. *J. Clin. Microbiol.*, **13**, 438–443.
- <u>LUND M., NORDENTOFT S., PEDERSEN K. & MADSEN M. (2004). Detection of *Campylobacter* spp. in chicken fecal samples by
 real-time PCR. *J. Clin. Microbiol.*, **42**, 5125–5132.
 </u>
- LUND M., WEDDERKOPP A., WAINO M., NORDENTOFT S., BANG D.D., PEDERSEN K. & MADSEN M. (2003). Evaluation of PCR for
 detection of *Campylobacter* in a national broiler surveillance programme in Denmark. *J. Appl. Microbiol.*, **94**, 929–935.
- MARTIN K.W., MATTICK K.L., HARRISON M. & HUMPHREY T.J. (2002). Evaluation of selective media for *Campylobacter* isolation when cycloheximide is replaced with amphotericin B. *Lett. Appl. Microbiol.*, **34**, 124–129.
- MAYR A.M., LICK S., BAUER J., THARIGEN D., BUSCH U. & HUBER I (2010). Rapid detection and differentiation of *Campylobacter jejuni*, *Campylobacter coli* and *Campylobacter lari* in food, using multiplex real-time PCR. *J. Food Prot.*, **73**, 241–250.
- MUGHINI-GRAS L., PIJNACKER R., COIPAN C., MULDER A.C., VELUDO A.F., DE RIJK S., VAN HOEK A.H.A.M., BUIJ R., MUSKENS G.,
 KOENE M., VELDMAN K., DUIM B., VAN DER GRAAF-VAN BLOOIS L., VAN DER WEIJDEN C., KUILING S., VERBRUGGEN A., VAN DER
 GIESSEN J., OPSTEEGH M., VAN DER VOORT M., CASTELIJN G.A.A., SCHETS F.M., BLAAK H., WAGENAAR J.A., ZOMER A.L. & E.
 FRANZ (2021). Sources and transmission routes of campylobacteriosis: A combined analysis of genome and exposure data.
 J. Infect., 82, 216–226. https://doi.org/10.1016/J.JINF.2020.09.039
- MUGHINI-GRAS L., SMID J.H., WAGENAAR J.A., DE BOER A.G., HAVELAAR A.H., FRIESEMA I.H., FRENCH N.P., BUSANI L. & VAN
 PELT W. (2012). Risk factors for campylobacteriosis of chicken, ruminant, and environmental origin: a combined case control and source attribution analysis. *PLoS One*, 7(8):e42599. doi: 10.1371/journal.pone.0042599.
- MULDER A.C., FRANZ E., DE RIJK S., VERSLUIS M.A.J., COIPAN C., BUIJ R., MÜSKENS G., KOENE M., PIJNACKER R., DUIM B., VAN
 DER GRAAF-VAN BLOOIS L., VELDMAN K., WAGENAAR J.A., ZOMER A.L., SCHETS F.M., BLAAK H. & MUGHINI-GRAS L. (2020).
 Tracing the animal sources of surface water contamination with *Campylobacter jejuni* and *Campylobacter coli*. Water Res.,
 187, 116421. doi: 10.1016/j.watres.2020.116421.
- NEWELL D.G., MUGHINI-GRAS L., KALUPAHANA R.S. & WAGENAAR J.A. (2017). *Campylobacter* epidemiology sources and
 routes of transmission for human infection. *Campylobacter:* Features, Detection, and Prevention of Foodborne Disease.
 <u>85-110</u>. Elsevier, Amsterdam, Netherlands.
- NEWELL D.G. & WACENAAR J.A. (2000). Poultry infections and their control at the farm level. *In: Campylobacter,* Second
 Edition, Nachamkin I. & M.J. Blaser, eds. ASM Press, Washington DC, USA, 497–509.
- OLSEN J.E., ABO S., HILL W., NOTERMANS S., WERNARS K., GRANUM P.E., POPVIC T., RASMUSSEN H.N. & OLSVIK O. (1995).
 Probes and polymerase chain reaction for the detection of food-borne bacterial pathogens. *Int. J. Food Microbiol.* 28, 1–
 78.
- 417 ON S.L.W. (1996). Identification methods for Campylobacters, Helicobacters, and Related organisms. *Clin. Microbiol. Rev.*,
 418 9, 405–422.
- 419 ON S.L.W. & JORDAN P.J. (2003). Evaluation of 11 PCR assays for species-level identification of *Campylobacter jejuni* and 420 *Campylobacter coli. J. Clin. Microbiol.*, **41**, 330–336.
- 421 RICKE S.C., FEYE K.M., CHANEY W.E., SHI Z., PAVLIDIS H. & YANG Y. (2019). Developments in Rapid Detection Methods for 422 the Detection of Foodborne *Campylobacter* in the United States. *Front Microbiol.*, **9**, 3280. doi: 10.3389/fmicb.2018.03280.
- Scallan E., Hoekstra R.M., Anculo F.J., Tauxe R.V., Widdowson M.A., Roy S.L., Jones J.L., Griffin P.M. (2011).
 Foodborne illness acquired in the United States major pathogens. *Emerg. Infect. Dis.*, **17**, 7–15.
- 425 SJOCREN E., LINDBLOM G.B. & KAIJSER B. (1987). Comparison of different procedures, transport media, and enrichment 426 media for isolation of *Campylobacter* species from healthy laying hens and humans with diarrhea. *J. Clin. Microbiol.*, **25**,
- 427 1966–1968.

- STEELE T.W. & McDERMOTT S.N. (1984). The use of membrane filters applied directly to the surface of agar plates for the isolation of *Campylobacter jejuni* from feces. *Pathology*, **16**, 263–265.
- STEINHAUSEROVA I., CESKOVA J., FOJTIKOVA K. & OBROVSKA I. (2001). Identification of thermophilic *Campylobacter* spp. by
 phenotypic and molecular methods. *J. Appl. Microbiol.*, **90**, 470–475.
- TANG Y., MEINERSMANN R.J., SAHIN O., WU Z., DAI L., CARLSON J., PLUMBLEE LAWRENCE J., GENZLINGER L., LEJEUNE J.T. &
 ZHANG Q. (2017). Wide but Variable Distribution of a Hypervirulent Campylobacter jejuni Clone in Beef and Dairy Cattle in
 the United States. Appl. Environ. Microbiol., 83. https://doi.org/10.1128/AEM.01425-17
- TAST-LAHTI E., ROSENDAL T., JERNBERG T. & SKARIN H. (2022). Optimal detection of *Campylobacter* spp. in swine caecal
 contents the impact of selective media, time between sampling and start of analysis and the number of colonies
 confirmed. Abstract book FoodMicro2022, 27th International ICFMH (International Committee on Food Microbiology and
 Hygiene) Conference, Athens, Greece.
- 439 VANDAMME P. (2000). Taxonomy of the family Campylobacteraceae. *In: Campylobacter*, Second Edition, Nachamkin I. &
 440 M.J. Blaser, eds. ASM Press, Washington DC, USA, 3–26.
- VIDAL A.B., RODGERS J., ARNOLD M. & CLIFTON-HADLEY F. (2013). Comparison of different sampling strategies and laboratory
 methods for the detection of *C. jejuni* and *C. coli* from broiler flocks at primary production. *Zoonoses Public Health*, **60**,
 412–425.
- WAGENAAR J.A., NEWELL D.G., KALUPAHANA R.S. & MUGHINI-GRAS L. (2023). Campylobacter. Animal Reservoirs, Human
 Infections, and Options for Control. *In:* Zoonoses: Infections Affecting Humans and Animals, Sing A., eds. Springer, Cham.
 https://doi.org/10.1007/978-3-030-85877-3_6-1
- WANG G., CLARK C.G., TAYLOR T.M., PUCKNELL C., BARTON C., PRICE L., WOODWARD D.L. & RODGERS F.G. (2002). Colony
 multiplex PCR assay for identification and differentiation of *Campylobacter jejuni, C. coli, C. lari, C. upsaliensis*, and *C. fetus* subsp. fetus. J. Clin. Microbiol., 40, 4744–4747.
- WORLD HEALTH ORGANIZATION (WHO) (2013). The global view of campylobacteriosis: report of an expert consultation,
 Utrecht, Netherlands, 9–11 July 2012, WHO, Food and Agriculture Organization of the United Nations, World Organisation
 for Animal Health, eds. WHO, Geneva, Switzerland.
- 453 http://www.who.int/iris/handle/10665/80751
- 454 *
 455 *
 456 NB: There is a WOAH Reference Laboratory for campylobacteriosis
 457 (please consult the WOAH Web site for the most up-to-date list:
 458 https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).
 459 Please contact the WOAH Reference Laboratories for any further information on diagnostic tests and reagents for campylobacteriosis
 461 NB: FIRST ADOPTED IN 2004. MOST RECENT UPDATES ADOPTED IN 2017.

5

6

7

1

Annex 18. Item 5.1. - Chapter 3.10.8. Toxoplasmosis

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023

CHAPTER 3.10.8.

TOXOPLASMOSIS

SUMMARY

8 Toxoplasmosis is a zoonotic infection of animals caused by the protozoan parasite Toxoplasma gondii. This parasite has the potential to infect all warm-blooded animals. Although infection does not result in clinical 9 illness in the majority of animal species, in some it causes acute life-threatening disease. In some animals, 10 11 particularly small ruminants, Toxoplasma infection may manifest itself as a disease of pregnancy by multiplying in the placenta and fetus. In these animals it can result in abortion or the birth of weak offspring. 12 Human infections are generally asymptomatic, but they can cause abortion in pregnant women, ocular 13 disease, hydrocephalus or intracranial calcifications in congenitally infected children, ocular toxoplasmosis 14 in immunocompetent individuals, and serious symptoms and even death in severely immunocompromised 15 16 patients.

17Toxoplasma gondii is an obligate intracellular parasite that has a sexual cycle in Felidae and a two-stage18asexual cycle in all warm-blooded animals. Globally the T. gondii population structure is diverse and the19various genotypes are associated with the extent of virulence in particular hosts. In the acute phase of20infection, tachyzoites multiply in host cells to cause varying degrees of tissue destruction. With the onset of21an immune response, tachyzoites transform into bradyzoites that multiply slowly in cells to produce tissue22cysts.

Detection of the agent: In aborted fetuses and placenta, T. gondii is often difficult to find histologically, but is more likely to be seen in tissue sections of brain and placenta. Parasitic stages can be identified by immunohistochemistry, while nucleic acid-based assays might be used to confirm presence of parasite DNA in tissues and may allow genotyping of the parasite in biological specimens. In-vitro isolation of T. gondii from host samples is expensive, time consuming and rarely used.

The sexual part of the life-cycle of T. gondii takes place exclusively in epithelial cells of the feline intestine and can result in the excretion of large numbers of oocysts in the faeces. Oocysts may remain viable in the environment for many months. Oocysts of T. gondii morphologically_resemble those from Hammondia hammondi, a related but non-virulent parasite that also uses cats as definitive hosts. Nucleic acid-based molecular tests are available to distinguish between these related parasites.

Serological tests: Among the easy-to-perform serological tests, the indirect fluorescent antibody test (IFAT) and the direct agglutination test (DAT) allow the titration of sera and the establishment of appropriate cutoffs to ensure diagnostic sensitivity and specificity. The IFAT can be used to differentiate IgM and IgG antibodies. The DAT is fast and requires no complex laboratory facilities. Enzyme-linked immunosorbent assays (ELISA) require more sophisticated laboratory equipment but can process large numbers of samples
 and are easier to standardise.

Requirements for vaccines: A vaccine composed of live T. gondii tachyzoites is available commercially for use in sheep in certain countries. The vaccine is supplied as a concentrated suspension of tachyzoites with an approved diluent and delivery system. The vaccine must be handled strictly according to the manufacturer's instructions as it can be hazardous to the user and has a very short shelf life.

A. INTRODUCTION

Toxoplasma gondii is a zoonotic, obligate intracellular protozoan parasite with the capacity to infect all warm-blooded animals, including birds. Although clinical toxoplasmosis seldom occurs in the majority of animal species, acute lifethreatening disease has been reported in some animals. In small ruminants, in particular, it manifests itself as a disease of pregnancy by multiplying in the placenta and fetus. Acute, potentially fatal, infections have been recorded from a range of wild or zoo animals (Dubey, 2022). Infected humans often show no symptoms, but congenital toxoplasmosis, postnatally acquired ocular toxoplasmosis in immunocompetent individuals, or toxoplasmosis in severely immunocompromised patients represent serious threats (EFSA, 2018).

Toxoplasma gondii has a two-stage asexual cycle in warm-blooded animals and a sexual cycle in Felidae. A systematic review and meta-analysis have reported a global seroprevalence of 38% in domestic cats and 64% wild felids (Hatam-Nahavandi *et al.*, 2021). The genetic diversity of *T. gondii* is complicated; three archetypal clonal lineages (I, II, and III) prevail in Europe and North America; in South America, Asia, and Africa, much greater genetic diversity is apparent and, furthermore, fewer clonal and non-clonal lineages have been genotyped (Lorenzi *et al.*, 2016). Transport between continents via animal migration, including birds, and human activity such as trade may have contributed to the genetic population structures of *T. gondii* in different geographical regions (Shwab *et al.*, 2018).

In the asexual part of the lifecycle, the two developmental stages are the rapidly multiplying tachyzoite and the slowly 58 59 multiplying bradyzoite. In acute infection, tachyzoites actively penetrate host cells where they multiply, causing the cell to 60 rupture and release organisms locally and into the circulation. As the host develops immunity, the parasite retains its overall size and shape, but transforms into the bradyzoite stage and multiplies more slowly within tissue cysts to establish a 61 62 persistent infection. These microscopic tissue cysts occur most frequently in brain and skeletal muscle and represent the 63 quiescent stage of the parasite within the host. Viable tissue cysts within muscle (meat) are a significant source of human infection, and ingestion of bradyzoites in prey is probably the main route of infection to predators, including the felid 64 definitive host. In animals that succumb to acute infection, tachyzoites may be demonstrated in ascitic fluid or in lung 65 impression smears, as well as in tissue sections of the liver and other affected organs. 66

67 Abortions in sheep and goats due to T. gondii are of particular veterinary importance. Toxoplasmosis in small ruminants must be differentiated from diseases caused by other infectious agents, including infections with Chlamydophila abortus 68 (see Chapter 3.8.5 Enzootic abortion of ewes), Coxiella burnetii (see Chapter 3.1.17 Q fever), Brucella melitensis (see 69 70 Chapter 3.1.4 Brucellosis [Brucella abortus, B. melitensis and B. suis]), Campylobacter foetus (see Chapter 3.4.4 Bovine genital campylobacteriosis), Salmonella spp. (see Chapter 3.10.7 Salmonellosis), and the viruses that cause border 71 disease (see Chapter 3.8.1 Border disease), bluetongue (see Chapter 3.1.3), Wesselsbron's disease, and Akabane 72 73 disease (see Chapter 3.10.1). In pigs, Brucella suis (see Chapter 3.1.4) may also cause fetal death, mummification, and 74 abortion.

75 The sexual part of the lifecycle occurs in enteroepithelial cells of the feline definitive host, and results in the production of 76 T. gondii occysts. Following primary infection of a cat, occysts may be shed in the faeces for several days, with large numbers contaminating the environment; up to one billion oocysts from domestic cats, and probably similar numbers from 77 wild felids (Shapiro et al., 2019). The oocysts sporulate in the environment over the next 1-5 days (depending on aeration, 78 79 humidity, and temperature), at which time they become infective. The structure of T. gondii oocysts results in extreme resistance to environmental conditions, with the polymeric structure of the walls giving providing strength against 80 mechanical forces and protection against chemical agents (Shapiro et al., 2019). This results in prolonged survival, up to 81 82 18 months in water at 4°C and, once sporulated, can persist in damp soil for as long, at temperatures ranging from −20°C to 35°C. Sporulated oocysts are 11 × 13 µm in diameter and each contains eight sporozoites, four in each of two sporocysts 83 (Dubey, 2022). When a susceptible animal ingests sporulated oocysts, the sporozoites are released to penetrate the 84 85 intestinal lining, become tachyzoites, and establish an infection.

86 1. Human health risks

43

Toxoplasma gondii is a zoonotic parasite and readily infects people. While human infection, as determined by seropositivity,
 is moderately common globally (local prevalence ranges from under 10% to over 90% [Pappas *et al.*, 2009]), clinical illness
 is relatively uncommon. The immunosuppressed are particularly at risk of developing clinical illness. In patients being

90 treated with immunosuppressive drugs, toxoplasmosis may occur due to new infection or activation of chronic infection. In 91 addition, the parasite can pose a serious threat to an unborn child if the mother becomes infected for the first time while 92 pregnant. The T. gondii genotype is also relevant, and outbreaks of clinical infection with some non-archetypal exotic 93 strains have occurred in people with no apparent immune deficiency. Toxoplasmosis usually manifests as general malaise, 94 fever, and lymphadenopathy. However, more severe symptoms may occur, including ocular problems, such as retinochoroiditis, potentially resulting in loss of vision, pneumonitis, and also toxoplasmic encephalitis. The main burden of 95 human disease, based on disability-adjusted life years (DALYs), ranks as having a high contribution to disease burden 96 97 globally (Torgerson et al., 2015).

As with animal infections, people may be infected by ingestion of bradyzoites in raw or lightly cooked meat containing live 98 T. gondii tissue cysts or by ingestion of sporulated oocysts. These may be as contaminants of water or of raw or lightly 99 cooked fresh produce; less commonly, people can be infected by ingestion of tachyzoites in non-heat-treated milk. In 100 addition, transmission via blood transfusion or organ transplantation is also possible. Outbreaks of both waterborne and 101 foodborne toxoplasmosis have been described (EFSA, 2018; Shapiro et al., 2019). The largest outbreak to date occurred 102 in Santa Maria, Brazil in 2018, and is considered to be due to contamination of the water supply with oocysts of a virulent 103 strain of T. gondii: at least 930 confirmed cases occurred, among which 8% required hospitalisation, with three fetal deaths. 104 10 abortions, and 29 cases of congenital transmission, with 19 infants with ocular lesions (Dubey, 2021). 105

106 Clearly, *T. gondii* represents a human health risk and all laboratory manipulations with live organisms should be handled 107 with appropriate measures determined by risk analysis as described in Chapter 1.1.4 *Biosafety and biosecurity: Standard* 108 *for managing biological risk in the veterinary laboratory and animal facilities.*

109

110

B. DIAGNOSTIC TECHNIQUES

 Table 1. Test methods available for the diagnosis of toxoplasmosis and their purpose

			Р	urpose		
Method	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies	Confirmation of clinical cases	Prevalence of infection – surveillance	Immune status in individual animals or populations post- vaccination
		Detection an	d identificatio	n of the agent		
PCR (including nested and real- time PCR)	_	_	_	++	+	-
LAMP	-	-	_	++	+	_
Histopathology	-	-	-	+	_	-
Immmuno- histochemistry	-	_	_	+	-	-
	Detection of immune response					
IFAT ^(a)	+	+	+	++	++	+
ELISA ^(a)	+	+	+	+++ ^(a)	+++	++
DAT/MAT	+	+	+	+	++	+

116

117

Key: +++ = recommended for this purpose; ++ recommended but has limitations;

+ = suitable in very limited circumstances; - = not appropriate for this purpose.

PCR = polymerase chain reaction; LAMP = loop mediated isothermal amplification; IFAT = indirect fluorescent antibody test; ELISA = enzyme-linked immunosorbent assay; DAT = direct agglutination test; MAT = modified agglutination test.
 (a)In IFAT and ELISA, detection of *Toxoplasma*-specific IgG and IgM antibodies may permit some discrimination between acute and chronic cases of infection. In ELISA, assays assessing the avidity of an IgG response to *T. gondii* may provide information regarding how recently the tested animals have experienced a primary *T. gondii* infection.

118 1. Detection of the agent

119 **1.1. Histopathology**

In animals that die with acute toxoplasmosis, focal mononuclear inflammation, with or without focal necrosis, may be
 seen in a number of tissues, including the liver, heart, and lungs. The latter may be oedematous. Lymph nodes may
 have undergone expansion and there may or may not be focal necrosis with or without haemorrhage. Typically, *T. gondii* tachyzoites may be demonstrable in association with necrosis and inflammation. In fatal cases, tachyzoites
 may be demonstrated in ascitic fluid or in lung impression smears.

In cases of abortion and stillbirth in small ruminants, affected placental cotyledons typically contain large foci of coagulative necrosis that may have become mineralised with time. Any associated inflammation is characteristically slight and nonsuppurative. Well-preserved samples of placental cotyledons may show moderate oedema of the mesenchyme of the foetal villi, with a diffuse hypercellularity due to the presence of large mononuclear cells. Small numbers of intracellular and extracellular stages are sometimes visible, usually on the periphery of a necrotic area or in a villus that is in the early stages of infection. The *T. gondii* tachyzoites appear ovoid, 2–6 μm long, with nuclei that are moderately basophilic and located centrally or towards the posterior end.

In the fetal brain, primary and secondary lesions may develop. Microglial foci, typically with a necrotic and sometimes 132 mineralised centre, and often associated with a mild focal lymphoid meningitis, represent a fetal immune response 133 134 following direct damage by local parasite multiplication. Toxoplasma gondii tissue cysts are only rarely found, usually at the periphery of these lesions. Focal leukomalacia is also common and is considered to be due to fetal anoxia in 135 late gestation caused by advanced lesions in the placentome preventing sufficient oxygen transfer from mother to 136 fetus. Such foci most commonly occur in the cerebral white matter cores, but sometimes also in the cerebellar white 137 138 matter. Focal leukomalacia on its own suggests placental disease or acute insufficiency, but the two types of neuropathological change seen together are characteristic of *T. gondii* infection. 139

140 **1.2. Immunohistochemistry**

141 Confirmation of the identity of *T. gondii*-like structures in tissue sections from such cases, as well as from instances 142 of acute toxoplasmosis, may be achieved by immunohistochemistry that labels intact *T. gondii* or antigenic debris 143 using polyclonal or monoclonal *T. gondii* specific antibodies (Dubey, 2022). The antigen-antibody reaction can be 144 visualized by avidin-biotin-complex (ABC) or indirect immune-peroxidase and the peroxidase–antiperoxidase (PAP) 145 technique. The method is both convenient and sensitive and is used with fixed tissues (including archived tissues) 146 that may also exhibit a degree of decomposition, where isolation would not be appropriate or possible. However, 147 cross-reactions with related parasites like *Neospora caninum* are possible.

148 **1.3. Detection of oocysts**

Toxoplasma gondii oocysts can be detected in stools of felids, as well as contaminating different environmental matrices, such as soil and water, or food, such as molluscs and fresh produce. The low quantity or sparse distribution of oocysts in the contaminated matrix, as for water and fresh produce, means that an initial procedure to concentrate the oocysts from a large volume of sample is needed. Chemical flocculation (e.g. using ferric or aluminium sulphate or calcium carbonate), filtration by cellulose acetate or polycarbonate membranes or cartridge filters and flotation with sucrose or caesium chloride gradient have been widely used for water samples. Washing with appropriate buffer(s) and pelleting by centrifugation is often used for fresh produce (Shapiro *et al.*, 2019; Slana *et al.*, 2021).

Although the autofluorescence of *T. gondii* oocysts, pale blue under UV light, facilitates detection by microscopy, this
 property is shared with oocysts and sporocysts of other related coccidian parasites (e.g. *Hammondia hammondi*)
 (Lindquist *et al.*, 2003). As a commercially available antibody specific for *T. gondii* oocysts for microscopy detection
 is currently lacking, molecular assays are usually used to confirm *T. gondii* identification in field samples. Molecular
 methods are needed to assess or confirm the identity of oocysts observed.

161 **1.4. Molecular methods – detection of nucleic acids**

The presence of *T. gondii* (tachyzoites, tissue cysts, oocysts) can be assessed by detecting the parasite genomic DNA using several molecular techniques, including conventional polymerase chain reaction (PCR), nested PCR and loop-mediated isothermal amplification (LAMP) (Table 1). No standard methods are available and many of the published protocols are not yet sufficiently validated (for details refer to Chapter 1.1.6. *Principles and methods of validation of diagnostic assays for infectious disease* and Chapter 2.2.3. *Development and optimisation of nucleic acid detection assays*). 168 Appropriate protocols may allow detection of T. gondii DNA from circulating tachyzoites (acute infection) or bradyzoites in tissue cysts (latent infection) and in different biological samples, including animal and human 169 tissues (e.g. heart and skeletal muscle, placenta, brain) and body fluids (e.g. blood, urine, aqueous humour, 170 cerebrospinal fluid, amniotic fluid, milk). In addition, DNA from oocysts in stool (only felids), food and 171 environmental samples (fresh produce, water, soil) can be detected (Slana et al., 2021). No standard method 172 173 for extracting T. gondii DNA exists, but suitable DNA extraction protocols, based on both in-house protocols and commercial kits have been developed (Dzib Paredes et al., 2016). Sample preparation and DNA extraction 174 procedures are likely to have a considerable impact on the sensitivity of the test. Sensitivity is generally higher 175 176 in DNA-poor matrices than DNA-rich ones (e.g. tap water vs meat samples). Moreover, inhibitors of DNA amplification differ and are related to sample type. Inhibition of DNA amplification can be avoided by DNA 177 178 extraction optimised for sample type or by using appropriate additives (e.g. bovine serum albumin) during DNA amplification. A specific concentration of the parasite stage (e.g. oocysts) or its DNA (e.g. by magnetic capture) 179 from the matrix might be required prior to DNA extraction as reported for oocysts from water, fresh produce, or 180 181 faeces (Slana et al., 2021). A validated protocol for DNA extraction from meat and meat products using a commercial kit is available at the website of the European Union Reference Laboratory for Parasites (EURLP)⁴⁸. 182 An example of in-house method for DNA extraction from pig tissue (Jauregui et al., 2001) is reported below. 183

1.4.1. DNA extraction from animal tissue

Procedure

184

185

186

187 188

191

192

193

194

195 196

197

198

199 200

201

203

204 205

206

207

208

209

210

211

212

213

214

215

216 217

- i) Homogenate sample (e.g. 50 g of brain or tongue, or 1 g of muscle) in a blender with 5 volumes of sterile saline solution (phosphate-buffered saline [PBS]: 300 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, 1.7 mM NaH₂PO₄).
- 189ii)Digest sample with an equal volume of warm (37°C) pepsin-HCl (1.4 mg of pepsin and 10 mg190of NaCl per ml in 0.1 N HCl) for 1 hour at 37°C in a shaking water bath.
 - iii) Neutralise mixture by two washes with 0.1 M Tris buffer (pH 8.0).
 - iv) Centrifuge mixture aliquots for 10 minutes at 1180 g.
 - v) For each aliquot, digest the post centrifugation pellet overnight at 55°C with DNA digestion buffer (0.5% sodium dodecyl sulphate [SDS], 25 mM ethylene diamine tetra-acetic acid [EDTA], 100 mM NaCl, 20 mM Tris-HCl [pH 8.0], and proteinase K [0.1 mg/ml final concentration]).
 - vi) Extract with one volume of phenol-chloroform-isoamyl alcohol (25:24:1).
 - vii) Precipitate DNA in 0.3 M sodium acetate (final concentration) with 2.5 volumes of 100% ethanol.
 - viii) Resuspend DNA pellets in TE buffer (10 mM Tris-HCl, 1 mM EDTA). Store DNA at –20°C until use.

PCR-based assays are commonly applied for the molecular detection of *T. gondii* genomic DNA (Dzib Paredes *et al.*, 2016; Robert *et al.*, 2021; Slana *et al.*, 2021).

202 1.4.2. DNA extraction from oocysts

Although DNA detection is considered highly specific, cross reactivity has been observed between *T. gondii* and *H. hammondi*, a non-zoonotic coccidian that also uses felids as definitive hosts and cannot be differentiated based on oocyst morphology. A real-time PCR targeting a repetitive element of *H. hammondi* (HhamREP-529) has been demonstrated to be highly sensitive and efficient in distinguishing between the two parasites (Schares *et al.*, 2021).

Detection of DNA from *T. gondii* oocysts may present additional challenges because of inhibitors in faecal matter, vegetable or water sediment, and difficulty of extracting DNA from the oocysts. Options for an efficient breaking of oocysts walls include bead-beating, freeze-thaw cycles, heating or chemical/enzymatic treatments (Slana *et al.*, 2021). An in-house method is detailed below for preparation of oocysts and extraction of DNA. An example of a validated method using a commercial kit and a bead-beating-based DNA extraction is available (Lalle *et al.*, 2018).

Procedure

Wash oocysts four times in 15 ml PBS (300 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, 1.7 mM NaH₂PO₄) in a 15 ml centrifugation tube, with centrifugation between washes (1100 g for 7 minutes without braking).

⁴⁸ https://www.iss.it/documents/5430402/5722370/MI 12 rev. 1.pdf/a82a4078-f511-affe-8f90-cabc397bc8ce?t=1620381672663

218 219	ii)	Incubate pellet (up to 0.5 ml) in 2 ml 5.75 % sodium hypochlorite (sodium hypochlorite, aqueous solution, $\ge 4\%$ as active chlorine) for 30 minutes at 37°C.
220	iii)	Add double-distilled H ₂ O up to 15 ml.
221 222	iv)	Centrifuge supernatant (1100 g for 7 minutes without use of brake) and mix the pellet with PBS. Wash the pellet three times with PBS (1100 <i>g</i> for 7 minutes without brake).
223 224	v)	After final centrifugation, re-suspend pellet in 1 ml PBS, transfer into a 1.5 ml reaction tube and spin down (1100 g for 7 minutes without brake).
225 226	vi)	Remove as much supernatant as possible and apply three freeze–thaw cycles (10 minutes at –20°C followed by 2 minutes at room temperature) to the pellet.
227 228	vii	 Re-suspend pellet in 100 μl OOC lysis buffer (600 mM EDTA, 1.3% [v/v] N-lauroylsarcosine, 2 mg/ml proteinase K, pH 9.5) for 45 minutes, at 65°C.
229 230 231	viii	 Add 400 μl OOC-CTAB buffer (2% [v/v] cetyl-trimethyl ammonium bromide, 1.4 M NaCl, 0.2 % [v/v] mercaptoethanol, 20 mM EDTA, 100 mM tris[hydroxymethyl]aminomethane) for 60 minutes at 60°C.
232 233	ix)	Mix with 500 μl phenol/chloroform/isoamyl alcohol (25/24/1) by inverting 50 times. Centrifuge for 7 minutes at 13,000 <i>g</i> .
234 235	x)	Transfer supernatant to fresh tube and mix with 500 μ l phenol/chloroform/isoamyl alcohol (25/24/1) by inverting 50 times. Centrifuge for 7 minutes at 13,000 g .
236 237	xi)	Transfer supernatant to a fresh tube and add 0.04 volumes of 4 M NaCl and 2–3 volumes of – 20° C cold 96% (v/v) ethanol to precipitate DNA (keep at least 20–30 minutes at – 20° C).
238	xii) Centrifuge for 15 minutes at 13,000 <i>g</i> . Decant the supernatant.
239	xiii) Wash the pellet using 70% (v/v) ethanol and centrifuge for 15 minutes at 13,000 g .
240	xiv	 Discard the ethanol solution and air dry the pellet.
241	xv) Solve DNA in double-distilled water for at least 12 hours at 4°C.
242	xv	i) Use 2.5–10 μl aliquots for PCR (see Section B.1.2 above).
243 244		e same PCR-based and LAMP assays detailed in Section B.1.2 have been also used for oocyst tection, with B1 gene and the 529RE being targets of choice (Slana <i>et al.</i> , 2021).
245	An	other important issue is the possibility of combining detection with information on oocyst viability.
246	Bio	passays, currently the only definitive way to detect viable oocysts, are expensive and relatively few
247 248		poratories have the necessary facilities. Reverse transcription (RT) real-time PCR (real-time RT- CR) or propidium monoazide-based real-time PCR have shown some promise for assessing oocyst
249		bility in complex sample matrices (Kim <i>et al.</i> , 2021; Rousseau <i>et al.</i> , 2018).
250	1.4.3. Nu	ucleic acid detection methods
251	Alt	hough single copy genes (e.g. SAG1, SAG2, SAG3, GRA6, and GRA7) have been used, multi-
252		py genes or genetic elements (e.g. 18S rDNA, B1, ITS1, 529RE) are preferred as they provide a
253 254		gher sensitivity (Dzib Paredes <i>et al.</i> , 2016; Slana <i>et al.</i> , 2021). For instance, 35 copies of the B1 ne and 200–300 copies of the 529 bp repetitive element (529RE) are present in the <i>T. gondii</i>
255		nome, and 10 to 100-fold higher sensitivity is generally observed in amplification targeting the
256		9RE compared with B1, although this also reflects the type of assay and sample being analysed
257 258		elaz <i>et al.</i> , 2015). In addition, some strains may have partially lost, or have a mutated, 529RE, and s could compromise diagnostic sensitivity (Wahab <i>et al.</i> , 2010).
259		proventional PCR targeting B1 was the first to be used in clinical diagnostics in people (Burg et al.,
260 261		89). To obtain details on <i>T. gondii</i> genotype (e.g. for outbreak investigation, infection source cing) the methods of choice are multi-locus PCR combined with restriction fragment length
261		lymorphism (PCR-RFLP) or sequencing and multi-locus microsatellite typing (Ajzenberg <i>et al.</i> ,
263		10; Su <i>et al.</i> , 2006).
264	То	increase sensitivity, several nested PCR assays have been implemented (Dzib Paredes et al.,
265		16). The reaction consists of two successive rounds of amplification. The product of the first
266		nplification serves as template for the second amplification, using one or two internal primers. The
267 268		k of cross- and carry-over contamination and false positives is increased with nested PCR, and ecautions should be taken to mitigate the risk (Dzib Paredes <i>et al.</i> , 2016).
	F.,	

269 270 271 272 273 274 275 276 277 278 279 280	There are several real-time PCR protocols and real-time PCR in combination with a hydrolysis probe is the most frequently applied (Slana <i>et al.</i> , 2021). This has largely improved both sensitivity and specificity of detection of <i>T. gondii</i> DNA, with the advantage of avoiding post-amplification manipulations and thus limiting the risk of carry-over contamination. Although sensitivity can be satisfactory with both conventional and real-time PCR using pure genomic <i>T. gondii</i> DNA, assay specificity might be affected when testing field samples. Conventional PCR can result in non-specific amplification, whereas this is not detected by real-time PCR due to the probe detection, despite the amplified target being the same. Furthermore, real-time PCR can be multiplexed and simultaneous amplification of an internal amplification control (a heterologous DNA fragment) can be used to monitor for the presence of inhibitors. In addition, amplifying two <i>T. gondii</i> -specific targets at once may increase sensitivity. Another advantage of real-time PCR is the possibility of quantification of <i>T. gondii</i> DNA.
281 282 283 284 285 286 287	A selective enrichment of target DNA combined with real-time PCR (i.e., magnetic capture PCR) has been reported to increase <i>T. gondii</i> detectability in animal samples (Gisbert Algaba <i>et al.</i> , 2017). The principle relies on separating and concentrating <i>T. gondii</i> DNA from sample DNA by specific DNA probes, complementary to the targeted parasite 529RE genomic region, which are conjugated to magnetic beads and followed by real-time PCR (Gisbert Algaba <i>et al.</i> , 2017). However, this method is expensive, time consuming, and requires further expertise, so might be not suitable for routine analysis or large surveys.
288 289 290 291 292 293 294 295 296 297 298	As an alternative to PCR, LAMP has been considered for <i>T. gondii</i> DNA detection, and diagnostic purposes, in environmental, veterinary, and human samples. LAMP takes advantage of a DNA polymerase (originally Bst) having both high strand displacement and replication activities. Nucleic acid amplification is performed under isothermal conditions (60–65°C), without the need of a DNA denaturation step. Both, B1 and 529RE have been widely used as targets in different LAMP assays, and LAMP is reported as comparable to real-time PCR for the detection of <i>T. gondii</i> in blood and animal tissues (Robert <i>et al.</i> , 2021). Although LAMP provides an opportunity for development of point-of-care testing or implementation of molecular tests in settings with limited facilities, there are several drawbacks including design of appropriate primers and the high risk of carry-over contamination. Commercial assays for both real-time PCR and LAMP are available for clinical diagnosis of toxoplasmosis.
299 300 301 302 303	Overall, the reported sensitivity of published molecular methods can be as low as one (or even less) genome equivalent per reaction. However, this largely depends on sample type, DNA extraction, copies of the targeted gene or sequence, amplification and detection reagents, procedures and platforms. The lack of accepted standard methods prevents robust comparison of sensitivity and specificity of the currently applied molecular tests.

304 2. Serological tests

There are several serological tests available for the detection of *T. gondii* antibodies (Table 1). All serological tests have limitations in diagnostic sensitivity and specificity and need proper validation to ensure confidence in results (refer to Chapter 1.1.6. *Principles and methods of validation of diagnostic assays for infectious disease* and Chapter 2.2.1. *Development and optimisation of antibody detection assays*).

The Sabin-Feldman dye test (DT) is a reference serological test for *T. gondii* antibody in humans (Dubey, 2022). Although the DT appears both specific and sensitive in humans, it is not extensively validated in other species. In addition, it is potentially hazardous as live parasite is used, is expensive, and requires a high degree of technical expertise.

312 2.1. Preparation of antisera and antigens

Antisera to *T. gondii* and conjugated antisera for use in IFAT and ELISA, to allow screening of a variety of animal species, may be obtained commercially. International standards for animal sera are not available.

Below are protocols for the preparation of tachyzoite antigen for use in the IFAT and ELISA. Tachyzoites may be grown in tissue culture and retained as whole parasites for the IFAT, or prepared as soluble antigen for the ELISA.

317	2.2.	Preparation of frozen stabilates of <i>T. gondii</i> tachyzoites			
318		2.2.1.	Tes	t procedure	
319 320 321			i)	Produce tachyzoites in tissue cell culture as described. Suitable <i>T. gondii</i> strains, able to multiply in cell culture, like the RH strain are available in a number of repositories; e.g. at the American Tissue Cell Culture Collection (ATCC ⁴⁹).	
322 323 324			ii)	Centrifuge three times at 500 g for 5 minutes and resuspend tachyzoites in Iscove's modified Dulbecco's medium (IMDM) or any other cell culture medium suitable to cultivate <i>T. gondii.</i> Final concentration of the tachyzoite suspension should be approximately 1.5×10^8 tachyzoites/ml.	
325 326 327 328			iii)	Combine dimethyl sulphoxide (DMSO), normal horse serum (free from antibody to <i>T. gondii</i>) and the tachyzoite suspension to give these final concentrations: 10% DMSO, 20% normal horse serum, 70% tachyzoite suspension; this gives a final concentration of approximately 1×10^8 tachyzoites/ml.	
329			iv)	Allow the preparation to stand on the bench for 1 hour (4-10°C; optimally on ice).	
330			v)	Dispense into 1-ml aliquots using screw-topped tubes appropriate for liquid nitrogen storage.	
331 332			vi)	Put the tubes into a small container, wrap in thick insulating material (e.g. paper towels) and place in –70°C freezer to allow the tachyzoites to freeze gradually.	
333			vii)	The next day transfer to liquid nitrogen, keeping well insulated while transferring.	
334 335			viii)	These stabilates may then be used for tissue culture growth of the parasite. When removing from storage, thaw the sample rapidly in a water bath (37°C).	
336 337 338			ix)	Centrifuge three times at 500 g for 5 minutes and resuspend tachyzoites in Iscove's modified Dulbecco's medium (IMDM) or any other cell culture medium suitable to cultivate <i>T. gondii</i> and add suspension to cell culture.	
339	2.3.	Produ	ction	of <i>Toxoplasma</i> tachyzoites in cell culture	
340		2.3.1.	Tes	t procedure	
341 342 343			i)	<i>Toxoplasma gondii</i> can be grown and maintained in tissue culture by twice-weekly passage in African green monkey kidney (Vero) cells. Other cell lines (e.g. MARC145 cells) are also suitable. Cell lines are available from repositories (e.g. ATCC).	
344 345			ii)	Cells and parasite are grown in IMDM supplemented with 50 IU/ml penicillin, 50 μ g/ml streptomycin, and 2% fetal bovine serum; there are other cell culture media suitable as well.	
346 347			iii)	Tachyzoites are harvested from tissue culture flasks by scraping the cell monolayer using a sterile cell scraper.	
348 349 350 351			iv)	Using 25 cm ² vented tissue culture flasks that have each been seeded with 1×10^5 Vero cells, add tachyzoites at the rate of two tachyzoites per monolayer cell and incubate at 37°C in a 5% CO ₂ humidified chamber. Harvest, when 2/3 of the cell layer was destroyed by tachyzoite multiplication, usually after 3–4 days.	
352	2.4.	Preparation of whole tachyzoites for IFAT and agglutination			
353		2.4.1.	Tes	t procedure	
354			i)	Produce 4 × 10^7 /ml suspension of <i>T. gondii</i> tachyzoites in PBS.	
355			ii)	Add formaldehyde (40%) to give a final concentration of 0.2% (v/v).	
356 357			iii)	Incubate at 4°C overnight and divide into aliquots in suitable tubes and store frozen until required (–20°C).	
358					

⁴⁹ American Type Culture Collection, P.O. Box 1549, Manassas, Virginia 20108, United States of America.

359	2.5. Proc	luction	of soluble antigen for ELISA
360	2.5.1	. Tes	t procedure
361		i)	Produce a suspension of <i>T. gondii</i> tachyzoites in PBS.
362 363		ii)	Centrifuge at 2000 g for 15 minutes, retain the pellet and resuspend it in nine times its volume of distilled water.
364		iii)	Lyse the tachyzoites by freezing and thawing three times.
365		iv)	Sonicate the antigen preparation for 20 seconds at 4°C at an amplitude of 20 microns.
366		v)	Remove any cellular debris by centrifugation at 10,000 g for 30 minutes at 4°C.
367 368		vi)	Retain the supernatant and store at -20° C until required; protein estimation should be between 2 and 4 µg/ml.
369	2.6. Indii	rect flu	orescent antibody test
370 371 372 373 374 375 376 377	Toxoplasma antibody is specific sec commerciall find some sp nuclear antil	tachyz added, condary y availat pecies-s bodies.	cent antibody test (IFAT) (Dubey, 2022) is a simple and widely used method. Whole, killed oites are incubated with diluted test serum, the appropriate fluorescent labelled secondary and the result is then viewed with a fluorescence microscope. Fluorescent-labelled species-antibodies are available commercially, the method is relatively inexpensive, and kits are ble. However, the results are read by eye, so subjective variation may occur. It may be difficult to pecific conjugates and there is a risk of possible cross-reactivity with rheumatoid factor and anti-The following is a protocol for carrying out an IFAT for anti- <i>Toxoplasma</i> IgG antibodies in sheep es minor modifications for testing different species or for measuring IgM antibody.
378	2.6.1	. Tes	t procedure
379 380		i)	Clean the required number of multi-well immunofluorescence assay slides (e.g. 10–21 well- slides with wells of 4–6 mm in diameter are suitable) and allow to dry.
381 382		ii)	Apply 5 μ I of a whole tachyzoite preparation (Section B.2.4.1 above) on to each well and allow to air dry.
383		iii)	Fix in methanol for 10 minutes.
384		iv)	Wash twice (10 minutes for each wash) in 0.3 M PBS, pH 7.4.
385		V)	Prepare serial dilutions of the test sera in PBS (e.g. 1/16, 1/32, etc. up to 1/1024).
386 387 388		vi)	Add 5 µl of the given test sheep serum (diluted in PBS) to each well. Ensure that positive and negative control sera are included in each test as well as a 'PBS-only' sample. Incubate for 30 minutes at room temperature.
389		vii)	Wash twice (10 minutes each time) in PBS.
390 391 392		viii)	Add 5 μ I of an appropriate dilution of rabbit-anti-sheep IgG conjugated to fluorescein isothiocyanate, diluted in 0.2% filtered Evans blue dye in PBS (filtered through a 0.45 μ m sterile filter), to each well and incubate for 30 minutes at room temperature.
393		ix)	Wash three times for 10 minutes each time in PBS.
394		x)	Mount the slides under cover-slips with buffered glycerol (nine parts PBS, one part glycerol).
395		xi)	Examine using a fluorescence microscope, fitted with ×20 and ×40 objective lenses.
396 397 398 399 400 401 402 403 404		the (nor leas In a lam shou fluor	In a negative test serum result, the whole parasites will appear red due to the autofluorescence of Evans blue dye. They may also present with a green fluorescent cap at the parasite pole hspecific polar fluorescence). With a positive test serum, the parasites will fluoresce red and at t 80% of them within a given well will be surrounded by an unbroken band of green fluorescence. In adult sheep/goat a positive titre could be defined as $\geq 1/64$ and a negative titre as $\leq 1/32$. For b/kid and fetal sera, respective titres could be defined as $\geq 1/32$ and $\leq 1/16$. These cut off values uld be validated locally as results may vary between laboratories, depending on, e.g., the rescence microscope and the operator. Optimally, each slide should include positive control and ative controls.

405 **2.7. Modified agglutination test**

The modified agglutination test (MAT) (Dubey, 2022) is both sensitive and specific. Formalinised Toxoplasma 406 tachyzoites are added to U-shaped well microtiter plates and dilutions of test sera are then applied. Positive samples 407 will produce agglutination that can be graded, whereas negative samples will produce a 'button' of precipitated 408 409 tachyzoites at the bottom of the well. The test is simple and easy to perform although relatively large amounts of antigen are required. Kits are commercially available. It is important to treat sera with 0.2 2-mercaptoethanol to avoid 410 411 false positives due to non-specific IgM. The MAT has been used extensively for detection of T. gondii antibodies in 412 sera of many animal species and the procedure is detailed below. A commercially available latex agglutination test is also available, but this test is regarded relatively insensitive compared with MAT or IFAT. 413

414 **2.7.1. Ser**u

415

416 417

418

419

420

421

422

423

424

425

426

427

429

430 431

432

433

434 435

436

437 438

439 440

441

442 443

444

445

Serum-diluting buffer

- i) Dissolve 42.5 g NaCl, 1.54 g NaH₂PO₄, and 5.4 g Na₂HPO₄ in 900 ml deionised water.
 - ii) Adjust the pH to 7.2. Bring the volume to 1 litre with deionised water.
 - iii) Store in a refrigerator. This is the 5× stock solution.
- iv) Dilute this stock solution 1/5 to give 0.01 M PBS (working serum-diluting buffer: 1 part stock and 4 parts deionised water). PBS should be filtered through a 0.22 µM filter immediately prior to use.
- 2.7.2. Antigen-diluting buffer
 - i) Prepare a stock of borate buffer: dissolve 7.01 g sodium chloride, 3.09 g boric acid, 2.0 g sodium azide in 900 ml deionised water.
 - ii) Add 24 ml 1 N NaOH and adjust the pH to 8.9.
 - iii) Bring the volume to 1 litre. This is the stock solution and can be stored at room temperature.
 - iv) For the working antigen-diluting buffer, dissolve 0.4 g bovine serum albumin in 100 ml borate buffer. Store at 4°C.

428 2.7.3. Serum dilutions

 Dilute serum samples with working serum-diluting buffer (Section B.2.7.1 above) in small test tubes (1.2 ml in strips of 8 or 12) with a multichannel pipette, starting at 1/25. Note: Microtiter plates may also be used for making serum dilutions.

2.7.4. Preparation of antigen mixture

For each plate, mix 2.5 ml working antigen-diluting buffer (see Sections B.2.4.1 and B.2.7.2 above), 35 μl 2-mercaptoethanol, 50 μl Evans blue dye solution (2 mg/ml water) and 0.15 ml antigen (formalin-fixed whole parasites).

2.7.5. Agglutination procedure

Agglutination is done in U-bottom 96-well microtiter plates.

- i) Pipette 25 µl antigen mixture to each well immediately after mixing.
- ii) Pipette 25 µl serum dilutions into the wells and mix gently with the antigen by repeated pipetting action.
- iii) A positive control should be included in each plate. The control should have a titre of 1/200, and two-fold dilutions from 1/25 to 1/3200 should be used.
 - iv) Cover the plates with sealing tape and incubate overnight at 37°C.
 - v) Read results using a magnifying mirror. A blue button at the bottom of the well means negative. A clear bottom means positive.

446 **2.8. Enzyme-linked immunosorbent assay**

447 One of the first *T. gondii* enzyme-linked immunosorbent assay (ELISA) (Voller *et al.*, 1976) used a soluble antigen 448 preparation made from *T. gondii* RH strain tachyzoites (as described below) and layered into wells in an ELISA 449 microtiter plate. Test sera are added, followed by a species-specific secondary antibody conjugated with a reactive 450 enzyme, such as horseradish peroxidase. Protein A/G conjugates were used to replace species-specific antibody 451 conjugates, making ELISAs applicable to more than one animal species. Any conjugated enzyme causes a colour 452 change in the substrate that is directly related to the amount of bound antibody, and which can be read with a 453 spectrophotometer at the absorbance wavelength specific to the substrate used. The assay is simple, can readily test 454 a large number of samples, and is easy to perform with the chosen anti-species conjugate. Defined anti-species 455 conjugates, substrates, and whole kits are commercially available. The ELISA is well suited for analysing large 456 numbers of samples. A large number of species-specific or multi-species ELISAs are commercially available to detect 457 *T. gondii* antibodies.

To improve the specificity of the conventional ELISA, native purified *T. gondii*-specific antigens have been used (Basso *et al.*, 2013). In addition, severam recombinant antigens have been established, and these seem suitable for replacing native antigens for serological diagnostic tests. For many of these recombinant antigen ELISAs, thorough validation is lacking.

462 Clinically, there is a need to distinguish between recent (acute) and long-standing (chronic) infections. With the 463 conventional ELISA, detection of *Toxoplasma*-specific IgG and IgM antibodies, along with IgA, may permit some 464 discrimination between acute and chronic *T. gondii* infection. Assays assessing the avidity of an IgG response to 465 *T. gondii* have been applied in sheep and pigs. However, such avidity tests were used for research purpose only.

466

C. REQUIREMENTS FOR VACCINES

467 Currently there is only one commercially available live vaccine, which is licensed for use in breeding sheep in some regions (Europe and New Zealand) to reduce the effects of T. gondii infection (e.g. early embryonic death, abortion). It consists of 468 469 \geq 10⁵ tachyzoites of the S48 strain of *T. gondii* that has been attenuated by multiple passages in mice. The vaccine 470 stimulates effective protective immunity for at least 18 months following a single intramuscular injection given at least 4 weeks prior to mating and only for use in healthy, non-pregnant female sheep. Despite the acknowledged importance of 471 472 human toxoplasmosis, human vaccines are currently unavailable, and the vaccine for sheep has disadvantages, such as a short shelf-life (10 days), strict storage conditions, and, as a live vaccine, potential risk to operators. Information on the 473 production details of this vaccine and QC requirements are not available. 474

Although the importance of a killed or non-live vaccine is acknowledged – for vulnerable humans (e.g. women before they are pregnant), for reducing abortions in sheep, for reducing tissue cysts in meat animals (pigs, cattle, chickens, etc.), and for limiting oocyst shedding from kittens – to date this remains elusive (Innes *et al.*, 2019). However, with recent scientific advances, including availability of genetic, transcriptomic, and metabolomic data, the potential for developing knockout variants, and other new technologies suggest new possibilities for development of such a vaccine (Mevelec *et al.*, 2020; Zhang *et al.*, 2022).

481

REFERENCES

AJZENBERG D., COLLINET F., MERCIER A., VIGNOLES P. & DARDE M.L. (2010). Genotyping of *Toxoplasma gondii* isolates with
 15 microsatellite markers in a single multiplex PCR assay. *J. Clin. Microbiol.*, 48, 4641–4645.

BASSO W., HARTNACK S., PARDINI L., MAKSIMOV P., KOUDELA B., VENTURINI M.C., SCHARES G., SIDLER X., LEWIS F.I. & DEPLAZES
P. (2013). Assessment of diagnostic accuracy of a commercial ELISA for the detection of *Toxoplasma gondii* infection in pigs compared with IFAT, TgSAG1-ELISA and Western blot, using a Bayesian latent class approach. *Int. J. Parasitol.*, 43, 565–570.

BELAZ S., GANGNEUX J.P., DUPRETZ P., GUIGUEN C. & ROBERT-GANGNEUX F. (2015). A 10-year retrospective comparison of
 two target sequences, REP-529 and B1, for *Toxoplasma gondii* detection by quantitative PCR. *J. Clin. Microbiol.*, **53**, 1294–
 1300.

491 DUBEY J.P. (2021). Outbreaks of clinical toxoplasmosis in humans: five decades of personal experience, perspectives and 492 lessons learned. *Parasit. Vectors*, **14**, 263.

493 DUBEY J.P. (2022). Toxoplasmosis of Animals and Humans, Third Edition. CRC Press, Boca Raton, USA.

494 DZIB PAREDES G.F., ORTEGA-PACHECO A., ROSADO-AGUILAR J.A., ACOSTA-VIANA K.Y., GUZMÁN-MARÍN E. & JIMÉNEZ-COELLO M.

495 (2016). Toxoplasma gondii in meat for human consumption – A brief review of the most described strategies for its detection
 496 and quantification, significance, prevention and control of food related diseases. IntechOpen, London, UK.

- EUROPEAN FOOD SAFETY AUTHORITY (EFSA) PANEL ON BIOLOGICAL HAZARDS (BIOHAZ), KOUTSOUMANIS K., ALLENDE A.,
 ALVAREZ-ORDÓÑEZ A., BOLTON D., BOVER-CID S., CHEMALY M., DAVIES R., DE CESARE A., HERMAN L., HILBERT F., LINDQVIST R.,
 NAUTA M., PEIXE L., RU G., SIMMONS M., SKANDAMIS P., SUFFREDINI E., CACCIÒ S., CHALMERS R., DEPLAZES P.,
 DEVLEESSCHAUWER B., INNES E., ROMIG T., VAN DER GIESSEN J., HEMPEN M., VAN DER STEDE Y. & ROBERTSON L. (2018). Public
 health risks associated with food-borne parasites. *EFSA Journal*, **16**, e05495.
- GISBERT ALGABA I., GEERTS M., JENNES M., COUCKE W., OPSTEEGH M., COX E., DORNY P., DIERICK K. & DE CRAEYE S. (2017).
 A more sensitive, efficient and ISO 17025 validated Magnetic Capture real time PCR method for the detection of archetypal
 Toxoplasma gondii strains in meat. *Int. J. Parasitol.*, **47**, 875–884.
- HATAM-NAHAVANDI K., CALERO-BERNAL R., RAHIMI M.T., PAGHEH A.S., ZAREAN M., DEZHKAM A. & AHMADPOUR E. (2021).
 Toxoplasma gondii infection in domestic and wild felids as public health concerns: a systematic review and meta-analysis.
 Sci. Rep., **11**, 9509.
- 508 INNES E.A., HAMILTON C., GARCIA J.L., CHRYSSAFIDIS A. & SMITH D. (2019). A One Health approach to vaccines against 509 *Toxoplasma gondii. Food Waterborne Parasitol.*, **15**, e00053.
- 510 JAUREGUI L.H., HIGGINS J., ZARLENGA D., DUBEY J.P. & LUNNEY J.K. (2001). Development of a real-time PCR assay for 511 detection of *Toxoplasma gondii* in pig and mouse tissues. *J. Clin. Microbiol.*, **39**, 2065–2071.
- 512 KIM M., SHAPIRO K., RAJAL V.B., PACKHAM A., AGUILAR B., RUEDA L. & WUERTZ S. (2021). Quantification of viable protozoan 513 parasites on leafy greens using molecular methods. *Food Microbiol.*, **99**, 103816.
- LALLE M., POSSENTI A., DUBEY J.P. & POZIO E. (2018). Loop-Mediated Isothermal Amplification-Lateral-Flow Dipstick (LAMP-LFD) to detect *Toxoplasma gondii* oocyst in ready-to-eat salad. *Food Microbiol.*,**70**, 137-142.
- 516 LINDQUIST H.D., BENNETT J.W., HESTER J.D., WARE M.W., DUBEY J.P. & EVERSON W.V. (2003). Autofluorescence of 517 *Toxoplasma gondii* and related coccidian oocysts. *J. Parasitol.*, **89**, 865–867.
- LORENZI H., KHAN A., BEHNKE M.S., NAMASIVAYAM S., SWAPNA L.S., HADJITHOMAS M., KARAMYCHEVA S., PINNEY D., BRUNK B.P.,
 AJIOKA J.W., AJZENBERG D., BOOTHROYD J.C., BOYLE J.P., DARDE M.L., DIAZ-MIRANDA M.A., DUBEY J.P., FRITZ H.M., GENNARI
 S.M., GREGORY B.D., KIM K., SAEIJ J.P., SU C., WHITE M.W., ZHU X.Q., HOWE D.K., ROSENTHAL B.M., GRIGG M.E., PARKINSON
 J., LIU L., KISSINGER J.C., ROOS D.S. & SIBLEY L.D. (2016). Local admixture of amplified and diversified secreted
 pathogenesis determinants shapes mosaic *Toxoplasma gondii* genomes. *Nat. Commun.*, 7, 10147.
- MEVELEC M.-N., LAKHRIF Z. & DIMIER-POISSON I. (2020). Key Limitations and New Insights Into the Toxoplasma gondii
 Parasite Stage Switching for Future Vaccine Development in Human, Livestock, and Cats. *Front. Cell. Infect. Microbiol.*,
 10 (697) doi: 10.3389/fcimb.2020.607198.
- 526 PAPPAS G., ROUSSOS N. & FALAGAS M.E. (2009). Toxoplasmosis snapshots: global status of *Toxoplasma gondii* 527 seroprevalence and implications for pregnancy and congenital toxoplasmosis. *Int. J. Parasitol.,* **39**, 1385–1394.
- 528 ROBERT M.G., BRENIER-PINCHART M.-P., GARNAUD C., FRICKER-HIDALGO H. & PELLOUX H. (2021). Molecular diagnosis of 529 toxoplasmosis: recent advances and a look to the future. *Expert Rev. Anti-Infe.*, **19**, 1529–1542.
- ROUSSEAU A., LA CARBONA S., DUMETRE A., ROBERTSON L.J., GARGALA G., ESCOTTE-BINET S., FAVENNEC L., VILLENA I., GERARD
 C. & AUBERT D. (2018). Assessing viability and infectivity of foodborne and waterborne stages (cysts/oocysts) of *Giardia duodenalis*, *Cryptosporidium* spp., and *Toxoplasma gondii*: a review of methods. *Parasite*, 25, 14.
- SCHARES G., GLOBOKAR VRHOVEC M., TUSCHY M., JOERES M., BARWALD A., KOUDELA B., DUBEY J.P., MAKSIMOV P. & CONRATHS
 F.J. (2021). A real-time quantitative polymerase chain reaction for the specific detection of *Hammondia hammondi* and its
 differentiation from *Toxoplasma gondii*. *Parasit. Vectors*, **14**, 78.
- 536 SHAPIRO K., BAHIA-OLIVEIRA L., DIXON B., DUMÈTRE A., DE WIT L.A., VANWORMER E. & VILLENA I. (2019). Environmental 537 transmission of *Toxoplasma gondii*: Oocysts in water, soil and food. *Food Waterborne Parasitol.*, **15**, e00049-e00049.

SHWAB E.K., SARAF P., ZHU X.Q., ZHOU D.H., MCFERRIN B.M., AJZENBERG D., SCHARES G., HAMMOND-ARYEE K., VAN HELDEN
P., HIGGINS S.A., GERHOLD R.W., ROSENTHAL B.M., ZHAO X., DUBEY J.P. & SU C. (2018). Human impact on the diversity and
virulence of the ubiquitous zoonotic parasite *Toxoplasma gondii*. *Proc. Natl Acad. Sci. USA*, **115**, E6956-E6963.

SLANA I., BIER N., BARTOSOVA B., MARUCCI G., POSSENTI A., MAYER-SCHOLL A., JOKELAINEN P. & LALLE M. (2021). Molecular
 methods for the detection of *Toxoplasma gondii* oocysts in fresh produce: An extensive review. *Microorganisms*, 9 doi:
 10.3390/microorganisms9010167.

544 SUC., ZHANG X. & DUBEY J.P. (2006). Genotyping of *Toxoplasma gondii* by multilocus PCR-RFLP markers: a high resolution 545 and simple method for identification of parasites. *Int. J. Parasitol.*, **36**, 841–848.

TORGERSON P.R., DEVLEESSCHAUWER B., PRAET N., SPEYBROECK N., WILLINGHAM A.L., KASUGA F., ROKNI M.B., ZHOU X.N.,
FEVRE E.M., SRIPA B., GARGOURI N., FURST T., BUDKE C.M., CARABIN H., KIRK M.D., ANGULO F.J., HAVELAAR A. & DE SILVA N.
(2015). World Health Organization Estimates of the Global and Regional Disease Burden of 11 Foodborne Parasitic
Diseases, 2010: A Data Synthesis. *PLoS Med*, **12** (12), e1001920.

- 550 VOLLER A., BARTLETT A. & BIDWELL D.E. (1976). Enzyme immunoassays for parasitic diseases. *Trans. Royal Soc. Trop.* 551 *Med. Hyg.*, **70**, 98–106.
- 552 WAHAB T., EDVINSSON B., PALM D. & LINDH J. (2010). Comparison of the AF146527 and B1 repeated elements, two real-time 553 PCR targets used for detection of *Toxoplasma gondii*. J. Clin. Microbiol., **48**, 591–592.
- 554 ZHANG Y., LI D., LU S. & ZHENG B. (2022). Toxoplasmosis vaccines: what we have and where to go? *NPJ Vaccines*, 7, 131.

555 556	* *
557 558	NB: At the time of publication (2024) there were no WOAH Reference Laboratories for toxoplasmosis (please consult the WOAH Web site:
559	https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).
560	NB : First adopted in 2004. Most recent updates adopted in 2017.

Annex 19. Analysis of the questionnaire for Reference Laboratories

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4-8 September 2023



Kindly click on the image above to gain access to the entire presentation.

Annex 20. List of Main Focus Areas and Specialties for WOAH Collaborating Centres

2

13

14

15

16

17 18

19

20

22

32

34

46 47

The role of WOAH Collaborating Centres is anchored to the WOAH's founding mandate and to the Seventh Strategic Plan
 (2021–2025)⁵⁰.

5 1. Animal health management

6 WOAH has the responsibility to collect, analyse and disseminate relevant scientific information, especially on disease 7 control methods, and to provide expertise in the control of animal diseases including zoonotic diseases, as well as 8 <u>health threats</u> at the animal-human-ecosystems interface, while taking into account <u>under</u> the "One Health" <u>approach</u> 9 concept whenever possible. This topic covers issues primarily, but not exclusively, related to Sections 2 and 4 of the 10 *Terrestrial* and *Aquatic Codes* and to Parts 3 <u>of the *Terrestrial Manual*</u> and <u>Part</u> 2 of both the *Terrestrial* and *Aquatic* 11 *Manuals*, respectively. The topic is expected to assist WOAH and its Members to fulfil the core missions of the 12 organisation.

- Disease control prevention, risk assessment, early warning and preparedness
- Species related (e.g. molluscs, bees, camelids)
- Preventing animal disease Biosecurity and disease prevention along the value-chain biosecurity
- Emerging animal diseases (early detection, warning and response)
- Animal health emergencies
- Zoonotic diseases
 - Epidemiology, modelling, surveillance
 - Social and economic implications of animal diseases <u>control</u>
- Biothreat reduction
 - Impact of climate change on animal health

23 2. Animal production

WOAH's founding mandate has evolved and has been adapted to Members' needs, it now includes improving the safety of food of animal origin from hazards originating in animal production, and establishing standards and guidelines for animal welfare through a science-based approach and promote their application. This topic corresponds to this mandate and more specifically to Section 7 of the *Terrestrial* and *Aquatic Codes* on animal welfare, and the relevant provisions on food and feed safety in the chapters in Section 6 on Veterinary Public Health of the *Terrestrial Code* (Chapters 6.1, 6.2, 6.3, 6.5, 6.12, 6.13) and Chapter 4.8 <u>4.9</u> of the *Aquatic Code*.

- 30 Animal welfare
- Animal production food safety
 - Sustainable animal production (including integrated health management)
- Safety of animal feed
 - Climate change and impacts

35 3. Laboratory expertise

This topic covers issues related to management and operation of veterinary diagnostic laboratories. It corresponds essentially to provisions of Chapters 1.1.1 to 1.1.7 of the *Terrestrial Manual*, as well as Chapter 2.1.2, and to Chapters 1.1.1 and 1.1.2 of the *Aquatic Manual*. Beyond WOAH standards, the topic is expected to assist WOAH and its Members to follow the recommendations of the first two International Conferences on Biological Threat Reduction, as well as to contribute to the Seventh WOAH Strategic Plan and commitment to modern technology.

- Biorisk management
- Quality management systems
- Biobanking and reference collections
- Genomics and bioinformatics
- 45 Laboratory information systems technology
 - Validation procedures for diagnostic tests of laboratory methods
 - Development and application of innovative technologies

⁵⁰ https://www.woah.org/en/document/seventh-strategic-plan/

48 **4.** Training and education

It is part of the WOAH's founding mandate to improve the legal framework, competency and resources of national Veterinary Services, and particularly their global public good components. This topic covers the scientific and technical veterinary knowledge and skills needed for veterinarians, animal health professionals and veterinary paraprofessionals to implement WOAH Standards. The topic primarily, but not exclusively, corresponds to provisions of the Section 3 of the *Terrestrial* and *Aquatic Codes*. The topic is also expected to assist the WOAH and its Members to follow-up on the recommendations of the first two-International Conferences of Veterinary Education.

- Veterinary <u>education</u> (under<u>- and post-graduate</u>) education
 - Veterinary-Post-graduate training and education (scientific and technical) and capacity building
 - Veterinary specialisation and Laboratory or epidemiological expertise in infectious diseases
 - Capacities-Capacity building of Veterinary Services or Aquatic Animal Health Services

59 5. Veterinary products

56

57

58

65

66 67

68

85

This topic corresponds to Chapters 1.1.8 to 1.1.10, and most of the specific recommendations included in the Part 2 of the *Terrestrial Manual*. Progress made on vaccines, diagnostics and the development of new drugs is believed to contribute to the global efforts against antimicrobial resistance. As for antimicrobial resistance, the topic also corresponds to Chapters 6.1 to 6.4 of the *Aquatic Code*, Chapters 6.6 to 6.10 of the *Terrestrial Code*, and Chapter 2.1.1 of the *Terrestrial Manual*.

- Vaccines, diagnostics (kits), and drugs
- <u>Managing</u> antimicrobial agents resistance
- <u>Alternatives to antimicrobials</u>
 - New technologies

69 6. <u>Wildlife health and biodiversity</u>-Environment and climate change

70 WOAH provides expertise to Members in understanding and managing the effects of environmental and climate changes on animal health and welfare. Climate change is likely to increase pressure on animal production, and 71 72 provide newly suitable conditions for invasive pests and pathogens. The risk of emergence of new pathogens has increased as a consequence of global changes in the way food is produced, moved and consumed. This topic is 73 expected to address animal health issues, including aquatic animals, connected to wildlife, biodiversity, climate 74 change, and emerging risks. Wildlife plays a vital role in maintaining healthy and functioning ecosystems, thus 75 76 contributing to the preservation of biodiversity. Wildlife is an asset, supporting livelihoods through the provision of 77 income, whether it be through tourism or as a source of food. Importantly, wildlife has a positive effect on human wellbeing, contributing to education, physical and mental health, social values, culture and spirituality. This topic is 78 79 expected to address animal health issues, including terrestrial and aquatic animals, connected to wildlife, biodiversity, 80 and emerging risks.

- 81 Threats to livestock or wildlife health and welfare and biodiversity
- 82 Impact of climate change and biodiversity on wildlife
- 83 Disease related (including vector-borne)
- <u>Wildlife disease epidemiology, modelling, and surveillance</u>
 - Transboundary diseases at the ecosystem/human/domestic animal health interface
- Drivers for emerging risks
- Wildlife's role in the epidemiology of diseases affecting livestock and humans, and in disease emergence at the human–animal interface

© World Organisation for Animal Health (WOAH), 2023

This document has been prepared by specialists convened by the World Organisation for Animal Health (WOAH). Pending adoption by the World Assembly of Delegates, the views expressed herein can only be construed as those of these specialists.

All WOAH publications are protected by international copyright law. Extracts may be copied, reproduced, translated, adapted or published in journals, documents, books, electronic media and any other medium destined for the public, for information, educational or commercial purposes, provided prior written permission has been granted by the WOAH.

The designations and denominations employed and the presentation of the material in this publication do not imply the expression of any opinion whatsoever on the part of the WOAH concerning the legal status of any country, territory, city or area or of its authorities, or concerning the delimitation of its frontiers and boundaries.

The views expressed in signed articles are solely the responsibility of the authors. The mention of specific companies or products of manufacturers, whether or not these have been patented, does not imply that these have been endorsed or recommended by the WOAH in preference to others of a similar nature that are not mentioned.