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 7 December 2023 to be considered at the February 2023 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

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List of Annexes

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 step-wise 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.

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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 Annexes 1 and 2 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 ELISA[2] (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.

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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 FMD[3] 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*.

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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 Annex 4 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 Annex 5 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 Annex 6 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 Annex 7 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 PCR[4], 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*

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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 Annex 12 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 POC[5] 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 Annex 13 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 ASF[6] 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 Annex 16 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.

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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 PD$_{50}$ 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 ToR[7] 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)

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7   ToR: Terms of Reference

Federal Research Institute for Animal Health
Naumburger Str. 96a – 07743 Jena,
GERMANY
Tel.: (49-3641) 804.2243; E-mail: herbert.tomaso@fli.de
Website: Institute of Bacterial Infections and Zoonoses (IBIZ): Friedrich-Loeffler-Institut (fli.de)
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: pdebenedictis@izsvenezie.it
Website: https://www.izsvenezie.it/
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: Epidemiology, Parasites and Vectors (arc.agric.za)
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 Annex 19 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 (Collaborating Centre - Procedures for Designation). 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 5-year 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, PPR[8] 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 overview of ASF diagnostic tests for field application (POC tests).

Peste des petits ruminants

The WOAH PPR Reference Laboratory network continues to regularly update its website 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/IAEA[9] 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 FAO[10] 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 WOAH website 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 continue to support international initiatives, in particular the United Against Rabies Forum activities, and have 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 developed guidance 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.

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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 Annex 20.

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 AFNOR[11], on behalf of CEN/TC[12] 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 standards[13], antigen standards[14] and PCR assays[15], 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.

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11 AFNOR: Association française de normalisation
12 CEN/TC: European Committee for Standardization/Technical Committee
13 https://www.woah.org/app/uploads/2021/03/a-guideline-antibody-standards.pdf
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 Animal Health Forum (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 Strategic Challenges in the Global Control of High Pathogenicity Avian Influenza 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 Resolution 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-TADs[16] 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 adopted Resolution, 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.

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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 February 2023 WHO[18] 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 avian influenza report, they collected 795 avian influenza virus sequences of H5, 34 of H7, and 305 of H9 subtypes. Additionally, for the swine influenza report 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 OFFLU avian influenza matching (AIM) initiative 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

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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 RVCM[19] holdings around the world, with the exception of diagnostic materials and vaccines. All seven FAO-WOAH designated RHFs[20] 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 EuFMD[21].

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 VICH[22] 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 Biologic 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.

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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 report 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 IHSC[23]-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.

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<div align="right">…/Annexes</div>

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23 IHSC: International Horse Sports Confederation

Annex 1. Adopted Agenda

MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION

Paris, 4–8 September 2023

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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 PD_{50} 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 risk-based 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 IHSC[24]-WOAH collaboration agreement and consultancy project in Asia
    11.10. Update on the virtual biobank project

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24 IHSC: International Horse Sports Confederation

Annex 2. List of Participants

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4–8 September 2023

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MEMBERS OF THE COMMISSION

Prof. Emmanuel Couacy-Hymann
(President)
Professor of Virology
CNRA/LIRED
Abidjan
CÔTE D'IVOIRE

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

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

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

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

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

Ms Sara Linnane
Scientific Officer
Science Department

Dr Gounalan Pavade
Scientific Coordinator
Science Department

Dr Charmaine Chng
Deputy Head
Science Department

Dr Mariana Delgado
Scientific Secretariat Officer
Science Department

Annex 3. Work Programme for the WOAH Biological Standards Commission

MEETING OF THE BIOLOGICAL STANDARDS COMMISSION

Paris, 4–8 September 2023

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Subject	Issue	Status and Action
Updating the *Terrestrial Manual*	1) Circulate the chapters approved by the BSC to Members for first-round comment	October 2023
	2) 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 *Terrestrial Manual*	On-going
	a) Finalise the template for the validation report for current and future tests recommended in the *Terrestrial Manual* and make available to Reference Laboratory experts for submission for new or existing tests	December 2023
	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 *Test methods available and their purpose* 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
	6) Develop criteria for removing chapters for nonlisted diseases and assess those chapters against the criteria	For February
	7) Review new developments in diseases causing significant global impacts (e.g. avian influenza, African swine fever) and prioritise those chapters	On-going
Collaborating Centres	1) Implementation of the adopted SOPs:	
	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
	2) Review the designations of those Centres that completed 5 years	February 2025
	3) 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
Reference Laboratories	1) Put under-performing labs on watch list	On going
	2) Implement the new system for evaluating annual reports and provide list of assigned reports to BSC members	For October 2023
	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	1) Follow up with the three newly launched Reference Laboratory networks (ASF, PPR and rabies)	On-going
Standardisation/ Harmonisation	1) Project to extend the list of WOAH-approved reference reagents	On-going
	2) 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
	3) 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, Workshops and Meetings with participation by BSC Members	1) Biosafety research roadmap	On-going
	2) ISWAVLD OIE Seminar: develop a theme and programme and speakers	June 2025 in Canada
Performance	1) Engage with the ongoing processes around performance issues with Reference Laboratories	On-going
Develop laboratory standards for emerging diseases	1) Discuss the *Terrestrial Code* chapter once adopted with the aim of introducing a corresponding chapter for the *Terrestrial Manual*	After May 2024
Case definitions	1) Follow up the implementation of the SOPs for case definitions	On-going

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2   **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

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6   # C H A P T E R   1 . 1 . 5 .

7
8   # QUALITY MANAGEMENT IN VETERINARY TESTING LABORATORIES

9   ## SUMMARY

10   _Valid laboratory results are essential for diagnosis, surveillance, and trade. Such results are ~~achieved by the~~_
11   _~~use~~ assured through implementation of ~~good~~ a management ~~practices, valid~~ system that supports accurate_
12   _and consistent test and calibration methods~~, proper techniques, quality control and quality assurance, all~~_
13   _~~working together within a quality management system~~. Laboratory quality management includes technical,_
14   _managerial, and operational elements of ~~testing~~ performing, interpreting and ~~the interpretation of~~ reporting_
15   _test results. A quality management system enables the laboratory to demonstrate both competency and an_
16   _ability to generate consistent technically valid results that meet the needs of its customers. ~~The need for~~_
17   _Mutual recognition and acceptance of test results for international trade, and the ~~acceptance~~ accreditation_
18   _of tests to international standards such as ISO/IEC [25] 17025~~:2005~~ (General Requirements for the_
19   _Competence of Testing and Calibration Laboratories) (ISO/IEC, ~~2005~~ 2017b) requires ~~good~~ suitable_
20   _laboratory quality management systems. This chapter is not intended to reiterate the requirements of_
21   _ISO/IEC 17025, nor has it been endorsed by accreditation bodies. Rather, it outlines the important issues_
22   _and considerations a laboratory should address in the design and maintenance of its quality management_
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_
26   _necessary for the effective delivery of diagnostic services, and highlights the critical elements that should be_
27   _established as minimum requirements._

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

30   To ensure that the quality management system is appropriate and effective, the design must be carefully ~~thought out~~
31   planned and, where accreditation is sought, must address all criteria of the appropriate quality standard. The major
32   ~~categories of~~ consideration~~s~~ and ~~the~~ their associated key issues and activities ~~within each of these categories~~ are outlined
33   in the following eight sections of this chapter.

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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:

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 /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;

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 Role 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 Role in assisting with, confirming, or overseeing the work of other laboratories (e.g. as a reference laboratory);

viii) Business goals of the laboratory, including the need for any third-party recognition or accreditation.

2. Standards, guides, and references

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, and for all WOAH Reference Laboratories, the use of ISO/IEC 17025 (ISO/IEC, 2005 2017b) or equivalent will be is 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 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 American Association for Laboratory Accreditation (A2LA), etc. Technical and international organisations such as AOAC International (The Scientific Association Dedicated to Analytical Excellence; formerly the Association of Official Analytical Chemists) and the International Organization for Standardization (ISO) publish useful references, guides, application documents and standards that supplement the general requirements of ISO/IEC 17025. Other relevant documents may include guides and application documents providing interpretative criteria and recommendations for the application of ISO/IEC 17025 in the field of veterinary testing for both applicant and accredited facilities, e.g. https://www.nata.com.au/phocadownload/spec-criteria-guidance/animal-health/Animal-Health-ISO-IEC-17025-Appendix.pdf https://nata.com.au/files/2021/05/Animal-Health-ISO-IEC-17025-Appendix-effective-March2021.pdf; 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 a underpin a laboratory quality system, fulfilment of its requirements does not necessarily ensure or imply assure technical competence (in the areas listed in Section 3 *Accreditation*). Conformance to the requirements of ISO 9001 is assessed by a certification body that is accredited to undertake such assessments by the national accreditation body to undertake such assessments. 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).

3. Accreditation

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

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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
85 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~~ requires:

89 i) Adequate facilities and environmental controls;

90 ii) ~~Has~~ Appropriately qualified and trained personnel with a depth of technical knowledge commensurate with
91   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;

100 x) ~~Has~~ Technically valid and validated test methods, procedures and specifications ~~that are~~, documented in accordance
101   with the requirements of the applicable standard or guidelines, e.g. Chapter 1.1.6 *Principles and methods of validation*
102   *of diagnostic assays for infectious diseases* ~~and~~, chapters 2.2.1 to 2.2.8 *Recommendations for validation of diagnostic*
103   *tests* and Special Issue of the *Scientific and Technical Review* (2021)[27];

104 xi) ~~Demonstrates~~ Demonstrable proficiency in the applicable test methods ~~used~~ (e.g. by regular participation in
105   proficiency ~~tests on a regular basis~~ testing schemes);

106 xii) Accurate assessment and control of the measurement of uncertainty in testing;

107 xiii) Good documentation practices, e.g. ALCOA+ principles (i.e. Attributable, Legible, Contemporaneous, Original,
108   Accurate, Complete, Consistent, Enduring, Available);

109 xiv) Non-conformance management process, including detection, reporting, risk-assessment and implementation of
110   effective corrective and preventive actions;

111 xv) Complaints management;

112 xvi) Adequate control of data and information;

113 xvii) Appropriate reporting and approval process;

114 xviii) Culture of continual improvement.

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: Conformity Assessment – General Requirements for Accreditation Bodies Accrediting Conformity
121 Assessment Bodies (ISO/IEC, ~~2004b~~ 2017a). 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).

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

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

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

151 vii) Availability of personnel, facilities and equipment;

152 viii) Availability of appropriate materials and reference standards (e.g. ~~standardised~~ reagents, ~~internal quality control~~
153   ~~samples~~ controls, reference cultures) ~~and~~

154 ix)   Access to proficiency testing schemes;

155 x)   The quality ~~assurance~~ control processes necessary for materials, reagents and media;

156 xi)   The validation status, e.g. access to field samples from infected and non-infected animals, technical complexity and
157   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 ~~part~~ element of quality management focused on providing confidence that ~~quality~~ defined
161 requirements ~~will be~~ are fulfilled. The requirements may be internal or defined in an accreditation or certification standard.
162 QA is process-oriented and ~~ensures~~ provides the ~~right things are being done in the right way~~ appropriate inputs to prevent
163 problems arising.

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

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

173 All laboratories should, where possible, participate in external proficiency testing schemes appropriate to ~~their testing.~~
174 ~~Participation~~ the suite of tests provided; participation in such schemes is a requirement for accredited laboratories. This
175 provides an independent assessment of the testing methods used ~~and~~ as well as the level of staff competence. If such
176 schemes are not available, valid alternatives may be used, such as ring trials organised by reference laboratories, inter-
177 laboratory testing, use of certified reference materials or internal quality control samples, replicate testing using the same
178 or different methods, retesting of retained items, ~~and~~ or correlation of results for different characteristics of a specimen.

179 Providers and operators of proficiency testing programmes should be accredited to ISO/IEC 17043 – Conformity
180 Assessment – General Requirements for Proficiency Testing (ISO/IEC, 2010).

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

7. Test methods

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

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

196 In the veterinary profession laboratories, other standard methods (published in international, regional, or national
197 standards) or fully validated methods (having undergone a full collaborative study and that are published or issued by an
198 authoritative technical body such as the AOAC International) may be preferable to use, but may not be available. Many
199 veterinary laboratories develop or modify methods, and most laboratories have test systems that use non-standard
200 methods, or a combination of standard and non-standard methods. In veterinary laboratories, even with the use of standard
201 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, verification and validation of test methods.

205 This *Terrestrial Manual* provides more detailed and specific guidance on test selection, optimisation, standardisation, and
206 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*
207 that deal with the development and optimisation of fundamentally different assays such as antibody, antigen and nucleic
208 acid detections tests, measurement uncertainty, statistical approaches to test validation, selection and use of reference
209 samples and panels, validation of diagnostic tests for wildlife, and comparability experiments after changes in a validated
210 test method.

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

7.1. Selection of the test method

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

7.1.1. Considerations for the selection of a test method

217   i) International acceptance;

218   ii) Scientific acceptance;

219   iii) Appropriate or current technology;

220   iv) Suitable performance characteristics (e.g. analytical and diagnostic sensitivity and specificity,
221   repeatability, reproducibility, isolation rate, limits of detection, precision, trueness, and
222   uncertainty);

223   v) Suitability of the test in the species and population of interest;

224   vi) Sample type (e.g. serum, tissue, milk) and its expected quality or state on arrival at the laboratory;

225   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   xv)   Availability of reference standards, reference materials and proficiency testing schemes. (See
234   also chapter 2.2.6.).

7.2. Optimisation and standardisation of the test method

236 Once the method has been selected, it must be set up at the laboratory. Additional optimisation is necessary, whether
237 the method was developed in-house (validation) or imported from an outside source (verification). Optimisation
238 establishes critical specifications and performance standards for the test process as used in a specific laboratory.

7.2.1. Determinants of optimisation

240   i)   Critical specifications for equipment, ~~instruments~~ consumables, ~~and~~ reagents (e.g. chemicals,
241   biologicals), reference standards, reference materials, and internal controls;

242   ii)   Robustness – critical control points and acceptable ranges, attributes or behaviour at critical
243   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   viii)   Evidence of technical competence for those ~~who~~ performing the test ~~processes~~ methods,
250   authorising test results and interpreting results.

7.3. Validation of the test method

252 Test method validation evaluates the test for ~~its~~ fitness for ~~a given use~~ purpose by establishing ~~test~~ performance
253 characteristics such as sensitivity, specificity, and isolation rate; and diagnostic parameters such as positive or
254 negative cut-off, repeatability, reproducibility and titre of interest or significance. Validation should be ~~done~~ performed
255 using an optimised, documented, and fixed procedure. The extent and depth of the validation process will depend on
256 logistical and risk factors. ~~It~~ and may involve any number of activities and amount of data, with subsequent data
257 analysis using appropriate statistical methods (Chapter 1.1.6.). Acknowledging diagnostic test validation science as
258 a key element in the effective detection of infectious diseases, WOAH recently published a Special Issue representing
259 an up-to-date compilation of the relevant standards (WOAH and non-WOAH) and guidance documents for all stages
260 of diagnostic test validation and proficiency testing, including design and analysis, as well as clear, complete and
261 transparent reporting of validation studies in the peer-reviewed literature (Colling & Gardner, 2021). It is important to
262 note that the current version of ISO 17025:2017 specifies that personnel must be authorised to perform validation
263 and related activities, which means that training in validation and verification methods, including results interpretation,
264 is likely to become more important to prove competence (Colling & Gardner, 2021).

7.3.1. Activities that validation might include

266   i)   Field or epidemiological studies, including disease outbreak investigations and testing of samples
267   from infected and non-infected animals;

268   ii)   Development of testing algorithms for specific purposes, e.g. surveillance, outbreak
269   investigations, etc.;

270   iii)   Repeat testing in the same laboratory to establish the effect of variables such as operator,
271   reagents, equipment;

272   iv)   Comparison with other, preferably standard methods and with reference standards (if available);

273   v)   Collaborative studies with other laboratories using the same documented method. Ideally
274   organised by a reference laboratory and including testing a panel of samples of undisclosed
275   composition or titre with expert evaluation of results and feedback to ~~the~~ participants to estimate
276   reproducibility;

277   vi)   Reproduction of data from an accepted standard method, or from a ~~reputable~~ peer-reviewed
278   publication (verification);

279   vii)   Experimental infection ~~or disease outbreak~~ studies;

280   viii)   Analysis of internal quality control data.

281   Validation is always a balance between cost, risk, and technical possibilities. There may be cases where
282   ~~quantities such as~~ only basic accuracy and precision can ~~only~~ be ~~given~~ determined, e.g. when the disease is
283   not present in a ~~simplified way~~ country or region. Criteria and procedures for the correlation of test results for
284   diagnosis of disease status or for regulatory action must be developed. The criteria and procedures developed
285   should account for screening methods, retesting and confirmatory testing.

286   ~~Test validation is covered in chapter 1.1.6.~~

~~7.4. Uncertainty of the test method~~

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

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.

295   Laboratories must estimate the MU for each test method resulting in a quantitative measurement ~~included in their~~
296   ~~scope of accreditation~~, and for any methods used to calibrate equipment, included in their scope of accreditation
297   (ISO/IEC 17025, ~~2005~~ 2017b).

298   Tests can be broadly divided into two groups: quantitative (e.g. biochemical assays, enzyme-linked immunosorbent
299   assays [ELISA], titrations, real-time polymerase chain reaction [PCR], pathogen enumeration, etc.); and qualitative
300   (bacterial culture, parasite identification, virus isolation, endpoint PCR, immunofluorescence, etc.).

301   The determination of MU is well established in *quantitative* measurement sciences (ANSI, 1997). It may be given as
302   a numeric expression of reliability and is commonly shown as a stated range. Standard deviation (SD) and confidence
303   interval (CI) are examples of the expression of MU, for example the optical density result of an ELISA expressed as
304     $\pm n$ SD, where n is usually 1, 2 or 3. The confidence interval (usually 95%) gives an estimated range in which the
305   result is likely to fall, calculated from a given set of test data. For quantitative measurements, example for a top-down
306   or control-sample approach are provided for an antibody ELISA in chapter 2.2.4, and by the Australian government
307   webpage[28]. An example for a quantitative PCR (TaqMan) assay is provided by Newberry & Colling (2021).

308   The ISO/IEC 17025 requirement for "quality control procedures for monitoring the validity of tests" implies that the
309   laboratory must use quality control procedures that cover all major sources of uncertainty. There is no requirement
310   to cover each component separately. Laboratories may establish acceptable specifications, criteria, ranges, etc., at
311   critical control points for each component of the test process. The laboratory can then implement appropriate quality
312   control measures at these critical points, or seek to reduce or eliminate the uncertainty effect of each component.

7.4.1. Potential sources of uncertainty include:

314   i)   Sampling;

315   ii)   Contamination;

316   iii)   Sample transport and storage conditions;

317   iv)   Sample processing;

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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)	Reagent quality, preparation and storage;
319	vi)	Type of reference material;
320	vii)	Volumetric and weight manipulations;
321	viii)	Environmental conditions;
322	ix)	Equipment effects;
323	x)	Analyst or operator bias;
324	xi)	Biological variability;
325	xii)	Unknown or random effects.

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 source of uncertainty is introduced (the uncertainty of the correction). This must be assessed as part
330 of the MU estimate.

331 The application of the principles of MU to *qualitative* testing is less well defined. The determination
332 and expression of MU has not been standardised for veterinary (or medical, food, or environmental)
333 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 may preclude metrologically and statistically valid calculation of uncertainty of measurement. In such
336 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 Analytical Chemistry) teach courses or provide guidance on MU for laboratories seeking
342 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 ~~establish acceptable specifications, criteria, ranges, etc., at critical control points for each component~~
347 ~~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.~~

| 363 | ~~Systematic errors or bias determined by validation must be corrected by changes in the method,~~ |
| 364 | ~~adjusted for mathematically, or have the bias noted as part of the report statement.~~ |

365	~~If an adjustment is made to a test or procedure to reduce uncertainty or correct bias then a new~~
366	~~source of uncertainty is introduced (the uncertainty of the correction). This must be assessed as part~~
367	~~of the MU estimate.~~

368 Additional information on the analysis of uncertainty may be found in the Eurachem Guides to
369 Quantifying Uncertainty in Measurement (Eurachem, 2012) and Use of ~~uncertainty information in~~
370 ~~compliance assessment~~ Uncertainty Information in Compliance Assessment (Eurachem, 2007).

7.5. Implementation and use of the test method

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

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

8. Strategic planning

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

388 i)   Attendance at conferences, organisation of in-house or external meetings on diagnostics and quality
389   management;

390 ii) ~~Participation in~~ <u>Membership of</u> local and international organisations;

391 iii) ~~Participation in writing~~ <u>Contribution to</u> national and international standards (e.g. on ILAC and ISO committees);

392 iv) <u>Maintenance of</u> current awareness ~~of publications, writing~~ <u>through review of</u> and ~~reviewing publications about~~
393   ~~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>.

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

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29 NCSL: The National Conference of Standards Laboratories.
30 CITAC: The Cooperation of International Traceability in Analytical Chemistry.

3 **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

4 **Paris, 4–8 September 2023**

5 _____

6 # C H A P T E R   1 . 1 . 9 .

7 # TESTS FOR STERILITY AND FREEDOM FROM
8 # CONTAMINATION OF BIOLOGICAL MATERIALS
9 # INTENDED FOR VETERINARY USE

10 ## 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*
14 *derivation and substances containing such derivatives. Where chemical or physical treatments are*
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*
20 *preferred reference is master cell stocks. While the onus for ensuring safety of a product remains with the*
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*
25 *examples of contamination of vaccines with extraneous agents: a) Foamy virus (Spumaretroviridae) was*
26 *identified as a contaminant of primary monkey kidney cultures used for vaccine production in the early 1950,*
27 *b) In the 1960s it was shown that yellow fever live attenuated vaccines prepared in chicken embryo*
28 *fibroblasts were infected with avian leukosis virus (ALV). c) Calicivirus was found in Chinese hamster ovary*
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*
32 *1 and 2) are one of the most common extraneous agents in veterinary and human vaccines, g) RD114 is a*
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*
36 *(SV40). SV40 was found to cause cancer in animals and is associated with human brain, bone and lung*
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 *Section G. Protocol examples below. Control of contamination with transmissible spongiform*
43 *encephalopathy (TSE) agents is not covered in this chapter because standard testing and physical*

44   *treatments cannot be used to ensure freedom from these agents.* <u>*Detection methods are described in*</u>
45   <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 fit<u>s</u> 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 <u>various</u> culture methods, <u>including use</u>*
59   *<u>of embryonated eggs, which are</u> supported by cytopathic effects (CPE) detection<u>/embryo death</u>, fluorescent*
60   *antibody techniques and* ~~*other*~~ *suitable (<u>fit for purpose),</u> methods such as polymerase chain reaction (PCR)*
61   *and <u>antigen detection ELISA</u> (enzyme-linked immunosorbent assay). As is explained in more detail in this*
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>.*

65   ~~*Avian materials and vaccines are required to be inoculated on to primary avian cell cultures or eggs for the*~~
66   ~~*detection of avian viruses. A combination of general tests, for example to detect haemadsorbing,*~~
67   ~~*haemagglutinating and CPE-causing viruses and specific procedures aimed at the growth and detection of*~~
68   ~~*specific viruses is recommended to increase the probability of detection. Assays to detect other*~~
69   ~~*contaminants, such as bacteria, fungi, protozoa, rickettsia and mycoplasma are also described.*~~

70   ~~*Procedures applied*~~ <u>*Testing procedures*</u> *should be validated and found to be "fit for purpose" following*
71   *Chapter 1.1.6. Validation of diagnostic assays for infectious diseases of terrestrial animals, where possible.*

72   <u>*It is a requirement of many regulators, that a laboratory testing report notes the use of validated procedures*</u>
73   <u>*and describes the validated procedures in detail including acceptance criteria. This gives the regulator*</u>
74   <u>*transparency in the procedures used in a testing laboratory.*</u>

75   <u>*The validation assessment of an amplification process in cell culture should include documentation of the*</u>
76   <u>*history of permissive cell lines used, reference positive controls and culture media products used in the*</u>
77   <u>*process of excluding adventitious agents, to ensure the process is sound and is not compromised. The*</u>
78   <u>*validation assessment should give information (published or in-house) of the limitations that may affect test*</u>
79   <u>*outcomes and an assessment of performance characteristics such as analytical specificity and sensitivity of*</u>
80   <u>*each cell culture system, using well characterised, reference positive controls.*</u>

81   *It is <u>the</u> responsibility of the submitter to* ~~*assure*~~ <u>*ensure*</u> *a representative selection and number of items to*
82   *be tested.* ~~*The principles of*~~ *Appendix 1.1.2.1 Epidemiological approaches to sampling: sample size*
83   *calculations of Chapter 1.1.2 Collection, submission and storage of diagnostic specimens* ~~*apply*~~ <u>*describes*</u>
84   <u>*the principles to be applied.*</u> ~~*Adequate transportation is described in*~~ *Chapter 1.1.2 and Chapter 1.1.3*
85   *Transport of biological materials <u>describe transportation requirements</u>.*

A. AN OVERVIEW OF TESTING APPROACHES

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

91 • Primary materials must be collected from sources shown to be free from contamination and handled in such a way as
92   to minimise contamination and the opportunities for any contaminants to multiply (Figure 1).

93 • Materials that are not sterilised and those that are to be processed further after sterilisation must be handled aseptically.
94   Such materials will require further assessment of freedom of contaminants at certain stages of production to assure
95   freedom of adventitious agents.

96 • Materials that can be sterilised without their biological activities being affected unduly must be sterilised by a method
97   effective for the pathogens ~~concerned~~ of concern. The method must reduce the level of contamination to be
98   undetectable, as determined by an appropriate sterility test study. ~~(See Section D.1. below).~~ If a sterilisation process is
99   used, it shall be validated to demonstrate that it is fit for purpose. Suitable controls will be included in each sterilisation
100   process to monitor efficiency.

101 • The environment in which any aseptic handling is carried out must be maintained in a clean state, protected from
102   external sources of contamination, and controlled to prevent internal contamination. Rules governing aseptic
103   preparation of vaccines are documented in Chapter 2.3.3 *Minimum requirements for the organisation and management*
104   *of a vaccine manufacturing facility*.

105 **Figure 1. ~~Testing algorithm~~ Risk assessment flowchart for vaccine production.**

Sourcing and testing of raw materials,
e.g. porcine trypsin, serum etc.

↓

Testing bulk products for adventitious agents,
e.g. master cell seed, master virus seed, bovine serum albumin

↓

Monitoring of manufacturing processes for
freedom of contamination (GMP)

↓

Evaluation of the clearance of specific viruses by the
manufacturing process, e.g. inactivation studies and safety testing

106

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

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

123 More recently metagenomic high throughput sequencing (HTS) workflows have shown potential for quality control of
124 biological products (van Borm *et al*., 2013) and vaccines (Baylis *et al.,* 2011; Farsang & Kulcsar, 2012; Neverov &
125 Chumakov, 2010; Onions & Kolman, 2010; Victoria *et al*., 2010) in particular for the identification and characterisation of
126 unexpected highly divergent pathogen variants (Miller *et al*., 2010; Rosseel *et al*., 2011) that may remain undetected using
127 targeted diagnostic tests. Nevertheless, targeted assays, e.g. amplification in cell culture followed by polymerase chain
128 reaction (PCR) may be superior to HTS for specific agent detection (Wang *et al*., 2014) due to lack of sensitivity of HTS at
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.

134 Given the availability of new technologies, there will be future opportunities and challenges to determine presence of
135 extraneous agents in biologicals intended for veterinary use for industry and regulators. Problems can arise when the
136 presence of genome positive results are interpreted as evidence for the presence of contamination (Mackay & Kriz,
137 2010). When using molecular technologies, it is important to understand the correlation between genome detection and
138 detection of live ~~virus~~ agent. It cannot be assumed that detection of genome corresponds to the presence of an infectious
139 agent.

B. LIVING VIRAL VACCINES FOR ADMINISTRATION BY INJECTION, OR THROUGH DRINKING WATER, SPRAY, OR SKIN SCARIFICATION

142 1. Materials of animal origin ~~shall~~ should be ~~(a)~~ sterilised, ~~or (b)~~ and obtained from healthy animals that, in so far as is
143 possible, should be shown to be free from pathogens that can be transmitted from the species of origin to the species
144 to be vaccinated, or any species in contact with them by means of extraneous agents testing.

145 2. Seed lots of virus, any continuous cell line and biologicals used for virus growth ~~shall~~ should be shown to be free from
146 ~~viable~~ bacteria, fungi, mycoplasmas, protozoa, rickettsia, and extraneous viruses ~~and other pathogens~~ that can be
147 transmitted from the species of origin to the species to be vaccinated or any species in contact with them. ~~There may~~
148 ~~be some exceptions for a limited number of non-pathogenic bacteria and fungi to be present in live viral vaccines~~
149 ~~produced in eggs and administered through drinking water, spray, or skin scarification.~~

150 For ~~the~~ production of vaccines in embryonated chicken eggs and the quality control procedures for these vaccines, it
151 is recommended ~~(required in many countries)~~ that eggs from specific pathogen-free birds should be used.

152 3. Each batch of vaccine ~~shall~~ should pass tests for freedom from extraneous agents that are consistent with the
153 importing country's requirements for accepting the vaccine for use. Some examples of published methods that
154 document acceptable testing ~~procedures~~ processes in various countries include: ~~(US) Code of Federal Regulations~~
155 ~~(2015); European Pharmacopoeia (2014); European Commission (2006); World Health Organization (WHO) (1998;~~
156 ~~2012) and Department of Agriculture (of Australia) (2013).~~

157 • Code of Federal Regulations, Title 9 (9CFR) (of the United States of America) (2015).
158 • Department of Agriculture, Forest and Fisheries (Australia) (2013).
159 • Department of Agriculture, Forest and Fisheries (Australia) (2021b) Live Veterinary Vaccines.
160 • European Medicines Agency Sciences Medicines Health (2016).
161 • European Pharmacopoeia, 10th Edition (2021).
162 • World Health Organization (WHO) (1998; 2012).

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

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

175 ~~1. Section B applies.~~

176 ~~2. A limited number of contaminating, non-pathogenic bacteria and fungi may be permitted (see Section I.2.2 *General*~~
177 ~~*Procedure for testing live viral vaccines produced in eggs and administered through drinking water, spray, or skin*~~
178 ~~*scarification for the presence of bacteria and fungi*).~~

D C. INACTIVATED VIRAL AND BACTERIAL VACCINES

179

180 1. Each batch of vaccine shall pass a test for inactivation of the vaccinal ~~virus~~ seed and should include inactivation studies
181   on representative extraneous agents if the virus or bacterial seed has not already been tested and shown to be free
182   from extraneous agents. An example of a simple inactivation study could include assessment of the titre of live vaccine
183   before and after inactivation and assessing the $\log_{10}$ drop in titre during the inactivation process. This would give an
184   indication of the efficacy of the inactivation process. There is evidence that ~~virus~~ titration tests may not have sufficient
185   sensitivity to ensure complete inactivation. In these circumstances, a specific innocuity test would need to be developed
186   and validated to be fit for increased sensitivity. To increase sensitivity more than one passage would be required
187   depending on the virus or bacteria of concern. An example of this approach can be found at:
188   https://www.aphis.usda.gov/animal_health/vet_biologics/publications/memo_800_117.pdf (accessed 25 July 2023).

189 2. If studies on representative extraneous agents are required, then spiking inactivated vaccine with live representative
190   agents and following the example of an inactivation study ~~as in D.1 above would~~ could be useful. The inactivation
191   process and the tests used to detect live ~~virus~~ agent after inactivation must be validated and shown to be suitable for
192   their intended purpose.

193   In addition, each country may have ~~particular~~ its own requirements for sourcing or tests for sterility as detailed in
194   Section B above.

E. D. LIVING BACTERIAL VACCINES

195

196 1. See Section B ~~applies~~.

197 2. Seed lots of bacteria shall be shown to be free from other bacteria as well as fungi and mycoplasmas, protozoa,
198   rickettsia, and extraneous viruses. Agents required for exclusion will be dependent on the country accepting the vaccine
199   for use. Use of antibiotics to 'inactivate' the living bacterial seed or vaccine prior to exclusion of viruses and fungi is
200   recommended to ensure testing in culture is sensitive. ~~Interference testing is recommended to ensure that the~~
201   ~~antibiotics used do not affect the growth of the extraneous virus or fungi that is being excluded. Sonication may also~~
202   ~~be useful~~

203   ~~Interference testing is required to ensure that antibiotics used (or sonication) does not affect the growth of extraneous~~
204   ~~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
208   concentration of antibiotics can be determined in a dilution experiment such as documented in 9CFR Section 113.25(d).
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
211   would require ~~validation~~ verification to ensure the process does not affect the sensitivity of exclusion of extraneous
212   agents of concern.

213 3. Sonication of a living bacterial seed may be useful when excluding specific viral agents. Once again, the inactivation
214   procedure would require a verification process to ensure the adventitious virus being excluded is not affected by the
215   treatment. Use of a suitable reference virus control during the exclusion process would be required.

216 4. Direct PCR techniques may be useful when culturing processes fail to be ~~sensitive~~ successful in detecting extraneous
217   bacteria from live bacterial seeds or vaccines.

~~F. INACTIVATED BACTERIAL VACCINES~~

218

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

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

1. Section B.~~1~~ <u>applies for sera/diagnostic agents that are not inactivated. Section C</u> applies for ~~non-~~inactivated sera<u>/diagnostic agents</u>.

2. Some countries require quarantine, health certification, and tests for specific diseases to be completed for all serum <u>and plasma</u> 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). <u>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.</u>

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).~~

4. ~~Inactivated serum, Section D applies.~~

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 bacterium is used.~~

~~H.~~ <u>F.</u> 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 <u>(2015),</u> FAO and <u>Department of Agriculture Forest and Fisheries (2021a; 2021b),</u> though ~~many such~~ <u>some</u> guidelines <u>may</u> give more detail in regard~~ ~~to the food safety aspect.

~~J.~~ <u>G.</u> PROTOCOL EXAMPLES

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

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

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 be detected further<u>,</u> by sensitive and specific diagnostic tests such as FAT or PCR ~~if~~<u>as</u> required. General detection systems can include haemabsorbance and CPE by immunohistochemistry staining methods. The example procedures for ~~sterility~~ <u>detection of contamination</u> testing and general detection of ~~viable~~ <u>virus, fungi, protozoa and</u> bacteria <u>(including rickettsia and</u> mycoplasma~~, fungi and viruses~~<u>)</u> described below are derived from standards such as the 9CFR (2015), European Pharmacopoeia, ~~(2014)~~ <u>10th Edition (2021)</u>, European ~~Commission (2006), WHO~~ <u>Medicines Agency Sciences Medicines Health (2016), Department of Agriculture, Forest and Fisheries (Australia) (2013) and World Health Organization</u> (1998; 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 <u>of concern</u>.

General procedures ~~will~~<u>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~~ <u>Fisheries</u> are able to address such agents in offering sensitive testing approaches based on reputable publications. <u>A</u> ~~CVMP~~ <u>reflection paper published</u> ~~written~~ <u>by the</u> ~~European Medicines Agency Sciences Medicines Health~~

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

271 Exclusion of specific agents requires procedures that maximise sensitivity by providing ideal amplification and detection of
272 the pathogen in question. Extraneous agents, for example, Maedi Visna virus~~, bovine immunodeficiency virus,~~ (and other
273 retroviruses), *Trypanosoma evansi* and porcine respiratory coronavirus are difficult to culture even using the most sensitive
274 approaches. In these circumstances, application of molecular assays directly to the biological material ~~in question to~~
275 ~~assess,~~ assessing for the presence of nucleic acid from adventitious agents offers an alternative. ~~Refer to Table 1.~~
276 ~~Consideration must be noted as described in Section A.6 as~~, though detection of the presence of non-viable and host
277 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
279 use, for example PCV-1 in a rotavirus vaccine (WHO, 2015). BVDV is well known for its presence in many bovine
280 associated biologicals, including cell culture. More recently, non-CPE pestivirus, BVD type 3 (HoBi-like) are found in foetal
281 calf serum and cell culture. Classical Swine fever has contaminated various porcine cell lines used for African swine fever
282 and FMDV diagnosis, and thus the potential for contamination of porcine based vaccines. PEDV is linked to spray-dried
283 porcine plasma used for feed. This is not an exhaustive list of agents of concern or by any means required for exclusion
284 by every country based on risk, they are ~~just~~ examples of infectious agents that are not culturable using general culturing
285 procedures and require ~~a more~~ use of specialised culturing processes and specific detection ~~process by means of the~~
286 ~~indirect fluorescent antibody test, PCR or ELISA, where applicable~~ processes. Notably, some subtypes of an agent type
287 may be detectable by general methods, and some may require specialised testing for detection. For example, bovine
288 adenovirus subgroup 1 (serotypes 1, 2, 3 and 9) can be readily isolated using general methods (Vero cells) however bovine
289 adenovirus subgroup 2 (serotypes 4, 5, 6, 7, 8 and 10) are not readily isolated and required specialised methods for
290 isolation.

291 **Table 1.** ~~Some~~ Examples of infectious agents of veterinary importance
292 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	Rhabdoviruses (e.g. rabies virus)	Some fungi (e.g. *Histoplasma*)

2. Example of detection of bacteria and fungi ~~contamination~~

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.

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

2.1.1. Diluent A

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

2.1.2. Diluent B

306 Diluent B is for oil-adjuvanted products or materials: Add 1 ml polysorbate 80 to 1 litre Diluent A, adjust
307 the pH to 7.1 ± 0.2, dispense into containers in 100 ml quantities, and sterilise by steam.

308 If the biological being tested has antimicrobial properties, the membrane is washed three times after
309 sample application with approximately 100 ml of the appropriate diluent (A or B). The membrane is then
310 transferred ~~whole~~ to culture media, aseptically cut into equal parts and placed in media, or the media is

transferred to the membrane in the filter apparatus. If the test sample contains merthiolate as a preservative, fluid thioglycolate medium (FTM) is used and the membranes are incubated at both 30–35°C and 20–25°C. If the test sample is a killed biological without merthiolate preservative, FTM is used at 30–35°C and soybean casein digest medium (SCDM) at 20–25°C. If the sample tested is a live viral biological, SCDM is used at both incubation temperatures. It has been suggested that sulfite-polymyxin-sulfadiazine agar be used to enhance the detection of *Clostridium* spp. when the membrane filtration technique 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 903, See https://www.aphis.usda.gov/animal_health/vet_biologics/publications (accessed 4 July 2022). To 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 microorganism	Incubation	
		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

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

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).

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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
352	according to the conditions specified.

353 To test for ability of the culture media to support growth in the presence of the test material, containers
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
357 organisms appears in all inoculated media containers within 7 days. In the event that growth is
358 evident, the organism should be identified to confirm that it is the organism originally added to the
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.

361 If the material being tested renders the medium turbid so that the presence or absence of microbial
362 growth cannot be readily determined by visual examination, 14 days after the beginning of incubation
363 transfer portions (each not less than 1 ml) of the medium to fresh vessels of the same medium and
364 then incubate the original and transfer vessels for not less than 4 days.

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~~
368 ~~fungal colony per dose for veterinary vaccines. From each container sample, each of two Petri dishes are inoculated~~
369 ~~with vaccine equal to ten doses if the vaccine is recommended for poultry, or one dose if recommended for other~~
370 ~~animals. To each plate 20 ml of brain–heart infusion agar are added containing 0.007 IU (International Units) of~~
371 ~~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.~~
372 ~~Colony counts are made at the end of each incubation period. An average colony count of all the plates representing~~
373 ~~a batch should be made for each incubation condition. If the average count at either incubation condition exceeds~~
374 ~~one colony per dose in the initial test, one retest to rule out faulty technique may be conducted using double the~~
375 ~~number of unopened final containers. If the average count at either incubation condition of the final test for a batch~~
376 ~~exceeds one colony per dose, the batch of vaccine should be considered unsatisfactory.~~

377 **2.3~~2~~.** **Example of** general procedure for testing seed lots of bacteria and live bacterial
378 biologicals for purity

379 Each seed lot of bacteria or batch of live bacterial biological should be tested for purity by inoculation of SCDM, which
380 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>
381 <u>sterile technique to avoid laboratory contamination</u>, a sterile pipette or syringe and needle is used to aseptically
382 transfer the quantity of biological directly into the two types of culture medium. The minimum ratio of inoculum to
383 culture medium is 1/15. <u>Both positive and negative controls are set up as well</u>.

384 If the inoculum or growth of the bacterial vaccine renders the medium turbid so that the absence of atypical microbial
385 growth cannot be determined by visual examination, subcultures should be made from all turbid tubes on day 3
386 through until day 11. Subculturing is done by transferring 0.1–1.0 ml to differential broths and agar and incubating for
387 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
389 lot of biological may be considered satisfactory for purity. If atypical growth is found but it can be demonstrated by a
390 <u>negative</u> control that the media or technique were faulty, then the first test ~~may~~ <u>should</u> be repeated. If atypical growth
391 is found but there is no evidence invalidating the test, then a retest ~~may~~ <u>should</u> be conducted. Twice the number of
392 biological containers and test vessels of the first test are used in the retest. If no atypical growth is found in the retest,
393 the biological could be considered to be satisfactory for purity but the results from both the initial and retest should
394 be reported for assessment by the individual countries relevant regulatory agency <u>if the laboratory is sure that the</u>
395 <u>first test result was not due to in-laboratory contamination</u>. If atypical growth is found in any of the retest vessels, the
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~~ <u>should</u> be repeated.

398 **2.4~~3~~.** ~~**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*~~

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

403	Inoculate 1.0 ml of prepared master or working ~~viral~~ live agent or cell seed material (not containing antibiotics) by
404	inoculating 50 µl of the test product into each of 10 flasks containing biphasic medium. At the same time 10 plates of
405	serum dextrose agar (SDA) are inoculated with 50 µl of inoculum and spread with a sterile bent glass Pasteur pipette
406	or hockey stick. An un-inoculated serum dextrose agar plate and a biphasic flask are also set up at the same time as
407	negative controls.

408	For assessment of inhibitory substances 50 µl of previously prepared master or working viral or cell seed material
409	and 10–100 CFU of *B. abortus* are inoculated on to duplicate SDA plates. Positive controls are prepared by inoculating
410	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% CO_2 environment. Plates are incubated with the agar |
| 412 | uppermost and flasks with the agar slope vertical. Flasks are incubated with the cap loose. |

413	Plates are checked for growth of colonies at days 4 and 8 of incubation. The biphasic medium is examined every 4 to
414	7 days for 28 days. After each examination of the flasks, they are tilted so that the liquid phase runs over the solid
415	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~~4.   ~~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
423	testing must be done before bacteriostatic or bactericidal agents are added. Five samples of each batch should be
424	tested; 5 ml or one-half of the container contents, whichever is the lesser, of the sample should be used to inoculate
425	100 ml of tryptose broth and tetrathionate broth. The inoculated broths should be incubated for 18–24 hours at 35–
426	37°C. Transfers from these broths should be made on to MacConkey and *Salmonella–Shigella* agar, incubated for
427	18–24 hours, and examined. If no growth typical of *Salmonella* is noted, the agar plates should be incubated an
428	additional 18–24 hours and again examined. If colonies typical of *Salmonella* are observed, further subculture on to
429	suitable differential media should be made for positive identification. Sensitive PCR tests are available for the
430	detection of *Salmonella* spp. in cultured material. If *Salmonella* is detected, the batch is determined to be
431	unsatisfactory.

| 432 | **3.   Example of detection of *Mycoplasma* ~~contamination~~** |

| 433 | **3.1.   An example of a ~~general~~ specific procedure for ~~detection~~ exclusion of *Mycoplasma*** |
| 434 | ***mycoides* 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~~
438	~~*Acholeplasma laidlawii*, *Mycoplasma arginini*, *M. fermentans*, *M. hyorhinis*, *M. orale*, and *M. synoviae* should be~~
439	~~used. The nutritive properties of the solid medium should be such that no fewer than 100 CFU should occur with each~~
440	~~test organism when approximately 100–200 CFUs are inoculated per plate. An appropriate colour change should~~
441	~~occur in the liquid media when approximately 20–40 CFUs of each test organism are inoculated. The ability of the~~
442	~~culture media to support growth in the presence of product should be validated for each product to be tested, and for~~
443	~~each new batch or lot of culture media.~~

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~~
446	~~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~~
447	~~sample being tested. Two plates are incubated at 35–37°C aerobically (an atmosphere of air containing 5–10% CO₂~~
448	~~and adequate humidity) and two plates are incubated anaerobically (an atmosphere of nitrogen containing 5–10%~~
449	~~CO₂ and adequate humidity) for 14 days. On day 3 or day 4 after inoculation, 0.25 ml from the liquid media are~~
450	~~subcultured on to two plates of solid media. One plate is incubated aerobically and the second anaerobically at 35–~~
451	~~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~~
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~~
453	~~for 10 days except for the 14-day subculture, which is incubated for 14 days. Liquid media is observed every 2–3~~
454	~~days and, if any colour change occurs, has to be subcultured immediately.~~

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.~~

~~Further detailed procedures can be found in Veterinary Medicinal Products, VICH GL34: 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% CO_2. 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 agar. The volume of the product is inoculated so that it is not more than 10% of the volume of the medium. The liquid medium is incubated at 37°C in 5–10% CO_2 and 100 µl of broth is subcultured on to agar at days 7, 14 and 21. The agar plates are incubated at 37°C in 5–10% CO_2 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 negative controls. For assessment of inhibitory substances, inoculate 1 ml of sample to be tested into 9 ml of the liquid medium and 100 µl on to solid medium and add 10–100 CFU of *Mmm*~~SC~~ to each. Prepare positive control by inoculating 9 ml of mycoplasma broth and a mycoplasma agar plate with 10–100 CFU of *Mmm*~~SC~~. Incubate as for samples and negative controls.

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 *Mmm*~~SC~~ 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 *Mmm*~~SC~~ 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.

If evidence of microbial growth is found in the test samples the contaminating bacterium will be identified and confirmed as *Mmm*~~SC~~ by specific PCR assay.

3.2   General testing for exclusion of *Mycoplasma* sp.

General testing for exclusion of *Mycoplasma* sp. that are less fastidious may require up to 28 days in culture, using general mycoplasma media. Some mycoplasmas cannot be cultivated, in which case the live biological sample will have to be tested using an indicator cell line such as Vero cells, DNA staining, or PCR methods.

Further detailed procedures can be found in Veterinary Medicinal Products, VICH GL34: http://www.ema.europa.eu/docs/en_GB/document_library/Scientific_guideline/2013/03/WC500140352.pdf

and

USDA SAM 910: https://www.aphis.usda.gov/animal_health/vet_biologics/publications/910.pdf, (both accessed 25 July 2023).

4.   Example of detection of rickettsia and protozoa

There are no general test procedures for exclusion of rickettsia or protozoa. Procedures to exclude specific agents of concern such as *Coxiella burnetti* (Q fever), *Ehrlichia canis, Trypanosoma evansi* and *Babesia caballi* can be found for

504 example, in the Review of Published Tests to detect pathogens in veterinary vaccines Intended for Importation into
505 Australia (Department of Agriculture [of Australia] [, Forest and Fisheries (2013)]). The review is based on the reading and
506 interpretation of applicable published papers from reputable journals and are regarded as examples of sensitive methods
507 for detection of specified agents.

4.1. An **Example of a specific test** protocol based on published methods for exclusion of *Babesia caballi* and *Theileria equi*

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

5. **Example of** detection of **virus** ~~viruses~~ in biological materials

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

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 ~~of~~ on 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 is completed.

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.~~

~~5.1. An example of general testing for the exclusion of viruses from virus and cell seed stocks 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
541 with the addition of 1 ml mono-specific antiserum.~~ The serum must be shown to be free from antibodies against any
542 agents for which the test is intended to detect. Antiserum ~~must~~ should be tested for nonspecific inhibiting affects. For
543 a general test, this can be difficult to ascertain. Serum should be of sufficiently high titre to neutralise the seed virus
544 effectively with the use of an approximately equal volume or less of serum. A microplate block titration is ~~used~~ useful
545 to determine the ~~titre~~ amount of the antiserum required to neutralise ~~the MVS~~ a known amount of ~~concern. The
546 antiserum~~ CPE causing virus seed. This is ~~allowed to neutralise the MVS at 37°C for 1 hour. The MVS and antiserum
547 mixture is then inoculated on to a 75 cm² flask with appropriate cells. If the MVS is known to be high-titred or difficult
548 to neutralise, the blocking antiserum can be added to the growth medium at a final concentration~~ done in the normal
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.

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

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: cytopathology

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 red-purple, 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 culture and then detection of that agent by means of fluorescence, antigen-capture ELISA or PCR; whichever is more sensitive. Specific testing is usually required when general procedures are not adequate for effective exclusion of more fastidious, viruses Some examples are listed in Table 1.

5.2. ~~An~~ Examples of specific ~~virus~~ agent exclusion testing ~~from~~ of biologicals used in the production 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 diarrhoea virus (PEDV) in Vero cells (CCL81, ATCC) to ensure the virus can enter host cells. Just confluent monolayers ~~(100%)~~ are required; as under confluent monolayers (<90%) are more sensitive to the presence of trypsin ~~and will be destroyed well before the 7 days required for each passage in culture~~. An over confluent or aging monolayer will not be sensitive for growth of PEDV. Maintenance media (MM) formulation consists of Earle's MEM (minimal essential medium) (with 5.6 M HEPES [N-2-hydroxyethylpiperazine, N-2-ethanesulphonic acid] and glutamine) + 0.3% Tryptose phosphate broth, 0.02% yeast extract and 4 µg/ml TPCK treated trypsin. The addition of the trypsin into ~~the~~ MM should occur on the day the media is to be used.

Prior to inoculation, confluent 75 cm^2 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~~ on 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.

Add 100 µl of diluted virus on to each of two duplicate wells. Rock inoculated plates to distribute the inoculum evenly over the surface of the monolayer. Incubate at 37°C with 5% CO_2 for 2 hours then add a further 1 ml volumes/well of MM.

After 7 days, 75 cm^2 monolayers have cells disrupted using two freeze–thaw cycles at –80°C. Positive control plates are read for end-point titres, and these are compared with virus in the presence of test 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 passage. Passages are repeated until a total of four passages are completed at which point cell lysates are assessed by PCR for detection of PEDV and day 7 monolayers in 24-well plates are fixed 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 presence of serum proteins and without trypsin present, PEDV ~~is unable to grow in cell culture~~ grows poorly, or not at all. Washing off the inoculum with two MM washes is required after an extended adsorption time of up to 4 hours to ensure acceptable sensitivity.

~~J~~ 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 should follow the requirements of the importing country. Requirements for each importing country should be accessible and published online. The manufacturer or the Veterinary Authority of the exporting country should make available detailed 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 substrates and component materials, the quality assurance procedures for each step in the process, final product testing regimes, and the pharmacopoeia with which the product must conform in the country of origin. They should also make available challenge organisms, their biotypes and reference sera, and other means of appropriate product testing.

For detailed examples of a risk-based assessment of veterinary biologicals for import into a country refer to:

- 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

657 • Department of Agriculture, Forest and Fisheries of Australia (2021b). Live veterinary vaccines Summary of information
658 required for biosecurity risk assessment, Version 6 and Inactivated veterinary vaccines, Version 8.

659 • Outline of the Regulatory System of Veterinary Drugs in Japan (2015) Assurance of the Quality, Efficacy, and Safety
660 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.

666 Genetically modified organisms (GMOs) are becoming more frequent in use with changes in manufacturing technologies
667 and specialised, time-consuming procedures need to be in place. A laboratory that accepts a GMO product for testing shall
668 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.

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

676 # ~~I.~~ 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*.

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

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

2   **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # C H A P T E R   2 . 2 . 4 .

6   # MEASUREMENT UNCERTAINTY

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 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 measurement process for detection of an analyte in a diagnostic sample is not entirely reproducible and hence there is no exact value that can be associated with the measured analyte. Therefore, the result is 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 tests. The approach described here is known as "top-down" or "control sample" because it uses a weak positive control sample and expresses the MU result at the cut-off, where it most matters. It is not a question of whether the measurement is appropriate and fit for whatever use to which it may be applied. It is not an alternative to test validation but is rightly considered a component of that process (see ~~the WOAH Validation Standard,~~ chapter 1.1.6 Section B.1.1 Repeatability).

A. THE NECESSITY OF DETERMINING MU

To assure compliance with ISO/IEC 17025~~-2005~~ 2017 requirements, national accreditation bodies for diagnostic testing 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 includes tests where numeric results are calculated and then ~~are~~ expressed as a positive or negative result at a cut-off 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% confidence interval (CI), relative standard deviation (RSD = SD / mean of replicates) and coefficient of variation (CV = RSD × 100%). Examples provided below assume normal distribution of data. The concept of MU does not apply to strictly binary (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 test method in a given laboratory. During assay development, repeatability is estimated by evaluating variation in results 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 infectious diseases of terrestrial animals*, Sections A.2.5 *Robustness* and B.1.1 *Repeatability*, and Chapter 2.2.6 *Selection and use of reference samples and panels*, Section ~~3.1~~ A.4.2). Typically, the variation in replicate results is expressed as RSD or CV. The significant feature is that repeatability studies can be used to define the expected precision of the assay in the detection of a range of analyte concentrations.

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 *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 results provide a continuous monitor relative to different aspects of repeatability, e.g. intra- and inter-assay variation, intra- and inter-operator variation and intra- and inter-batch variation, which, when subjected to statistical analysis, provide an expression of the level of robustness (precision) of a test procedure. The monitoring of assay quality control parameters 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, e.g. including sample preparation such as extraction steps or dilution of serum samples for an antibody enzyme-linked immunosorbent assay (ELISA).

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.

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 the assay threshold regarding the status of the test result as positive, negative, or inconclusive (as will be described in the following example). In this context, ~~low~~ weak positive samples, like those used in repeatability studies or as the ~~low~~ weak 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 detection for the assay. In this chapter, the application of MU with respect to cut-off (threshold) values, whether recommended by test-kit manufacturers or determined in the diagnostic laboratory, is described.

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.

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[32].

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 ~~low~~ weak positive control serum, the relative standard deviation (RSD), or coefficient of variation (CV), if expressed as a percentage, provides a convenient transformation:

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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/measurement-uncertainty (accessed 22 June 2023)

$$\text{RSD} \quad (X) \quad = \quad \text{SD} \quad (X) \quad / \quad (X)$$

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

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

91   The relative standard deviation becomes:

92
$$\text{RSD}(PI_L) = \text{SD}(PI_L) / (PI_L)$$

2.2. Example

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

98   ***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

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

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%)

103   Expanded uncertainty (U) is the statistic defining the interval within which the value of the measure and is believed
104   to lie within a specified level of confidence, usually 95%. Expanding the uncertainty is done by multiplying the RSD
105   (PI_L) by a factor of 2; this allows the calculation of an approximate 95% confidence interval around the threshold value
106   (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%

2.4. Interpretation

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

3. Example of MU calculation in molecular tests

3.1. Example

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

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

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

Mean = 33.36; Std Dev (SD) = 0.43; Assay n=10

3.2. Calculating uncertainty

From the limited data set,

RSD (PI$_L$) = 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 (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 Ct value = 37), assuming normally distributed data.

U (95%CI) = 2 × RSD = 0.0255

This estimate can then be applied at the threshold level

95% CI = 37± (37 × 0.0255) = 37 ± 0.94

The 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% CI) 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). Subtraction 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).

3.3. Interpretation of the results

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

B. OTHER APPLICATIONS

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

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

Scope and limitations of the top-down approach

Methods for quantifying uncertainty (addressing MU) for tests vary. When estimating MU for quantitative, biologically based 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, 2021; Standards Council of Canada, 2021; and footnote 1). The advantage of this method is that quality control data are generated during normal test runs and can be used to estimate the precision of the assay and express it at the cut-off. The application at the cut-off depends on the performance of the test at different analyte concentrations, e.g. variation is likely to increase at higher diluted samples. The top-down approach does not identify individual contributors to measurement uncertainty but rather provides an overall estimate. Measurement uncertainty does not replace test validation; however, the validation process includes assessments of repeatability through quality control samples which facilitate calculation of MU.

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205   *
206   *   *

207   **NB:** There is a WOAH Collaborating Centre for
208   Diagnostic Test Validation Science in the Asia-Pacific Region (please consult the WOAH Web site:
209   https://www.woah.org/en/what-we-offer/expertise-network/collaborating-centres/#ui-id-3).

210   Please contact the WOAH Collaborating Centre for any further information on validation.

211   **NB:** FIRST ADOPTED IN 2014.

2   **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # C H A P T E R   2 . 2 . 6 .

6
7 # SELECTION AND USE OF REFERENCE SAMPLES AND PANELS

8   ## INTRODUCTION

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

14   *Reference samples and panels are essential from the initial proof of concept in the development laboratory*
15   *through to the maintenance and monitoring of assay performance in the diagnostic laboratory and all of the*
16   *stages in between. The critical importance of reference samples and panels cannot be over-emphasised.*
17   *The wrong choice of reference materials can lead to bias and flawed conclusions right from development*
18   *through to validation and use. Therefore, care must be exercised in selecting reference samples and*
19   *designing panels.*

20   **Fig. 1.** *Reference samples and panels grouped based on similar characteristics and composition. The*
21   *topics and alphanumeric subheadings (e.g. Proof of concept, A.2.1) refer to the relevant section in* ~~the~~
22   ~~WOAH Validation Standard,~~ *Chapter 1.1.6* Validation of diagnostic assays for infectious diseases of
23   terrestrial animals.

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 tests</u>** B.1.4.	**Provisional recognition, B.**2.6 <u>7.</u>
<u>**ASe,** B.1.3.</u>	~~Reference samples and panels~~	**Biological modifications,** B.5.2.2.
Optimisation, A.2. ~~3~~ <u>2.</u>	**Group C**	**Group E**
~~**Robustness,** A.2.5.~~ **<u>Preliminary repeatability, A.2.8</u>**	**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 <u>7</u>.	**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.	
Technical modifications, B.5.2.1.		
Reagent replacement, B.5.2.3.		

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

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

~~Reference samples may be used for multiple purposes from the initial stages of development and 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 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 development and validation process. For assays that may target multiple species, the samples should be representative of the primary species of interest. It is critical that these samples reflect both the target analyte 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.

~~It is important to emphasise that, no matter~~ Whether reference samples are selected from natural sources or prepared in the laboratory, all selection criteria ~~or~~ <u>and</u> preparation procedures, as well as testing requirements, need to be fully described and put into document control. Not only is this good quality management practice, but it will provide both an enhanced level of continuity and confidence throughout the lifespan of the assay. <u>Summaries of the data to be collected and documented for reference material can be found in Figure 2. For more detail on best practice and quality standards for the documentation of provenance of reference material refer to Watson et al. (2021).</u>

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[33] https://www.techlab.fr/Commun/UK_Def_MRC.asp

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

A. GROUP A

The question of pooling of samples to create a reference sample is often asked. If reference material is harvested from a 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 is a good alternative but it is imperative to pool from animals that are in a similar phase of infection. This is particularly important for antibody detection systems. Pooling also addresses the issue of the larger quantities of reference material to 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 reactivity. There should be an assessment following pooling to ensure that unforeseen interference is not introduced by the pooling of multiple samples, for example differing blood types or antibody composition within the independent samples 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.

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

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

~~As mentioned above, all reference samples should be well characterised. This includes documentation on both the 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, vaccination history, herd history, etc. Wherever possible, the phase of infection should be noted. This could include details related to clinical signs, antibody profiles, pathogen load or shedding, etc. Equally important, tests that are used to determine disease/infection status need to be well documented (see Section E of this chapter for further explanation). In some cases, experimental infection/exposure may be the only viable option for the production of reference material. In this case, all of the above considerations plus the experimental protocol should be detailed.~~

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

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

1. Proof of concept (~~WOAH Validation Standard,~~ Chapter 1.1.6, Section A.2.1)

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

2. Operating range (~~WOAH Validation Standard,~~ Chapter 1.1.6, Section A.2.2 3) and analytical sensitivity (~~WOAH Validation Standard,~~ Chapter 1.1.6, Section B.1.3)

2.1. ~~Analytical approaches~~ Operating range and analytical sensitivity

119 The operating range of the assay ~~is~~ defines the lower and upper analyte detection limits and the interval ~~of analyte~~
120 ~~concentrations (amounts)~~ over which the method provides suitable accuracy and precision. ~~It also defines the lower~~
121 ~~and upper detection limits the assay. To establish this range,~~ The operating range is established by serial dilution, to
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
124 representative of the typical sample ~~matrix of samples~~ type taken from animals in the population targeted by the
125 assay. This includes antibody assays where ~~a high~~ replicates of a strong positive reference serum should be diluted
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.

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.

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

3. 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>)

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 application. At least three reference samples representing negative, ~~low~~ <u>weak</u> and ~~high~~ <u>strong</u> positive may be chosen from either natural or prepared reference samples. Optimisation experiments are rather exhaustive especially when assays with multiple preparatory and testing steps are involved. It is very important that a sufficient quantity of each reference 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 continuity of optimisation evidence.

Assessment of repeatability should begin during assay development and optimisation stages~~. Repeatability~~ <u>and</u> 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~~ <u>throughout</u> to provide continuity of evidence.

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

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

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 reference materials should be standardised. National reference standards are calibrated by comparison with an international standard reagent whenever possible. In the absence of an international standard, a national reference 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 within the responsible organisation. <u>In all cases, thorough documentation of reference material should be observed as summarised in Figure 2.</u> All of the standard reagents, whether natural or prepared, must be highly characterised through extensive analysis, and preferably the methods for their characterisation, preparation, and storage have been published in peer-reviewed publications (<u>Watson *et al.*, 2021</u>). These reference standards should also be both stable and innocuous.

Reference standards, especially antibody, are usually provided in one of two formats. They may be provided as a 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 from highly positive samples as a pre-dilution in a negative matrix in order to maximise the number of aliquots available. The drawback here is that there is no accounting for any potential matrix effect in the candidate assay as there is no matrix control provided. The other approach is to provide a negative and a ~~low~~ <u>weak</u> and ~~high~~ <u>strong</u> positive set of reference standards that are of known concentrations or reactivities and are within the operating range of the standard method that was used to prepare them. <u>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 diluted.</u> This compensates for any potentially hidden matrix effect. In addition, this set of three acts as a template for the selection and/or preparation of process controls (discussed below).

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

4.2. Working standards or process controls

Working standard reagent(s), commonly known as quality or process controls, are calibrated to international, national, or in-house standard reagents. They are selected or prepared in the local matrix which is found in the population for which the assay is intended. Ideally, negative and ~~low~~ <u>weak</u> and ~~high~~ <u>strong</u> positive working standards should be selected or prepared. Concentrations and/or reactivities should be within the normal operating range of the assay. Large quantities should be prepared, aliquoted and stored for routine use in each diagnostic run of the assay. The 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 random and/or systematic variability using various control charting methods. Their daily performance will determine whether or not an assay is in control and if individual runs may be accepted. As such, these working reference samples are critically important from a quality management standpoint.

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 <u>description of experiments and</u> statistical approaches ~~to assay precision in the face of technical modifications~~ <u>that are appropriate for comparability testing (Bowden & Wang, 2021; Reising *et al.*, 2021).</u>

In general, these approaches require the use of three reference samples, a negative<u>, a weak</u> and ~~a low and high~~ <u>strong</u> positive~~. Again these samples~~ <u>to represent the entire operating range of both assays. Samples</u> may be either natural or prepared. The important point to re-iterate here is that the same reference samples that were used in the developmental 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 reference samples have been well characterised. At the very least, if new reference samples are to be used, they should be selected or prepared using the same criteria or preparation procedures established for previous materials~~. Again~~ <u>as</u> this enhances the continuity of evidence.

6. Reagent replacement (~~WOAH Validation Standard,~~ Chapter 1.1.6, 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~~ <u>Replacement</u> reference reagent should be selected or prepared using the same criteria or preparation procedures established for previous materials~~. Again~~ <u>as</u> this enhances the continuity of evidence and confidence in the assay <u>and underlines the importance of documentation of reference material data (Figure 2)</u>.

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 ~~specific~~ <u>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, 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) infected/non-vaccinated, and iv) infected/vaccinated animals. These samples may be collected under field conditions but 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 <u>(Figure 2)</u>. Alternatively, it may be necessary to produce this material in experiments like those described in Section A.2.2 of this chapter, ~~but~~ <u>including</u> a combination of experimentally vaccinated and challenged animals. ~~It~~ <u>Application of the 3 Rs (replacement, reduction and refinement) aims to avoid or minimise the number of animals used in experiments. For enzyme-linked immunosorbent assays (ELISAs), it</u> is important to avoid use of the vaccine as capture antigen in the assay (e.g. indirect <u>ELISA</u> ~~enzyme-linked immunosorbent assay [I-ELISA]~~), because carrier proteins in the vaccine may stimulate non-specific antibody responses in vaccinated animals that may be detected in ELISA ~~leading to false positives in the assay. Similarly 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 need to be increased in order to collect sufficient reference material.~~<u>, leading to false positives in the assay.</u> Depending on the DIVA test, a single experiment could be designed to assess aspects of both ASe and ASp.

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
252   organisms that are capable of eliciting cross-reacting antibody. An attempt should be made to obtain reference samples
253   from documented cases of infections and/or organisms that may be cross-reactive. Depending on the type of assay, these
254   reference materials may represent the organism itself, host-derived samples, or genomic sequences. A profile for the
255   exclusivity of the assay should be established, and expanded on a continual basis as potentially cross-reactive organisms
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.

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

266   Some test methods or procedures are solely analytical tools and are usually applied used to further characterise an analyte
267   that has been detected in a primary assay, for example assays like. Examples are the virus neutralisation tests used to
268   type an isolated virus or characterise an antibody response and subtyping of haemagglutinin genes by polymerase chain
269   reaction of avian influenza virus. Such adjunct ancillary tests must be validated for analytical performance characteristics,
270   but and differ from to routine diagnostic tests because they do not require validation for diagnostic performance
271   characteristics. The analytical accuracy of these tests is often dependant on the use of reference reagents material. These
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.

C. GROUP C

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

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
292   these materials in bulk and prepare working aliquots from time to time. However, if storage space is available, it is
293   preferable to prepare and store large numbers of aliquots at one time because bulk quantities of analyte, undergoing
294   freeze–thaw cycles to prepare a few aliquots at a time, may be subject to degradation. Because this type of reference
295   material is consumed at a fairly high rate, they will need to be replaced or replenished on a continual basis. As potential
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
299   with respect to ASe, at least five animals in each group should be considered. For smaller host species, this number may
300   need to be increased in order to collect sufficient reference material.

1. Repeatability (~~WOAH Validation Standard,~~ Chapter 1.1.6, Section B.1.1) and ~~preliminary reproducibility~~ provisional assay recognition (~~WOAH Validation Standard,~~ Chapter 1.1.6, 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.

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 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 type of panel could also be used for a limited evaluation of reproducibility to enhance provisional acceptance status for the assay. The test method is usually assessed in ~~one~~ two or more laboratories with a high level of experience and proficiency 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 Stage 3 of assay validation. ~~Consult Chapter 2.2.4 for further explanation of the topic and its application.~~

2. Reproducibility (~~WOAH Validation Standard,~~ Chapter 1.1.6, Section B.3)

Reproducibility is an important measure of the precision of an assay when used in a cross-section of laboratories located in distinct or different regions or countries using the identical assay (protocol, reagents and controls). As the number of laboratories increases, so do~~es~~ the number of variables encountered with respect to laboratory environments, equipment differences and technical expertise. ~~These~~ An overview of the factors affecting testing reproducibility is provided in Waugh & Clark (2021). Reproducibility studies are a measure of an assay's capacity to remain unaffected by substantial changes 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 of 'blind' samples containing a minimum of 20 samples, representing negative and a range of positive samples. If selected negative and/or positive samples ~~in the panel~~ are duplicated~~,~~ in the panel then it may be possible to assess both assay reproducibility and within-laboratory repeatability ~~estimates may be augmented by replicate testing of these samples when used in the reproducibility studies~~.

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

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 testing is one measure of laboratory competence derived by means of an inter-laboratory comparison; implied is that participating laboratories are using the same (or similar) test methods, reagents and controls. Results are usually expressed qualitatively, i.e. either negative or positive, to determine pass/fail criteria. However, ~~for single dilution assays,~~ where semi-quantitative results ~~provide~~ are provided, additional ~~data for assessment of~~ analysis may assess non-random error among the participating laboratories. Refer to Johnson & Cabuang (2021) for an overview of proficiency testing and ring trials.

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 above, would be adequate. Proficiency testing is not unlike a continuous form of reproducibility assessment. However, 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 precision can be estimated for both the reproducibility and repeatability data if replicates of the same reference sample are included in this 'blind' panel. Consult Chapter 2.2.4 for further explanation of the topic and its application.~~ vary but a minimum of five samples, representing negative weak and strong positives, would be adequate.

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 *Comparability of assays after changes in a validated test method*, 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

352 days from the same animal, cannot be used as representative of individual animals in populations targeted by the assay,
353 because such samples violate the rule of independence of samples required for such studies.

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

1. Standard method comparison and provisional recognition (~~WOAH Validation Standard,~~ Chapter 1.1.6, Section<u>s</u> B.2.~~6~~ <u>5 and B.2.6</u>)

359 There are situations where it is not possible or desirable to fulfil Stage 2 of the Validation Pathway because appropriate
360 samples from the target population are scarce and animals are difficult to access (such as for exotic diseases). However,
361 a small but select panel of highly characterised test samples representing the range of analyte concentration should be
362 run in parallel in the candidate assay method and ~~by~~ a WOAH standard method, as published in the WOAH *Manual~~s~~*.
363 <u>Biobanks may be a useful resource in this context, providing well-characterised samples supported with metadata to</u>
364 <u>enhance transparency and provenance of samples used in method comparisons (Watson *et al.*, 2021).</u> 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
367 animals or animal products for exotic pathogens or confirmation of clinical signs, full validation beyond a test method
368 comparison may not be feasible or warranted.

369 Experience has shown that the greatest obstacle to continuing through Stage 2 of the Validation Pathway is the number of
370 defined samples required to estimate diagnostic performance parameters with a high degree of certainty (~~WOAH Validation~~
371 ~~Standard,~~ chapter 1.1.6, Section B.2). In some cases, provisional recognition by international, national or local authorities
372 may be granted for an assay that has not been completely evaluated past analytical stages. The different rationales for
373 provisional acceptance are well explained in ~~the WOAH Validation Standard,~~ <u>chapter 1.1.6</u>. In all cases however, sound
374 evidence must exist for comparative estimates of DSp and DSe based on a small select panel of well-characterised
375 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
380 animal. Consult Chapter 2.2.5 for statistical approaches to determining methods comparability using diagnostic samples.

2. Biological modifications (~~WOAH Validation Standard,~~ Chapter 1.1.6, 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
384 analyte as targeted in the original validated assay (e.g. from serum to saliva). At the very least, all of the analytical criteria
385 of the validation pathway must be re-assessed before proceeding. If the analytical requisites are met, the remaining
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
389 statistical approaches to determining methods comparability using diagnostic samples.

E. GROUP E

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

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

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

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

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

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

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

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

F. GROUP F

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

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

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

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

FURTHER READING

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NB: There is a WOAH Collaborating Centre for Diagnostic Test Validation Science in the Asia-Pacific Region (please consult the WOAH Web site: https://www.woah.org/en/what-we-offer/expertise-network/collaborating-centres/#ui-id-3). Please contact the WOAH Collaborating Centre for any further information on validation.

NB: FIRST ADOPTED IN 2014.

MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION

Paris, 4–8 September 2023

———————————

C H A P T E R 3 . 1 . 5 .

CRIMEAN–CONGO HAEMORRHAGIC FEVER

. . .

B. DIAGNOSTIC TECHNIQUES

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

Method	Purpose					
	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	–	+++	–	+++[b]	+[c]	–
Virus isolation in cell culture	–	–	–	+[b]	–	–
Detection of immune response						
IgG ELISA	+++	+	–	++[d]	+++	–
Competitive ELISA	+++	+	–	++[d]	+++	–
IgM ELISA	–	++	–	++[e]	–	–

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 CCHFV has been demonstrated by the detection of IgM antibodies specific to CCHFV using two different ELISAs based on two different antigens.

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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   # C H A P T E R   3 . 3 . 6 .

6   # AVIAN TUBERCULOSIS

7   # SUMMARY

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

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.*

17   Mycobacterium tuberculosis, *the agent that most commonly causes human tuberculosis* (gene IS61101) *is*
18   *~~less commonly~~ rarely the cause of infection in birds, and it is often ~~as a~~ the result of transmission from pet*
19   *bird owners.*

20   *Members of* M. avium *complex:* M. a. avium (serotypes 1–3; containing gene segments IS901 and IS1245),
21   M. avium *subsp.* hominissuis (serotypes 4–6, 8–11, and 21; lacking gene segment IS901 and containing
22   segment IS1245) *and* M. intracellulare (serotypes 7, 12–20, and 22–28; lacking both IS901 and IS1245) *can*
23   *also infect an extensive range of mammals such as swine, cattle, deer, sheep, goats, horses, cats, dogs,*
24   *and exotic species. In humans, all members of the* M. avium *complex and* M. genavense *~~are capable of~~*
25   *~~inducing~~ can induce a progressive disease that is refractory to treatment, ~~mostly~~ mainly in*
26   *immunocompromised patients.*

27   *~~All manipulations involving~~ Due to the contagious nature of this group of organisms, handling ~~of~~ open live*
28   *cultures or ~~of~~ material from infected birds must only be carried out ~~with~~ after an appropriate ~~biorisk~~*
29   *~~management~~ risk assessment and the implementation of biosafety measures designed to avoid infection.*

30   *Diagnosis of avian tuberculosis in birds depends on the demonstration of ~~the above mentioned~~ a*
31   *mycobacterial species in live or dead birds or the detection of an immune response, cellular or humoral,*
32   *culture examination, or gene segments ~~IS6110, IS901 and IS1245~~ by polymerase chain reaction (PCR) in*
33   *the excretions or secretions of live birds.*

34   ***Detection of the agent:*** *Where clinical signs of avian tuberculosis are seen in the flock, or typical*
35   *tuberculous lesions are present in birds at necropsy, the demonstration of acid-fast bacilli in smears or*
36   *sections made from affected organs is sufficient for a quick positive diagnosis. If acid-fast bacilli are not*
37   *found but typical tuberculous signs or lesions are present in the birds, a culture of the organism or PCR must*
38   *be attempted. ~~PCR could also be carried out directly~~ on tissue samples. Any acid-fast organism isolated*
39   *should be identified by nucleic-acid-based tests or chromatographical (e.g. high-performance liquid*
40   *chromatography [HPLC]) criteria; serotyping of isolates of* M. avium *complex members or PCR for 16S rRNA*
41   *gene followed by sequencing, or the presence of an amplicon for the insertion sequences IS6110, IS901,*
42   *and IS1245 could also be performed.*

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

46　　　　*In domestic fowl, the tuberculin test in the wattle is the test of choice. This test is less useful in other species*
47　　　　*of bird. ~~A better test, especially in waterfowl, is~~ The whole blood stained-antigen agglutination test is better,*
48　　　　*especially in waterfowl. It is more reliable and ~~has the advantage that it will~~ can give a result within a few*
49　　　　*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]).*

A. INTRODUCTION

54

55　　Several mycobacterial species can be involved in the aetiology of avian tuberculosis ~~and~~, also known as avian
56　　mycobacteriosis. Avian tuberculosis is most commonly ~~produced~~ caused by infection with *Mycobacterium avium* subsp.
57　　*avium* (serotypes 1, 2, and 3: containing specific gene segment IS*901* and nonspecific segment IS*1245*) and less frequently
58　　by *M. genavense* (Guerrero *et al.*, 1995; Pavlik *et al.*, 2000; Salamatian *et al., 2020; Sattar et al., 2021*; Tell *et al.*, 2001).
59　　Avian mycobacteriosis is also caused by other two members of the *M. avium* complex: *M. avium* subsp. *hominissuis*
60　　(serotypes 4–6, 8–11, and 21: lacking gene segment IS*901* and containing segment IS*1245* and mainly infecting humans
61　　and pigs) and *M. intracellulare* (serotypes 7, 12–20, and 22–28: lacking both gene segments IS*901* and IS*1245*) and by
62　　~~M. intracellulare, M. scrofulaceum, M. fortuitum, and~~ other potentially pathogenic mycobacterial species including
63　　*M. scrofulaceum and M. fortuitum*. Under some circumstances, an extensive range of mammalian species, such as swine,
64　　cattle, deer, sheep, goats, horses, cats, dogs, and exotic animals, can be infected by these mycobacterium species
65　　(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).
66　　*Mycobacterium tuberculosis* and *M. bovis* are ~~less common as causal~~ rarely the causative agents of tuberculosis in birds
67　　(Hoop, 2002; Lanteri *et al., 2011*; Peters *et al., 2007; Schmidt et al., 2022;* Tell *et al.*, 2001).

68　　*Mycobacterium avium* species with standing in nomenclature as of 2023[35] (Arahal *et al.*, 2023) consists of ~~four~~ three
69　　subspecies: *M. avium* subsp. *avium*, ~~M. avium subsp. hominissuis,~~ *M. avium* subsp. *silvaticum*, and *M. avium* subsp.
70　　*paratuberculosis* (Mijs *et al.*, 2002; Thorel *et al.*, 1990). The latter is the causal agent of Johne's disease, or
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.
76　　*paratuberculosis*, and *M. avium* subsp. *lepraemurium*. Further subdividing *M. avium* subsp. *avium* into three variants *M.*
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~~
80　　~~genome and produce a characteristic three-band pattern in IS1245 restriction fragment length polymorphism (RFLP) as~~
81　　~~described and standardised previously (Dvorska et al., 2003; Ritacco et al., 1998). This repetitive sequence is also present~~
82　　~~in M. a. silvaticum and RFLP analysis can help with identification. IS901 has only been detected in M. avium strains with~~
83　　~~serotypes 1, 2 and 3 (Pavlik et al., 2000; Ritacco et al., 1998) that are apparently more pathogenic to birds than other~~
84　　~~serotypes (Tell et al., 2001). On the basis of genetic and phenotypic differences it has recently been proposed to~~
85　　~~differentiate M. a. avium into two subspecies based on the target organism: M. a. hominissuis for human and porcine~~
86　　~~isolates and M. a. avium for bird type isolates (Mijs et al., 2002). Mycobacterium a. hominissuis has polymorphic multiband~~
87　　~~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~~
88　　~~noting that the typical features of bird-isolates, the three-band pattern in IS1245 RFLP and presence of IS901, have also~~
89　　~~been found in cervine and bovine isolates of M. a. avium.~~

90　　Avian tuberculosis in birds is most prevalent in gallinaceous poultry and ~~in~~ wild birds raised in captivity. Turkeys are ~~quite~~
91　　susceptible, but ducks, geese, and other water birds are comparatively resistant. The practices of allowing poultry to roam
92　　at large on the farm (free range) and of keeping the breeders for several years are conducive to the spread of the causal
93　　agent of avian tuberculosis among them. Infected individuals and contaminated environments (water and soil) are the ~~main~~
94　　primary sources of infection. The above-mentioned mycobacterial species causing avian tuberculosis can survive for
95　　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 <u>usually</u> 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~~<u>, including</u> sudden
99   death may occur, dyspnoea is less common, and granulomatous ocular lesions (Pocknell *et al*., 1996) ~~as well as~~ <u>and</u> skin
100 lesions have been reported. Under intensive husbandry conditions, sudden death may occur, often associated with severe
101 lesions in the liver; such lesions are easily observed at post-mortem examination (<u>Salamatian *et al.,* 2020;</u> Tell *et al*., 2001).

102 The primary lesions of avian tuberculosis in ~~birds~~ <u>poultry (chickens and turkeys)</u> are nearly always in the intestinal tract.
103 Such lesions take the form of deep ulcers filled with caseous material containing many mycobacterial cells, and these are
104 discharged into the lumen and appear in the faeces. Before the intestinal tract is opened, the ulcerated areas appear as
105 tumour-like masses attached to the gut wall~~, but~~<u>. Still,</u> when the intestine is opened, the true nature of the mass becomes
106 evident. Typical caseous lesions are nearly always found in the liver and spleen~~, and~~<u>;</u> these organs <u>are</u> usually ~~are~~ greatly
107 enlarged because of the formation of new tuberculous tissue. The lungs and other tissues are ordinarily free from lesions
108 even in advanced cases (<u>Salamatian *et al.,* 2020;</u> 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~~
111 ~~tuberculous lesions are found in the head and mesenteric lymph nodes on meat inspection after slaughter. Findings of~~
112 ~~tuberculous lesions that involve other organs (liver, spleen, lungs, etc.) are rare, usually occurring at the advanced stage~~
113 ~~of the disease. *Mycobacterium a. avium* accounted for up to 35% of the *Mycobacteria* isolated from such tuberculous~~
114 ~~lesions (Dvorska *et al.,* 1999; Pavlik *et al.,* 2003, 2005; Shitaye *et al.,* 2006). Unlike the other species mentioned previously,~~
115 ~~cattle are highly resistant to the causative agent of avian tuberculosis, and tuberculous lesions are detected in head lymph~~
116 ~~nodes, or occasionally in liver lymph nodes, only on meat inspection. *Mycobacterium a. avium* can be successfully isolated~~
117 ~~from tuberculous lesions in mesenteric lymph nodes from juvenile cattle: the isolation rate from cattle under 2 years of age~~
118 ~~was 34.4% in contrast to 13.0% from cattle over 2 years of age (Dvorska *et al.,* 2004).~~

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

131 It is essential to bear in mind that all members of *M. avium* ~~complex~~ and *M. genavense* are capable of giving rise to a
132 progressive disease in humans that is refractory to treatment, especially in immunocompromised individuals (<u>Narsana *et*</u>
133 <u>*al.,* 2023;</u> Pavlik *et al*., 2000; Tell *et al*., 2001). ~~Members of *Mycobacterium avium* complex are classed in Risk Group 2 for~~
134 ~~human infection and should be handled with appropriate measures~~ <u>All *Mycobacterium* species can cause infection in</u>
135 <u>people (Cowman *et al.,* 2019). Caution should be exercised by those working with birds in environments infected with</u>
136 <u>*Mycobacterium*, especially those immunosuppressed. All laboratory manipulations with live cultures or potentially</u>
137 <u>infected/contaminated material must be performed at an appropriate biosafety and containment level determined by</u>
138 <u>conducting a thorough risk assessment</u> as described in Chapter 1.1.4 *Biosafety and biosecurity: Standard for managing*
139 *biological risk in the veterinary laboratory and animal facilities*. ~~Biocontainment measures should be determined by risk~~
140 ~~analysis as described in Chapter 1.1.4.~~ <u>The CDC's online Manual for Biosafety in Microbiological and Biomedical</u>
141 <u>Laboratories is also a good reference</u>[36].

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36 https://www.cdc.gov/labs/pdf/SF__19_308133-A_BMBL6_00-BOOK-WEB-final-3.pdf

B. DIAGNOSTIC TECHNIQUES

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

Method	Purpose					
	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)	±	+++	±	–	++	–
PCR	++±	=+	++−	+++	+−	–
Detection of immune response						
Haemagglutination (stained antigen)	±	+++	±	=	++	=
Tuberculin test	++	+++	+	–	++	–

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.

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 detection of acid-fast bacilli (AFB) in smears or sections from affected organs, stained by the Ziehl–Neelsen method usually 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 due to large infecting doses giving rise to acute overwhelming disease, in which affected organs, most obviously the liver, have a 'morocco leather' appearance with fine greyish or yellowish mottling. In such cases AFB may not be found in such 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 to penetration of the stain. Recently, DNA probes and, polymerase chain reaction (PCR), and WGS techniques have been 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 genavense* is particularly fastidious and has special unique requirements for growth and identification (Shitaye *et al.,* 2010).

1.1. Culture

If there is a characteristic flock history and suggestive lesions are found at necropsy, but no AFB are seen in smears or sections, an attempt must be made to isolate the causative organism from the necropsy material. The liver or spleen is usually the best organ to use, but if the carcass is decomposed, bone marrow may prove more satisfactory as it could be less contaminated. As with the culture of *M. bovis*, non-sterile specimens need to be processed with 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 such as Lowenstein–Jensen, Herrold's medium, Middlebrook 7H10 and, 7H11, or Coletsos, with 1% sodium pyruvate added. It may occasionally be is necessary to incorporate mycobactin J, as it is used for the isolation of to isolate *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 'smooth' colonies within 2–4 weeks; rough variants do occur. Shorter incubation times can be achieved using the liquid culture BACTEC system or the automated fluorescent MGIT 960 culture system. *Mycobacterium a. avium* can also be detected in massively infected tissue by a conventional PCR, which also allows acceleration of the accelerates

175	pathogen <u>detection and</u> identification (Moravkova *et al.,* 2008). ~~Currently,~~ Direct detection and quantification of
176	*M. a. avium* using IS*901* quantitative real-time PCR can be considered ~~as~~ the ~~best~~ <u>fast and inexpensive</u> method
177	~~(despite its rather high cost per test)~~ (Kaevska *et al.,* 2010; Slana *et al.,* 2010).

178 For *M. genavense*, the optimal <u>medium is liquid media supplemented with Mycobactin J (an iron chelator) and then</u>
179 <u>plated onto a</u> solid medium ~~is~~ <u>such as</u> Middlebrook 7H11 ~~medium acidified to pH 6 and supplemented with blood and~~
180 ~~charcoal~~ (Realini *et al.*, 1999). The incubation period at 37°C <u>with 5–7% CO_2</u> should be extended for at least ~~6 months~~
181 <u>42 days. If samples are directly plated onto solid media, plates should be held for at least 12 weeks. Bacterial growth</u>
182 <u>should be prepared in a smear and stained using an acid-fast stain. All acid-fast organisms should be identified using</u>
183 <u>MALDI-TOF (matrix assisted laser desorption ionisation–time of flight [mass spectrometry]) or PCR (Buckwalter *et*</u>
184 <u>*al.,* 2016; Hall *et al.,* 2003; Shitaye *et al.,* 2010).</u>

185 Typing of mycobacteria to the species and subspecies level requires a specialised laboratory. Conventional
186 biochemical tests for species identification are lengthy and fail to distinguish between *M. avium* and *M. intracellulare*.
187 Thus, a miscellaneous group of mycobacteria that includes both species is usually classified under the ~~denomination~~
188 ~~of~~ *M. avium* complex <u>denomination</u>. Seroagglutination, ~~which is~~ based on <u>the</u> sugar residue specificity of surface
189 glycopeptidolipids, allows ~~classification~~ <u>the parsing</u> of *M. avium* complex organisms into 28 serovars (Wolinsky &
190 Schaefer, 1973). More sophisticated typing methods directed at cell-wall-specific targets are currently available, such
191 as enzyme-linked immunosorbent assays with monoclonal antibodies to major serovars, ~~and~~ high-performance liquid
192 chromatography ~~(HPLC)~~<u>,</u> <u>and WGS. Based on DNA–rRNA hybridisation</u> serovars 1 to 6, 8 to 11, and 21 ~~are currently~~
193 <u>have been</u> ascribed to *M. a. avium* and *M. a. hominissuis*, and serovars 7, 12 to 20, and 25 to *M. intracellulare*.
194 However, no consensus was achieved on other serovars, and some isolates cannot be <u>serotyped</u> (Inderlied *et al*.,
195 1993). <u>For final species and subspecies identification, the current methods are WGS and bioinformatic analysis of</u>
196 <u>isolates obtained from sick birds.</u> Avian tuberculosis in birds is <u>commonly</u> caused by *M. a. avium* types 1, 2, or 3. If
197 the isolate is not one of these three serotypes, further <u>molecular</u> identification tests ~~(IS*901* PCR)~~ must be ~~carried out~~
198 <u>conducted</u> in a specialised laboratory. However, it should be ~~borne in mind~~ <u>noted</u> that ~~superficial~~ tuberculous lesions
199 in ~~caged~~ <u>pet</u> birds, especially psittacines, may be caused by *M. tuberculosis,* and ~~IS*6110* PCR should be used for~~
200 precise identification <u>should always be attempted</u> (Hoop, 2002; Lanteri *et al.,* 2011; Peters *et al.,* 2007; Schmidt *et*
201 *al.,* 2008; Tell *et al.,* 2001).

1.2. Nucleic acid recognition methods

203 Specific and reliable genetic tests for speciation ~~are currently~~ <u>have been</u> available (Saito *et al*., 1990)~~.~~<u>, including</u>
204 commercial nucleic ~~acid~~ hybridisation probes ~~have become a 'gold standard'~~ <u>reference method</u> for ~~distinction between~~
205 <u>distinguishing</u> *M. avium*<u>,</u> ~~and~~ *M. intracellulare* ~~cultures.~~ <u>and *M. genavense*</u> ~~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
207 have been described that fail to react with specific *M. avium* and *M. intracellular* probes (Soini *et al*., 1996).
208 Nevertheless, identification errors were reported due to the cross-reactivity, which may have serious consequences
209 (van Ingen *et al*., 2009). Various in-house molecular methods have been reported ~~for the identification of~~ <u>to identify</u>
210 mycobacterial cultures, including ~~MAC.~~ <u>members</u> of the *Mycobacterium avium* complex. <u>The following gene segments</u>
211 <u>could be used to identify *Mycobacterium* isolates as *M. avium* in one multiplex PCR reaction: IS*900*, IS*901*, IS*1245*.</u>
212 <u>The isolates of *M. a. avium/M. a. silvaticum* are IS900−, IS901+, IS1245+, the isolates of *M. a. hominissuis* are</u>
213 <u>IS900−, IS901−, IS1245+, and the isolates of *M. a. paratuberculosis* are IS900+, IS901−, IS1245− (Kaevska *et al.,*</u>
214 <u>2010; Moravkova *et al.,* 2008).</u> A ~~multiplex~~ <u>16S rRNA</u> PCR <u>and sequencing</u> method for differentiating *M. avium* from
215 *M. intracellulare* and *M. tuberculosis* complex ~~has some advantages (Cousins *et al*., 1996).~~ <u>16S rRNA</u> ~~is currently~~
216 <u>commercially available. Similarly, many veterinary diagnostic laboratories commonly perform in-house PCR and</u>
217 <u>sequencing</u> (Kirschner *et al*., 1993) ~~may also be used~~. Culture-independent in-house molecular tests have been
218 developed ~~for the detection~~ <u>to detect</u> and ~~identification of~~ <u>identify</u> species belonging to the *M. avium* complex directly
219 from samples (Hall *et al.,* 2003; Kaevska *et al.,* 2010). <u>WGS of isolates has recently become the go-to molecular</u>
220 <u>method to identify mycobacterium isolates from birds with great accuracy. It enables, with the use of bioinformatic</u>
221 <u>tools, not only an accurate identification of species and subspecies, but also helps to determine the organism</u>
222 <u>relatedness within a flock or environment (Witte *et al.,* 2021). In recent years, veterinary diagnostic laboratories have</u>
223 <u>extensively adopted real-time PCR methods to detect *M. a. avium* directly from different specimens (faeces, tissues,</u>
224 <u>formalin-fixed tissues, and environmental samples). The technique rapidly detects fastidious and slow-growing</u>
225 <u>microorganisms, such as *M. a. avium* (Tell *et al.,* 2003a; 2003b).</u>

226 <u>Several commercial diagnostic PCR tests for detecting *M. a. avium* are available. Still, users should consider the skill</u>
227 <u>set and equipment necessary to perform such tests. Furthermore, it is important to determine the fitness for the</u>
228 <u>purpose of these tests before implementation. The interpretation of the results of these molecular tests also requires</u>
229 <u>veterinary expertise.</u>

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

233	~~1992) is possessed not only by the isolates of the above-named serotypes, but also by isolates, virulent for birds, that~~
234	~~could not be typed because agglutination occurred (Pavlik *et al.,* 2000). In epidemiological studies, a standardised~~
235	~~IS*901* RFLP methods replaced serotyping (Dvorska *et al.,* 2003).~~

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.

2.1. Tuberculin test

The tuberculin test is <u>the</u> most widely used test ~~in~~ <u>for</u> 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 intradermal inoculation in the wattle with 0.05 ml or 0.1 ml of tuberculin (containing approximately 2000 International Units [IU]), using a ~~very~~ fine needle of approximately <u>26 gauge,</u> 10 mm <u>long</u> ~~× 0.5 mm~~. The test is read after 48 hours ~~and~~<u>.</u> <u>A</u> positive reaction is any swelling at the site, from a small firm nodule approximately 5 mm in diameter to gross oedema extending into the other wattle and down the neck. ~~With practice,~~ <u>Even</u> very small wattles on immature birds can be inoculated successfully. However, ~~in immature birds~~ the comb may be used <u>in immature birds</u>, although <u>the</u> results are not ~~so~~ <u>as</u> reliable. Tuberculin testing of the wattle in turkeys is much less ~~reliable~~ <u>consistent</u> than in ~~the~~ domestic ~~fowl~~ <u>chickens</u>. Inoculation in the wing web has been recommended as ~~being~~ more efficient, but this is still not as good as ~~for domestic fowl~~ <u>in chickens</u>. Other birds may also be tested in the wing web, but results are not generally satisfactory. The bare ornamental skin areas on Muscovy ducks and some ~~species of~~ pheasant <u>species</u> can be used, but ~~reliability~~ <u>dependability</u> is doubtful, and interpretation <u>is</u> 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.

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.

2.2. Stained antigen test

<u>The stained-antigen agglutination test has been used with good results, especially in domestic and ornamental waterfowl. A drop (0.05–0.1 ml) of the antigen is mixed with the same volume of fresh whole blood, obtained by venipuncture, on a white porcelain or enamel tile. The mixture is rocked for 2 minutes and examined for agglutination. 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 blood-red colour. This test is especially useful for screening large flocks for immediate culling and therefore has 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.</u>

2.2.1. Preparation of the antigen

An antigen stained with 1% malachite green is used for the rapid whole blood plate agglutination test (Rozanska, 1965). The strain used to prepare the stained antigen must be smooth and not auto-agglutinate in saline suspension. It must conform to the characteristics of *M. a. avium*, preferably obtained from a culture collection, to guarantee its authenticity.

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 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 serotype 2 antigen will always detect serotype 3 infection and vice versa. Serotype 1 strains detect a wide spectrum of infections and frequently detect infections with mycobactin-dependent mycobacteria or *M. a. silvaticum*. There is no reason not to use a culture containing more than one 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.

The organism should be grown in a suitable liquid medium, such as Middlebrook 7H9 containing 1% sodium pyruvate. Good growth should be obtained in approximately 7 days. The liquid culture is used as a seed for bulk antigen preparation.

Antigen for agglutination tests is best obtained on a solid medium, such as Löwenstein–Jensen or 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
290 wash over the whole surface and provide sufficient moisture to keep the air in the bottle near 100%
291 humidity.

292 The bottles are incubated at 37°C, and good growth should be obtained in 14–21 days with most
293 strains. The antigen is harvested by adding sterile glass beads and twice the volume of sterile normal
294 saline (containing 0.3% formalin) as was used to inoculate the bottle. The bottle is then shaken gently
295 to wash off all the growth, and the washing is collected into a sterile bottle and re-incubated at 37°C
296 for 7 days. The killed bacilli are washed twice in sterile normal saline with 0.2% formalin by
297 centrifugation and re-suspension. This sequence is safer than the original method in which the
298 washing was carried out before the incubation that kills the organisms. Finally, bacilli are again
299 centrifuged and re-suspended in sterile normal saline containing 0.2% formalin and 0.4% sodium
300 citrate to a concentration of about 10^{10} bacteria per ml. This corresponds to a concentration ten times
301 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
303 suspension made from the culture washings is also re-examined microscopically (for likely
304 contaminants such as yeasts) and rechecked by culture to ensure that the formalin has killed the
305 mycobacteria.

2.2.2. Validation of the antigen

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

309 One or more batches for agglutinating antigen must be tested for efficacy in using serum from
310 naturally or artificially infected tuberculous birds by comparison with a standard preparation of known
311 potency. When using animals for research or reagent testing, approval of the procedures and the use
312 of animals by the institution's ethics committee should be sought before any testing occurs. The
313 potency relative to that of the standard preparation must not differ significantly from that declared on
314 the label. Each bottle of antigen must be tested with normal chicken serum (to detect
315 autoagglutination) and *M. a. avium* positive chicken serum of low and high antibody content. This
316 should be done, where possible, alongside a previous batch of stained antigens. Those bottles that
317 give satisfactory agglutination reactions with the antisera can now be pooled and the antigen stained.
318 This is done by adding 3 ml of 1% malachite green solution per 100 ml of suspension. The stained
319 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,
322 especially for autoagglutination.

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

Note on limitation of use

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

C. REQUIREMENTS FOR DIAGNOSTIC BIOLOGICALS

1. Background

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
332 *M. a. avium*. It is used by intradermal injection to reveal delayed hypersensitivity ~~as a means of identifying~~ to identify birds
333 infected with or sensitised to the same species of ~~tubercle bacillus~~ *Mycobacterium*. Importantly it is also used ~~as an~~ to aid

334 to differential diagnosis in the comparative intradermal tuberculin test for bovine tuberculosis (see Chapter 3.1.13). An
335 international standard preparation of PPD-A is being developed by WOAH to replace the former WHO Standard[37].

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

2. Outline of production and minimum requirements for tuberculin production

2.1. Characteristics of the seed

2.1.1. Biological characteristics of the master seed

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

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

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

2.2. Method of manufacture

2.2.1. Procedure

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

357 The organism is cultivated in modified Dorset-Henley's synthetic medium, then killed by heating in
358 flowing steam and filtered to remove cells. The protein in the filtrate is precipitated chemically
359 (ammonium sulphate or trichloroacetic acid [TCA] are used), washed, and resuspended. An
360 antimicrobial preservative that does not give rise to false-positive reactions, such as phenol (not more
361 than 0.5% [w/v]), may be added. Mercurial derivatives should not be used. Glycerol (not more than
362 10% [w/v]) or glucose (2.2% [w/v]) may be added as a stabiliser. The product is dispensed aseptically
363 into sterile neutral glass containers, which are then sealed to prevent contamination. The product
364 may be freeze-dried.

2.2.2. Requirements for ingredients

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

2.2.3. In-process controls

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

2.2.4. Final product batch tests

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37 PPD of *M. avium* tuberculin, WHO (1955) Technical Report Series, no.96, 11.
38 https://www.edqm.eu/en/d/234640?p_l_back_url=%2Fen%2Fsearch%3Fq%3Dpurified%2Bprotein%2Bderivative

i) Sterility

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 biological materials intended for veterinary use*).

ii) Identity

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* using a procedure similar to that described in Section C.2.2.4.iv. ~~In guinea pigs sensitised with *M. bovis*,~~ The potency of the preparation of avian tuberculin must be shown to be not more than 10% of the potency of the standard preparation of bovine tuberculin used in the potency test. The use of animals for this purpose should be reviewed and approved by your institution's ethical committee.

iii) Safety

Tuberculin PPD-A can be examined for freedom from living mycobacteria using the culture method described previously. This culture method, which does not require the 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 the safety of PPD. The use of animals for this purpose should be reviewed and approved by the institution's ethics committee. Two guinea-pigs, each weighing not less than 250 g and ~~that have~~ not ~~been treated~~ previously treated 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.

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~~ conducted according to the ~~specifications of the~~ European Pharmacopoeia (~~2000~~ 2024) specifications or the equivalent regulatory documents for each country or region.

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~~ International units – one IU is equal to the biological activity 0.02 μg of PPD – 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 PPD. 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 an 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.

433 At 24 hours, the <u>reactions'</u> diameters ~~of the reactions~~ are measured, and the results are calculated
434 using standard statistical methods, taking the diameters to be directly proportional to the logarithms
435 of the concentrations of the tuberculin~~s~~. The estimated potency must be not less than 75% and not
436 more than 133% of the potency stated on the label. The test is not valid unless the fiducial limits of
437 error ($p = 0.95$) are not less than 50% and not more than 200% of the estimated potency. If the batch
438 fails a potency test, the test may be repeated one or more times, provided that the final estimate of
439 potency and ~~of~~ fiducial limits is based on the combined results of all the tests.

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.

3. Requirements for authorisation/registration/licensing

3.1. Manufacturing process

444 The manufacturing process should follow the requirements of European Pharmacopoeia (~~2000~~ <u>2024</u>) or other
445 international standards.

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 concentration~~s~~ 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.

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.

3.3. Stability

456 During storage, liquid avian tuberculin should be protected from the light and held at a temperature of 5°C (±3°C).
457 Freeze-dried preparations may be stored at higher temperatures (~~but~~ not exceeding 25°C) <u>and</u> 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~~ <u>Following accepted practice, tuberculin should be</u> 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~~ <u>after</u>
461 the last satisfactory potency test: Liquid PPD tuberculin~~s~~: 2 years; lyophilised PPD<u>-A</u> tuberculin~~s~~: 8 years; HCSM
462 (heat-concentrated synthetic-medium) tuberculin~~s~~ <u>diluted</u>: 2 years. <u>Recent research on the temperature stability of</u>
463 <u>human, bovine, and avian tuberculin solutions has shown that they are stable for a year at 37°C. This should be</u>
464 <u>further explored as these products are used in the field in remote areas of the world where maintaining temperature</u>
465 <u>control is very difficult (Maes et al., 2011).</u>

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610 *
611 * *

612 **NB:** There is currently (2024) no WOAH Reference Laboratory for avian tuberculosis
613 (please consult the WOAH Web site for the current list:
614 https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3http://www.oie.int/).

615 **NB:** FIRST ADOPTED IN 1989 AS TUBERCULOSIS IN BIRDS. MOST RECENT UPDATES ADOPTED IN 2014.

2   **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # S E C T I O N   3 . 4 .

6   **B O V I N A E**

7   # C H A P T E R   3 . 4 . 1 .

8   # B O V I N E   A N A P L A S M O S I S

9   ## SUMMARY

10   ***Definition of the disease:*** *Bovine anaplasmosis results from infection with* Anaplasma marginale*. A second*
11   *species,* A. centrale*, has long been recognised and usually causes benign infections.* Anaplasma marginale
12   *is responsible for almost all outbreaks of clinical disease.* Anaplasma phagocytophilum *and* A. bovis*, which*
13   *infect cattle,* ~~have been recently~~ <u>are also</u> *included within the genus* ~~but they are not reported to~~<u>.</u> <u>Anaplasma</u>
14   <u>phagocytophilum</u> *can* cause ~~clinical~~ <u>self-limiting</u> *disease* <u>in cattle. There are no reports of disease associated</u>
15   <u>with</u> <u>A. bovis</u> <u>infection</u>*. The organism is classified in the genus* Anaplasma *belonging to the family*
16   Anaplasmataceae *of the order* Rickettsiales*.*

17   ***Description of the disease:*** *Anaemia, jaundice* <u>in acute, severe cases</u> *and* ~~sudden~~ <u>unexpected</u> *death are*
18   *characteristic signs of* <u>bovine</u> *anaplasmosis. Other signs include rapid loss of milk production and weight,*
19   *but the clinical disease can only be confirmed by identifying the organism. Once infected, cattle may remain*
20   *carriers for life, and identification of these animals depends on the detection of specific antibodies using*
21   *serological tests, or of rickettsial DNA using molecular amplification techniques. The disease is typically*
22   *transmitted by tick vectors, but mechanical transmission by biting insects or by needle can occur.*

23   ***<u>Detection</u>*** ~~***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 <u>spp.</u> *are available in some countries.*
30   Anaplasma phagocytophilum <u>*can only be observed in infected granulocytes, mainly neutrophils*</u> *and* A. bovis
31   *can only be observed* <u>in infected monocytes</u> ~~infecting granulocytes, mainly neutrophils~~*.*

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

35 *from blood retained in peripheral vessels. The latter are particularly ~~desirable~~ <u>useful</u> if post-mortem*
36 *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~~ <u>(I-ELISA)</u> is a reliable test used in many laboratories and*
44 *can be prepared in-house for routine diagnosis of anaplasmosis. <u>Finally, a displacement double-antigen</u>*
45 *<u>sandwich ELISA has been developed to differentiate between</u>* A. marginale *and* A. centrale *<u>antibodies.</u>*

46 ***Nucleic-acid-based tests*** *~~have been used~~ <u>are often used in diagnostic laboratories and</u> experimentally,*
47 *and are capable of detecting the presence of low-level infection in carrier cattle and tick vectors. A nested*
48 *<u>conventional polymerase chain reaction (PCR)</u> ~~reaction is necessary~~ <u>has been used</u> to identify low-level*
49 *carriers ~~using conventional polymerase chain reaction (PCR)~~<u>,</u> ~~and~~ <u>although</u> nonspecific amplification can*
50 *occur. ~~Recently,~~ Real-time PCR assays ~~with~~ <u>have</u> analytical sensitivity equivalent to nested conventional*
51 *PCR ~~have been described~~ <u>and are preferable in a diagnostic setting to reduce the risk of amplicon</u>*
52 *<u>contamination.</u>*

53 ***Requirements for vaccines:*** *Live vaccines are used in several countries to protect cattle against*
54 *~~A. marginale infection~~ <u>bovine anaplasmosis</u>. 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</u>* A. centrale *<u>leads to</u>*
56 *<u>infection and long-term persistence in many cattle. Vaccinated cattle are typically protected from disease</u>*
57 *<u>caused by</u>* A. marginale*<u>, but not infection.</u>*

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.*

62 Anaplasma centrale *vaccine is not entirely safe. A practical recommendation is to restrict its use, as far as*
63 *possible, to calves, as nonspecific immunity will minimise the risk of some vaccine reactions that may require*
64 *treatment with tetracycline or imidocarb. Partial immunity develops in 6–8 weeks and lasts for several years*
65 *after a single vaccination. In countries where* A. centrale *is exotic, it cannot be used as a vaccine against* A.
66 marginale.

A. INTRODUCTION

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

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

---

85 detected by these techniques and may lead to a misdiagnosis of bovine anaplasmosis. Visualisation of *A. marginale* bodies
86 in erythrocytes is therefore required for confirmation.

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

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

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

112 Intrastadial or transstadial transmission ~~is the usual mode~~ can occur, even in the one-host, *Rhipicephalus* species. Male
113 ticks may be particularly important as vectors, as they ~~can become persistently infected and serve as a reservoir~~ are most
114 likely to move between cattle searching for ~~infection~~ female ticks. Experimental demonstration of vector competence does
115 not necessarily imply a role in transmission in the field. However, *Rhipicephalus* species are clearly important vectors of
116 anaplasmosis in ~~countries such as~~ Australia ~~and countries in~~, many regions of Africa, and Latin America, ~~and some species~~
117 ~~of~~, *Dermacentor* spp. are efficient vectors in the United States of America (USA).

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.

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

B. DIAGNOSTIC TECHNIQUES

133

134

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

Method	Purpose					
	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)
Microscopic examination	–	+̶ =	–	+++	–	–
Detection of the agent[(a)]						
PCR	–	++ +̶	–	+++	–	–
Detection of immune response						
CAT[(b)]	–	–	–	–	+	+
C-ELISA[(b)]	+++	+̶ ++	+++	–	+++	+++
IFAT[(b)]	+	–	–	–	++	++
C̶F̶T̶	–	–	–	–	+̶	–
ddasELISA	=	=	=	=	=	++

135 Key: +++ = recommended for this purpose; ++ recommended but has limitations;
136 + = suitable in very limited circumstances; – = not appropriate for this purpose.
137 Agent id. = agent identification; CAT = card agglutination test; C̶F̶T̶ ̶=̶ ̶c̶o̶m̶p̶l̶e̶m̶e̶n̶t̶ ̶f̶i̶x̶a̶t̶i̶o̶n̶ ̶t̶e̶s̶t̶;̶
138 C-ELISA = competitive enzyme-linked immunosorbent assay; ddasELISA = displacement double-antigen, sandwich ELISA;
139 IFAT = indirect fluorescent antibody test; PCR = polymerase chain reaction.
140 [(a)]A combination of agent identification methods applied on the same clinical sample is recommended.
141 [(b)]These tests do not distinguish infected from vaccinated animals.

1. Detection of the agent

142

1.1. Microscopic examination

143

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

150 In contrast to *Babesia bovis, A. marginale* d̶o̶e̶s̶ infected erythrocytes do not accumulate in capillaries, so blood drawn
151 from the jugular or other large vessel is satisfactory. *Anaplasma marginale* replicate in the erythrocytes to form small
152 membrane-bound colonies, also termed inclusion bodies or initial bodies. B̶e̶c̶a̶u̶s̶e̶ ̶o̶f̶ ̶t̶h̶e̶ ̶r̶a̶t̶h̶e̶r̶ ̶i̶n̶d̶i̶s̶t̶i̶n̶c̶t̶i̶v̶e̶
153 m̶o̶r̶p̶h̶o̶l̶o̶g̶y̶ ̶o̶f̶ ̶*A̶n̶a̶p̶l̶a̶s̶m̶a̶* These initial bodies can be visualised on a blood smear, but are small and easily confused
154 with debris or stain precipitate (see Figure 1). Thus it is essential that smears are well prepared a̶n̶d̶ , including
155 ensuring slides are free f̶r̶o̶m̶ ̶f̶o̶r̶e̶i̶g̶n̶ ̶m̶a̶t̶t̶e̶r̶,̶ ̶a̶s̶ ̶s̶p̶e̶c̶k̶s̶ of debris c̶a̶n̶ ̶c̶o̶n̶f̶u̶s̶e̶ ̶d̶i̶a̶g̶n̶o̶s̶i̶s̶ and stain is recently filtered
156 (Watman #1 filter paper). Thick blood films as are used sometimes for the diagnosis of babesiosis are not appropriate
157 for the diagnosis of anaplasmosis, as *A̶n̶a̶p̶l̶a̶s̶m̶a̶ A. marginale* are difficult to identify once they become dissociated
158 from erythrocytes.

159
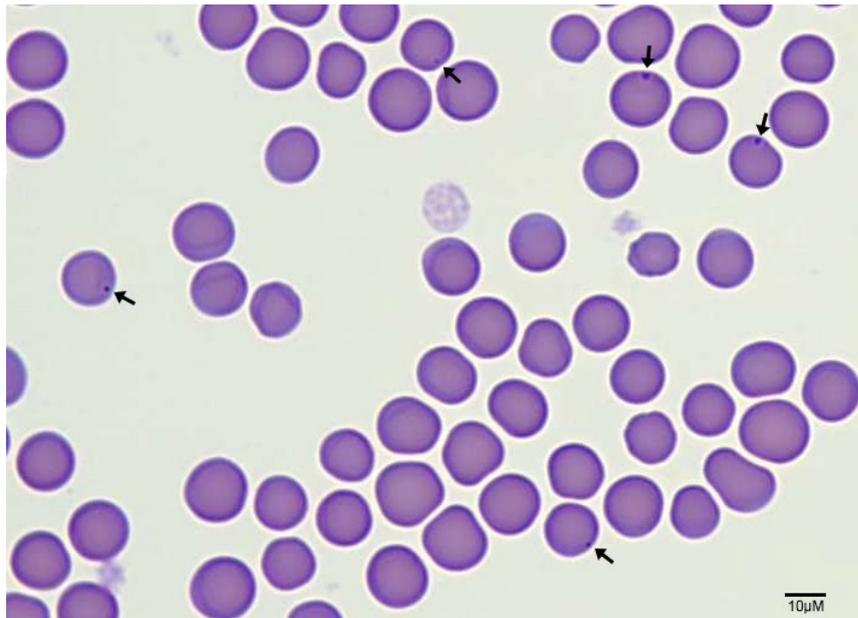
Figure. 1. _Anaplasma marginale initial bodies. A Diff-Quick stained blood smear from a bovine experimentally infected with A. marginale. Arrows point to the A. marginale initial bodies. Photo from S. Noh._

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 post-mortem examination because, under these circumstances, bacterial contamination of organ smears often makes identification of ~~Anaplasma~~ _A. 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.

Blood from organs, rather than organ tissues _per se_, is required for smear preparation, as the aim is to be able to examine microscopically intact erythrocytes for the presence of ~~Anaplasma~~ _A. marginale colonies_. Organ-derived blood smears can be stored satisfactorily at room temperature for several days.

Both blood and organ smears can be stained in 10% Giemsa stain for approximately 30 minutes after fixation in absolute methanol for 1 minute. After staining, the smears are rinsed three or four times with tap water to remove excess stain and are then air-dried. Conditions for Giemsa staining vary from laboratory to laboratory, but distilled water is not recommended for dilution of Giemsa stock. Water should be pH 7.2–7.4 to attain best resolution with Giemsa stain. Commercial stains that give very rapid staining of ~~Anaplasma~~ _A. marginale_ are available in some countries. Smears ~~are~~ must be examined under oil immersion at a magnification of ×700–1000.

Anaplasma marginale ~~appear as dense,~~ initial bodies are round~~ed 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)~~.

The percentage of infected erythrocytes varies with the stage and severity of the disease. Maximum rickettsaemias in excess of 50% may occur with _A. marginale._ Multiple infections of individual erythrocytes are common during 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.

1.2. Polymerase chain reaction

Nucleic-acid-based tests to detect *A. marginale* ~~infection~~ in ~~carrier~~ infected cattle have been developed although not ~~yet~~ 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 has been used to identify *A. marginale* carrier cattle with a capability of identifying as few as 30 infected erythrocytes per ml of blood, well below the lowest levels in carriers. However, nested PCR is time consuming as it requires two full PCR reactions, and poses significant quality control ~~and specificity~~ problems for routine use (Torioni De Echaide *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 *et al.*, 2007; Decaro *et al.*, 2008~~; Reinbold et al., 2010b~~). ~~Two~~ Advantages of ~~this technique~~ the real-time PCR, which uses a single closed tube for amplification and analysis, are reduced opportunity for amplicon contamination and a semi-quantitative assay result. Equipment and reagents needed for real-time PCR ~~is~~ are expensive~~, requires preventive maintenance,~~ and may be beyond the capabilities of some laboratories. ~~Real-time PCR assays may target one of several genes (Carelli et al., 2007; Decaro et al., 2008), or 16S rRNA (Reinbold et al., 2010b), and are reported to achieve a level of analytical sensitivity equivalent to nested conventional PCR (Carelli et al., 2007; Decaro et al., 2008; Reinbold et al., 2010b).~~

The most widely cited assays for the detection *A. marginale* in individual animals use a probe for increased specificity and are designed to detect *msp1b* (Carelli *et al.*, 2007) or *msp5* (Futse *et al.*, 2003) in genomic DNA extracted from whole blood. The assay based on detection of *msp1b* has been partially validated to detect the pathogen in individual animals and was used to define samples for the validation of a C-ELISA (Carelli *et al.*, 2007; Chung *et al.*, 2014). The analytical test performance of this assay is robust, and exclusivity testing confirmed other bacterial and protozoal tick-borne pathogens of cattle were not detected. The assay, evaluated using 51 blood samples from 18 cattle herds in three 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.

Msp5 has also been used as a target to detect *A. marginale* in cattle in field samples and more frequently in experimental samples (Futse *et al.*, 2003). *Msp5* is highly conserved among *A. marginale* strains and is a single copy gene, thus providing some advantages as a target for ensuring detection of widely variant strains of *A. marginale*. However, the related *Anaplasma* spp. and *Ehrlichia* spp. all have *msp5* orthologs with 50% identity to an *E. ruminantium* gene (NCBI accession: L07385.1), thus specificity must be determined in laboratory and field samples. Additionally, little work has been done to validate an *msp5*-based real-time PCR test for diagnostic purposes.

A third primer–probe set is designed to detect *A. marginale* using real-time, reverse transcriptase PCR. The primers amplify a 16sRNA gene segment from *A. marginale* and *A. phagocytophilum*, while the probe differentiates between the two species (Reinbold *et al.*, 2010b). The analytical performance of this assay is robust. However, the diagnostic sensitivity, specificity, and of particular importance with 16sRNA sequence-based tests, exclusivity for other tick-borne pathogens of cattle have not been evaluated. Additionally, this assay is designed for use following RNA extraction and reverse transcription, which is more laborious and expensive than DNA extraction. Bacterial RNA is rapidly degraded, and this may ultimately reduce diagnostic sensitivity of this assay.

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.* can also be used in a duplex reaction to detect and differentiate between *A. centrale* and *A. marginale* (Decaro *et al.*, 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 probes perform similarly in the single and duplex PCR (Carelli *et al.*, 2007). Using the same 51 field samples from cattle in Italy, the *A. centrale* assay had less analytical sensitivity compared with nested PCR and discordance in 4 of 51 samples between an *A. centrale* reverse line blot test and the duplex PCR assay.

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

Assay	**Reference**	**Oligonucleotides[(a)]**	**Sequence 5'–3'[(b)]**	**Amplicon size (bp)**	**NCBI accession number**
Real-time PCR	Carelli *et al.,* 2007	*Am_msp1b*_F	TTG-GCA-AGG-CAG-CAG-CTT	95	M59845
		*Am_msp1b*_R	TTC-CGC-GAG-CAT-GTG-CAT		
		*Am_msp1b*_PB	TCG-GTC-TAA-CAT-CTC-CAG-GCT-TTC-AT		
Real-time PCR	Futse *et al.,* 2003	*Am_msp5*_F	GCC-AAG-TGA-TGG-TGA-TAT-CGA	151	M93392
		*Am_msp5*_R	AGA-ATT-AAG-CAT-GTG-ACC-GCT-G		
		*Am_msp5*_PB	AAC-GTT-CAT-GTA-CCT-CAT-CAA		
Reverse-transcription real-time PCR	Reinbold *et al.,* 2010	*16S rRNA*_F[(c)]	CTC-AGA-ACG-AAC-GCT-GG	142	M60313
		*16S rRNA*_R[(c)]	CAT-TTC-TAG-TGG-CTA-TCC-C		
		*Am_16S rRNA*_PB[(d)]	CGC-AGC-TTG-CTG-CGT-GTA-TGG-T		
Real-time PCR[(d)]	Decaro *et al.,* 2008	*Ac_groEL*_F[(e, f)]	CTA-TAC-ACG-CTT-GCA-TCT-C	77	CP001759.1
		*Ac_groEL*_R[(e, f)]	CGC-TTT-ATG-ATG-TTG-ATG-C		
		*Ac_groEL*_PB[(e, f)]	TCA-TCA-TTC-TTC-CCC-TTT-ACC-TCG-T		

[(a)]*Am* denotes *A. marginale, Ac* denotes *A. centrale*, Pb denotes probe sequence.
[(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.
[(f)]Primers and probe amplify *A. centrale groEL*.

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 marginale* infections usually persist for the life of the animal. However, except for occasional small recrudescences, ~~Anaplasma~~ *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 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 assays to detect known infections of long-standing duration has been inadequately addressed.~~ An exception is a C-ELISA (see below), which ~~has been~~ was initially validated using true positive and negative animals defined by nested PCR (Torioni De Echaide *et al.,* 1998)~~, and the card agglutination assay, for which relative sensitivity and specificity in comparison with the C-ELISA has been evaluated (Molloy *et al.,* 1999)~~. And updated in 2014 (Chung *et al.,* 2014). Therefore, while most of the tests described in this section are useful for obtaining broad-based epidemiological data, caution is advised on their use for disease certification. The C-ELISA, I-ELISA and CAT are described in detail below.

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).

2.1. Competitive enzyme-linked immunosorbent assay

~~A C-ELISA using a recombinant antigen termed~~ Major surface protein 5 (MSP5) is an immunodominant protein expressed by A. *marginale*, A. *ovis*, and A. *centrale*. In *A. marginale* the gene is highly conserved making it a useful target 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 an MSP5-specific monoclonal antibody (mAb) has proven very sensitive and specific for detection of *Anaplasma*-infected animals (~~Hofmann-Lehmann *et al.*, 2004~~ Molloy *et al.*, 1999; Reinbold *et al.*, 2010b; Strik *et al.*, 2007). ~~All~~ ~~*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~~ ~~recent report~~ mAb used in the C-ELISA. This C-ELISA was updated in 2014 to improve performance by using glutathione S-transferase (GST) instead of maltose binding protein (MBP) as the tag on the rMSP5 (Chung *et al.*, 2014). This assay no longer requires adsorption to remove the antibodies directed against MBP, thus it is faster and easier than the previous version of the C-ELISA. The diagnostic sensitivity is 100% and the diagnostic specificity is 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 farms with no clinical history of bovine anaplasmosis. The 135 positive sera were from cattle positive for *A. marginale* using nested PCR and serology.

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 used in the assay did not react with *A. phagocytophilum* MSP5 in direct binding assays (Strik *et al.*, 2007). Cross 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~~ ~~negative sera from a non-endemic region, detecting acutely infected cattle as early as 16 days after experimental tick~~ ~~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~~ ~~were defined as true positive or negative using a nested PCR procedure, the rMSP5 C-ELISA had a sensitivity of~~ ~~96% and a specificity of 95% (Torioni De Echaide *et al.*, 1998)~~ *A. marginale* and *Ehrlichia* sp. BOV2010 isolated in 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 is available commercially that contains specific instructions. Users should follow the manufacturer's instructions. ~~In general, however, it is conducted~~ ~~as follows.~~

2.1.1. Kit reagents

~~A 96-well microtitre plate coated with rMSP5 antigen,~~

~~A 96-well coated adsorption/transfer plate for serum adsorption to reduce background binding,~~

~~100× Mab-peroxidase conjugate,~~

~~10× wash solution and ready-to-use conjugate-diluting buffer,~~

~~Ready-to-use substrate and stop solutions,~~

~~Positive and negative controls~~

2.1.2. Test procedure

~~i) Add 70 µl of undiluted serum sample to the coated adsorption/transfer plate and incubate at room temperature for 30 minutes.~~

~~ii) Transfer 50 µl per well of the adsorbed undiluted serum to the rMSP5-coated plate and incubate at room temperature for 60 minutes.~~

~~iii) Discard the serum and wash the plate twice using diluted wash solution.~~

~~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.~~

~~v) Discard the 1× diluted MAb-peroxidase conjugate and wash the plate four times using diluted wash solution.~~

~~vi) Add 50 µl per well of the substrate solution, cover the plate with foil, and incubate for 20 minutes at room temperature.~~

~~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.~~

~~viii) Immediately read the plate in the plate reader at 620, 630 or 650 nm.~~

2.1.3. Test validation

The mean ~~average~~ optical density (OD) of the negative control must range from 0.40 to 2.10. The ~~average per cent inhibition of the positive control must be ≥30%.~~

2.1.4. Interpretation of the results

~~The % inhibition is calculated as follows:~~

$$100 - \frac{\text{Sample OD} \times 100}{\text{Mean negative control OD}} = \text{Per cent inhibition}$$

~~% inhibition = 100[1 – (Sample OD ÷ Negative Control OD)]~~

~~Samples with <30% inhibition are negative. Samples with ≥30% inhibition are positive.~~

~~Specificity of the MSP5 C-ELISA may be increased by using a higher percentage inhibition cut-off value (Bradway *et al.*, 2001); however the effect of this change on sensitivity has not been thoroughly evaluated.~~

~~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 C-ELISA was faster, simpler, had a higher specificity and an improved resolution compared with the rMSP5-MBP C-ELISA with MBP adsorption (Chung *et al.*, 2014).~~

2.2. Indirect enzyme-linked immunosorbent assay

An I-ELISA was first developed using the CAT antigen, which is a crude *A. marginale* lysate (see below). ~~and it~~ The test can be implemented where the commercial C-ELISA is not available. Unlike the C-ELISA, most reagents, such as buffers and ready-to dissolve substrates, are available commercially in many countries. Any laboratory can prepare the antigen using local strains of *A. marginale*, though standardised methods have not been developed. I-ELISA uses small amounts of serum and antigen that ~~and the sensitivity and specificity of the test standardised with true positive and negative sera is as good as for the C-ELISA. As it~~ can be prepared in each laboratory. ~~Only the general procedure is described here (Barry *et al.*, 1986). For commercial kits, the manufacturer's instructions should be followed. In the case of in-house I-ELISA~~ The sensitivity and specificity of the test was 87.3% and 98.4–99.6% respectively, though this varied by laboratory (Nielsen *et al.*, 1996). For general methods, refer to Barry *et al.* (1986). ~~Initial bodies and membranes are obtained as for the complement fixation test (Rogers *et al.*, 1964). This antigen is treated with 0.1% sodium dodecyl sulphate for 30 minutes prior to fixing the antigen to the microtitre plate.~~ For each laboratory, the specific amount of antigen ~~has to~~ must be ~~adjusted~~ optimised to obtain the best reading and the least expenditure.

Alternatively, rMSP5 can be used as the antigen in this test. This eliminates the need for preparation and standardisation of antigen derived from splenectomised, *A. marginale* infected animals (Silva *et al.*, 2006). In a comparison between I-ELISA using the CAT antigen and rMSP with a histidine tag (rMSP5-HIS), these two I-ELISAs performed identically. In this comparison, IFAT was used as the gold standard test (Silva *et al.*, 2006).

Test results using the I-ELISA are available in about 4 to 5 hours. It is generally conducted as follows:

2.2.1. Test reagents

A 96-well microtitre plate coated with ~~crude~~ *A. marginale* antigen,

PBS/Tween buffer, (PBS 0.1 M, pH 7.2, Tween 20 0.05%),

Blocking reagent (e.g. commercial dried skim milk)

Tris buffer 0.1 M, MgCl$_2$, 0.1 M, NaCl, 005 M, pH 9.8

Substrate *p*-Nitrophenyl phosphate disodium hexahydrate

Positive and negative controls.

2.2.2. Test procedure (this test is run in triplicate)

i) Plates can be prepared ahead of time and kept under airtight conditions at –20°C.

ii) 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.

iii) Remove the lid and deposit 200 μl PBST20 solution in each well and incubate for 5 minutes at room temperature (RT).

iv) For one plate, dissolve 1.1 g of skim milk (blocking agent) in 22 ml of PBST20.

v) Remove the plate contents and deposit in each well 200 μl of blocking solution, put the lid on and incubate for 60 minutes at 37°C.

vi) Wash the plate three times for 5 minutes with PBST20.

vii) Dilute all serum samples including controls 1/100 in PBST20 solution.

viii) 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.

ix) Incubate plate at 37°C covered for 60 minutes.

x) Wash three times as described in subsection vi.

xi) Dilute 1/1000 anti-IgG bovine alkaline phosphatase conjugate in PBST20 solution. Add 200 μl of the diluted conjugate per well. Incubate the covered plate at 37°C for 60 minutes.

xii) Remove the lid and <u>wash three times as described in point vi above</u> ~~make three washes with PBST20~~.

xiii) Remove the contents of the plate and deposit 195 μl of 0.075% *p*-Nitrophenyl phosphate disodium hexahydrate in Tris buffer <u>in each well</u> and incubate for 60 minutes at 37°C.

xiv) The reaction is quantified by a microplate reader spectrophotometer, adjusted to 405 nm wavelength. The data are expressed in optical density (OD).

2.2.3. Data analysis

Analysis of results should take into account the following parameters.

i) The mean value of the blank wells.

ii) The mean value of the positive wells with their respective standard deviations.

iii) The mean value of negative wells with their respective standard deviations.

iv) The mean value of the blank wells is subtracted from the mean of all the other samples if not automatically subtracted by the ELISA reader.

v) 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.

Positive values are those above the cut-off calculated value which is the sum of the average of the negative and two times the standard deviation.

~~For purposes of assessing the consistency of the test operator, the error "E" must alsoo be estimated; this is calculated by determining the percentage represented by the standard deviation of any against their mean serum.~~

<u>As with all diagnostic tests, it is important to measure reproducibility. For more details see Chapter 2.2.4 *Measurement uncertainty.*</u>

2.3. <u>Displacement double-antigen sandwich ELISA to differentiate between *A. marginale* and *A. centrale* antibodies</u>

<u>In regions where vaccination with *A. centrale* is used to control bovine anaplasmosis, differentiation between *A. centrale*-vaccinated and *A. marginale*-infected animals may be useful. Because there is often high amino acid identity between *A. marginale* and *A. centrale* surface proteins, identifying unique targets for serological assays for this purpose is difficult. Epitopes from MSP5 (aa28-210, without the transmembrane region) that are not shared between *A. marginale* and *A. centrale* were used to develop a displacement double-antigen sandwich ELISA</u>

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
411 and allowed to incubate. Following washing, a combination of biotinylated and non-biotinylated recombinant proteins
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*
414 MSP5-coated well (ODAm) and the OD for the *A. centrale* MSP5 (ODAc) coated well for each animal is measured. If
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*.

418 For the detection of *A. marginale* the test has a diagnostic specificity of 98% and a diagnostic sensitivity of 98.9%.
419 For 702 field samples evaluated, 131 (19%) had an OD <0.2 in the ddasELISA and thus were excluded from the
420 analysis. Of those animals, 52% were nested PCR positive for *A. marginale*, 23% were nested PCR positive for *A.*
421 *centrale*, 4.6% were nested PCR positive for *A. marginale* and *A. centrale*, 20% were nested PCR negative for both,
422 suggesting the ddasELISA may lack sensitivity.

423 Of the 571 ddasELISA positive field samples, the agreement between the ddasELISA and nested PCR was 84% and
424 the kappa coefficient was 0.70 (95% CI: 0.635–0.754), indicating substantial agreement between tests. There was
425 agreement between the ddasELISA and nested PCR for 93% of the *A. marginale* ddasELISA positive samples and
426 86% of the *A. centrale* ddasELISA positive samples. Additionally, 36 nested PCR negative samples tested positive
427 for antibodies against *A. marginale* (*n*=28) or *A. centrale* (*n*=8) by ddasELISA. This test could not identify animals
428 with co-infections, meaning animals vaccinated with *A. centrale* that are then infected with *A. marginale*, which is not
429 uncommon.

430 Test results using the ddasELISA are available in 5–6 hours. It is conducted as outlined below, see Bellezze *et al.*,
431 2023 for more details.

2.3.1. Test reagents

433   i) A 96-well microtitre plate coated with either *A. marginale* or A. centrale recombinant protein

434   ii) PBS/Tween buffer (PBS (50mM sodium phosphate, 150 mM NaCL, pH 7.2) with 0.05% Tween-
435   20)

436   iii) Blocking reagent (PBS with 10% commercial dried skim milk)

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   vii) Chromogenic substrate (1 mM 2,2'-Azinobis [3-ethylbenzothiazoline-6-sulfonic acid]-
441   diammonium salt in0.05 M sodium citrate, pH 4.5, 0.0025% V/V H_2O_2 (100 µl/well).

442   viii) ELISA plate reader (405 nm reading)

443   ix) Positive and negative control sera for *A. marginale* and *A. centrale*

2.3.2. Test procedure

445   i) Plates are coated overnight.

446   ii) Block with blocking buffer for 1 hour at room temperature and wash three times with PBS/Tween
447   buffer.

448   iii) Add undiluted serum 100 ul/well and incubate for 1 hour at 25°C at 100 rpm.

449   iv) Wash three times with PBS/Tween buffer.

450   v) Add 100 µl of *A. marginale* MSP5-biotin (1 µg/ml) plus *A. centrale* MSP5 (10 µg/ml) to
451   *A. marginale* test wells. Add *A. centrale* MSP5-biotin (1 µg/ml) plus *A. marginale* MSP5
452   (10 µg/ml) in PBS/Tween buffer + 10% fat-free dried milk to *A. centrale* test wells.

453   vi) Incubate 1 hour at 25°C, 100 rpm and wash the plate five times with PBS/Tween buffer.

454   vii) To detect the bound protein–biotin complex, add streptavidin-HRP diluted in 1/500 in
455   PBS/Tween buffer with 10% dried milk for 1 hour at 25°C, 100 rpm.

456   viii) Wash five times with PBS/Tween buffer.

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

2.4. Card agglutination test

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

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:

2.4.1. Test procedure

479		i)	Ensure all test components are at a temperature of 25–26°C before use (this constant
480			temperature is critical for the test).
481		ii)	On each circle of the test card (a clear perspex/plastic or glass plate marked with circles that
482			are 18 mm in diameter), place next to each other, but not touching, 10 µl of bovine serum factor
483			(BSF), 10 µl of test serum, and 5 µl of CAT antigen[39]. Negative and low positive control sera
484			must be tested on each card.
485		iii)	BSF is serum from a selected animal with high known conglutinin level. If the conglutinin level
486			is unknown, fresh serum from a healthy animal known to be free from *Anaplasma* can be used.
487			The BSF must be stored at –70°C in small aliquots, a fresh aliquot being used each time the
488			tests are performed. The inclusion of BSF improves the sensitivity of the test.
489		iv)	Mix well with a glass stirrer. After mixing each test, wipe the stirrer with clean tissue to prevent
490			cross-contamination.
491		v)	Place the test card in a humid chamber and rock at 100–110 rpm for 7 minutes.
492		vi)	Read immediately against a backlight. Characteristic clumping of the antigen (graded from +1
493			to +3) is considered to be a positive result. The test is considered to give a negative result when
494			there is no characteristic clumping.

495 A latex card agglutination test, a relatively simple and rapid test platform, has been partially validated.
496 This test uses rMSP5-HIS rather than *A. marginale* lysate and does not require BSF. The
497 performance of this test was compared with that of the I-ELISA using rMSP5-HIS as the antigen. The
498 relative sensitivity was 95.2% and relative specificity was 91.86% (Ramos *et al*., 2014).

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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).

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 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 smears. A serious problem encountered with the test is nonspecific fluorescence. The reported sensitivity is 97.6% and specificity 89.6% (Gonzalez et al., 1978). Antigen made from blood collected as soon as adequate rickettsaemia (5–10%) occurs is most likely to be suitable. Nonspecific fluorescence due to antibodies adhering to infected erythrocytes can be reduced by washing the erythrocytes in an acidic glycine buffer before antigen smears are prepared. Infected erythrocytes are washed twice in 0.1 M glycine buffer (pH 3.0, centrifuged at 1000 *g* for 15 minutes 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 with other members of the *Anaplasmataceae* family, and specifically an *Ehrlichia* spp. identified as BOV2010 (Al-Adhami et al., 2011).

2.6.   Complement fixation test

The complement fixation test (CFT) was used extensively for many years; however, it has variable sensitivity (ranging from 20 to 60%), possibly reflecting differences in techniques for antigen production, and poor reproducibility. In addition, the CF assay fails to detect a significant proportion of carrier cattle (Bradway et al., 2001). It is also uncertain as 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 detecting infected animals.

C. REQUIREMENTS FOR VACCINES

1.   Background

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

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

2. Outline of production and minimum requirements for conventional vaccines

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.

Anaplasma centrale usually causes benign infections, especially if used in calves under 9 months of age. Severe reactions following vaccination have been reported when adult cattle are inoculated. The suitability of an isolate of *A. centrale* as a vaccine can be determined by inoculating susceptible cattle, monitoring the subsequent reactions, and then challenging the animals and susceptible controls with a virulent local strain of *A. marginale.* Both safety and efficacy can be judged by monitoring rickettsaemias in stained blood films and the depression of packed cell volumes of inoculated cattle during 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 ~~contaminants~~ pathogens 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, PCR, 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, and 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).

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
597	anticoagulant (5 International Units [IU] heparin/ml blood). The use of blood collection units for human
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 *et al.,* 2004).

| 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 *et al.,* 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: NaCl (7.00 g), $MgCl_2.6H_2O$
616	(0.34 g), glucose (1.00 g), Na_2HPO_4 (2.52 g), KH_2PO_4 (0.90 g), and $NaHCO_3$ (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. |

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 *et al.,* 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.

625　　iv)　Chilled vaccine should be kept refrigerated 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.

| 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 *et al.,* 2004). |

2.2.2.　　Requirements for substrates and media

636

637	*Anaplasma centrale* ~~cannot~~ <u>can</u> be cultured in ~~vitro~~ <u>*Rhipicephalus appendiculatus* and *Dermacentor variabilis*</u>
638	<u>cells lines, though antigen expression and immunogenicity of the cultured *A. centrale* need to be tested (Bell-</u>
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.

2.2.3.　In-process controls

641

642　　i)　Source and maintenance of vaccine donors

643	A source of calves free from natural infections of ~~*Anaplasma*~~ <u>*A. marginale*</u> 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	The calves should be maintained under conditions that will prevent exposure to infectious diseases
647	and to ticks and biting insects. In the absence of suitable facilities, the risk of contamination with the
648	agents of infectious diseases present in the country involved should be estimated, and the benefits
649	of local production of vaccine weighed against the possible adverse consequences of spreading
650	disease (Bock *et al.*, 2004).

651   ii)   Surgery

| 652 | Donor calves should be splenectomised to allow maximum yield of organisms for production of |
| 653 | vaccine. This is best carried out in young calves and under general anaesthesia. |

654   iii)   Screening of vaccine donors before inoculation

655	As for preparation of seed stabilate, donor calves for vaccine production should be examined for all
656	blood-borne infections prevalent in the vaccine-producing country, including *Babesia*, *Anaplasma*,
657	*Ehrlichia*, *Theileria* and *Trypanosoma*. This can be done by routine examination of stained blood films
658	after splenectomy, and preferably also by serology. Any calves showing evidence of natural infections
659	of any of these agents should be rejected. The absence of other infective agents should also be
660	confirmed. These may include the agents of enzootic bovine leukosis, bovine viral diarrhoea,
661	infectious bovine rhinotracheitis, ephemeral fever, Akabane disease, bluetongue, and foot and mouth
662	disease. The testing procedures will depend on the diseases prevalent in the country and the
663	availability of tests, but should involve serology of paired sera at the very least and, in some cases,
664	virus isolation, antigen, or DNA/RNA detection (Bock *et al.*, 2004; Pipano, 1981; 1997).

665   iv)   Monitoring of rickettsaemias following inoculation

666	It is necessary to determine the concentration of rickettsia in blood being collected for vaccine. The
667	rickettsial concentration can be estimated from the erythrocyte count and the rickettsaemia
668	(percentage of infected erythrocytes).

669   v)   Collection of blood for vaccine

670	All equipment should be sterilised before use (e.g. by autoclaving). Once the required rickettsaemia
671	is reached, the blood is collected in heparin using strict aseptic techniques. This is best done if the
672	calf is sedated and with the use of a closed-circuit collection system.

673	Up to 3 litres of heavily infected blood can be collected from a 6-month-old calf. If the calf is to live,
674	the transfusion of a similar amount of blood from a suitable donor is indicated. Alternatively, the calf
675	should be killed immediately after collection of the blood.

676   vi)   Dispensing of vaccine

677	All procedures are performed in a suitable environment, such as a laminar flow cabinet, using
678	standard sterile techniques. Use of a mechanical or magnetic stirrer will ensure thorough mixing of
679	blood and diluent throughout the dispensing process. Penicillin (500,000 IU/litre) and streptomycin
680	(370,000 µg/litre) are added to the vaccine at the time of dispensing.

681   2.2.4.   Final product batch tests

682	The potency, safety and sterility of vaccine batches cannot be determined in the case of chilled vaccine, and
683	specifications for frozen vaccine depend on the country involved. The following are the specifications for frozen
684	vaccine produced in Australia.

685   i)   Sterility and purity

| 686 | Standard tests for sterility are employed for each batch of vaccine and diluent (see Chapter 1.1.9 |
| 687 | *Tests for sterility and freedom from contamination of biological materials intended for veterinary use*). |

688	The absence of contaminants is determined by doing appropriate serological testing of donor cattle,
689	by inoculating donor lymphocytes into sheep and then monitoring them for evidence of viral infection,
690	and by inoculating cattle and monitoring them serologically for infectious agents that could potentially
691	contaminate the vaccine. Cattle inoculated during the test for potency (see Section C.2.2.4.iii) are
692	suitable for the purpose. Depending on the country of origin of the vaccine, these agents include the
693	causative organisms of enzootic bovine leukosis, infectious bovine rhinotracheitis, bovine viral
694	diarrhoea, ephemeral fever, Akabane disease, Aino virus, bluetongue, parainfluenza, foot and mouth
695	disease, lumpy skin disease, rabies, Rift Valley fever, contagious bovine pleuropneumonia,
696	Jembrana disease, heartwater, pathogenic *Theileria* and *Trypanosoma spp.*, *Brucella abortus*,

697 *Coxiella*, and *Leptospira* (Bock *et al.*, 2004; Pipano, 1981; 1997). Other pathogens to consider include
698 the causal agents of bovine tuberculosis and brucellosis as they may spread through contaminated
699 blood used for vaccine production. Most of these agents can be tested by means of specific PCR and
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.

2.3. Requirements for authorisation

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.

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.

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 *al.*, 2009). 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 *A. marginale*.

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.

3. Vaccines based on biotechnology

736 There are no vaccines based on biotechnology available for anaplasmosis.

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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.

2 **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

3 **Paris, 4–8 September 2023**

4 _____

5 # C H A P T E R   3 . 4 . 7 .

6 # BOVINE VIRAL DIARRHOEA

7 ## SUMMARY

8 *Cattle of all ages are susceptible to infection with bovine viral diarrhoea viruses (BVDV), including BVDV*
9 *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*
14 *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*
16 *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*
19 *diarrhoea, invariably leading to death. Mucosal disease can arise only in PI animals. Latent infections*
20 *generally do not occur following recovery from acute infection. However bulls may rarely have a persistent*
21 *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* virus (Pestivirus suis) *and ovine border disease virus̶e̶s̶ (Pestivirus ovis). BVD viruses are*
24 *classified into the distinct species* Pestivirus bovis (commonly known as BVDV type 1), Pestivirus tauri
25 *(BVDV type 2) and* Pestivirus brazilense *(BVDV type 3 or Hobi-like pestivirus). T̶h̶e̶ ̶t̶w̶o̶ ̶g̶e̶n̶o̶t̶y̶p̶e̶s̶ ̶(̶t̶y̶p̶e̶s̶ ̶1̶*
26 *a̶n̶d̶ ̶2̶)̶ ̶a̶r̶e̶ ̶c̶l̶a̶s̶s̶i̶f̶i̶e̶d̶ ̶a̶s̶ ̶s̶e̶p̶a̶r̶a̶t̶e̶ ̶s̶p̶e̶c̶i̶e̶s̶ ̶i̶n̶ ̶t̶h̶e̶ ̶g̶e̶n̶u̶s̶* Pestivirus. A̶ ̶t̶h̶i̶r̶d̶ ̶p̶u̶t̶a̶t̶i̶v̶e̶ ̶g̶e̶n̶o̶t̶y̶p̶e̶,̶ ̶B̶V̶D̶V̶ ̶t̶y̶p̶e̶ ̶3̶,̶*
27 *h̶a̶s̶ ̶a̶l̶s̶o̶ ̶r̶e̶c̶e̶n̶t̶l̶y̶ ̶b̶e̶e̶n̶ ̶p̶r̶o̶p̶o̶s̶e̶d̶.̶ 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*
29 *diagnostic virus isolation in cell cultures. PI animals can be readily identified by a variety of methods aimed*
30 *to detect viral antigens or viral RNA directly in blood and tissues. Virus can also be isolated by inoculation*
31 *of specimens onto susceptible cell cultures followed by immune-labelling methods to detect the replication*
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*
35 *persistently infected requires special attention to specimen transport and testing. RNA detection assays are*
36 *particularly useful because they are rapid, have very high sensitivity and do not depend on the use of cell*
37 *cultures.*

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

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*
45   *cows. Modified live virus vaccine should not be administered to pregnant cattle (or to their sucking calves)*
46   *due to the risk of transplacental infection. Live vaccines that contain cytopathic strains of BVDV present a*
47   *risk of inducing mucosal disease in PI animals. Inactivated viral vaccines are safe and can be given to any*
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*
50   *contamination of batches of fetal calf serum used as a culture medium supplement.*

A. INTRODUCTION

51

1.   Impact of the disease

52

53   Cattle of all ages are susceptible to infection with bovine viral diarrhoea viruses (BVDV). Distribution of the virus is world-
54   wide although some countries have recently eradicated the virus. BVDV infection results in a wide variety of clinical
55   manifestations, including enteric and respiratory disease in any class of cattle or reproductive and fetal disease following
56   infection of a susceptible breeding female. Infection may be subclinical or extend to severe fatal disease. Clinical
57   presentations and severity of disease may vary with different strains of virus. BVDV viruses also cause immune
58   suppression, which can render infected animals more susceptible to infection with other viruses and bacteria. The clinical
59   impact may be more apparent in intensively managed livestock. Animals that survive *in-utero* infection in the first trimester
60   of gestation are almost always persistently infected (PI). PI animals provide the main reservoir of the virus in a population
61   and excrete large amounts of virus in urine, faeces, discharges, milk and semen. The virus spreads mainly by close contact
62   between PI animals and other cattle. Virus shedding by acutely infected animals is usually less important. This virus may
63   also persist in the environment for short periods or be transmitted ~~with~~ via contaminated reproductive materials. Vertical
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,
66   stillbirths, teratogenic abnormalities or the birth of PI calves. Persistently viraemic animals may be born as weak, unthrifty
67   calves or may appear as normal healthy calves and be unrecognised clinically for a long time. However, PI animals have
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
70   leading to death. Mucosal disease can arise only in PI animals. It is important to avoid the trade of viraemic animals. It is
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
74   cannot be completely equated with 'safety'. Detection of PI animals must be specifically directed at detection of the virus
75   or its components (RNA or antigens). Latent infections generally do not occur following recovery from acute infection.
76   However, semen collected from bulls during an acute infection is likely to contain virus during the viraemic period and often
77   for a short time afterwards. Although extremely rare, some recovered bulls may have a persistent testicular infection and
78   excrete virus in semen, perhaps indefinitely.

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

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

2.   The causal agent

88

89   Bovine viral diarrhoea virus (BVDV) is a single linear positive-stranded RNA virus in the genus *Pestivirus* of the family
90   *Flaviviridae.* The genus contains a number of species including ~~the two genotypes of bovine viral diarrhoea virus (BVDV)~~
91   (types 1 [*Pestivirus bovis*]~~,~~ and 2 [*Pestivirus tauri*] and 3 [*Pestivirus brazilense*]) and the ~~closely~~ related classical swine
92   fever (*Pestivirus suis*) and ovine border disease viruses (*Pestivirus ovis*). Viruses in these ~~genotypes~~ pestivirus species
93   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~~ <u>species</u>
95 *Pestivirus bovis* and *tauri*, further subdivisions are discernible by genetic analysis (Vilcek *et al*., 2001). The two ~~genotypes~~
96 <u>species</u> may be differentiated from each other, and from other pestiviruses, by monoclonal antibodies (MAbs) directed
97 against the major glycoproteins E2 and Eᴿᴺˢ or by genetic analysis. Reverse-transcription polymerase chain reaction (RT-
98 PCR) assays enable virus typing direct from blood samples (Letellier & Kerkhofs, 2003; McGoldrick *et al*., 1999). Type 1
99 viruses are generally more common although the prevalence of type 2 strains can be high in North America. BVDV of both
100 ~~genotypes~~ <u>species</u> (*Pestivirus bovis* and *tauri*) may occur in non-cytopathic and cytopathic forms (biotypes), classified
101 according to whether or not microscopically apparent cytopathology is induced during infection of cell cultures. Usually, it
102 is the non-cytopathic biotype that circulates freely in cattle populations. Non-cytopathic strains are most frequently
103 responsible for disease in cattle and are associated with enteric and respiratory disease in any class of cattle or
104 reproductive and fetal disease following infection of a susceptible breeding female. Infection may be subclinical or extend
105 to severe fatal disease (Brownlie, 1985). Cytopathic viruses are encountered in cases of mucosal disease, a clinical
106 syndrome that is relatively uncommon and involves the 'super-infection' of an animal that is PI with a non-cytopathic virus
107 by a closely related cytopathic strain. The two virus biotypes found in a mucosal disease case are usually antigenically
108 closely related if not identical. Type 2 viruses are usually non-cytopathic and have been associated with outbreaks of
109 severe acute infection and a haemorrhagic syndrome. However some type 2 viruses have also been associated with a
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
112 infections are common following infection of non-pregnant animals with either ~~genotype~~ <u>virus species</u>.

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

3. Pathogenesis

3.1. Acute infections

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

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

3.2. *In-utero* infections

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

may appear to be normal at birth but fail to ~~grow normally~~ thrive. They remain PI for life and are usually seronegative, exceptions may be young calves that ingested colostrum containing antibodies. 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.

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 the~~ir~~ progeny of female PI animals will also always be PI. Animals being traded or used for artificial breeding should first be screened to ensure that they are not PI.

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.

3.5. Semen and embryos

Bulls that are PI usually have poor quality, highly infective semen and reduced fertility (McGowan & Kirkland, 1995). All bulls used for natural or artificial insemination should be screened for both acute and persistent BVDV infection. A rare event, possibly brought about by acute infection during pubescence, can result in persistent infection of the 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 donor cows that are PI with BVDV also ~~re~~present a potential source of infection, particularly as there are extremely high concentrations of BVDV in uterine and vaginal fluids. While oocysts without an intact zona pellucida have been 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. Female cattle used as embryo recipients should always be screened to confirm that they are not PI, and ideally, are seropositive or were vaccinated at least 4 weeks before first use.

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

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

4.1. Acute infections

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 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 of animals with respiratory disease or from rectum and faeces if there are enteric signs. Lung and spleen are preferred from dead animals. Viral RNA may be detected by real-time RT-PCR assays and have the advantages of high sensitivity and being able to detect genome from non-infectious virus. As the virus levels are very low, it is not usually practical to undertake virus isolation unless there is a need to characterise the strain of BVDV involved. Serology 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.

Confirmation that an abortion, stillbirth or perinatal death is caused by BVDV is often difficult to establish because 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 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 cases, emphasis should be placed on the detection of viral antigen by ELISA or RNA by real-time RT-PCR. For serology, both ELISAs and virus neutralisation test (VNT) are suitable though sample quality and bacterial contamination may compromise the ability to detect antibodies by VNT. Maternal serology, especially on a group of animals, can be of value, with the aim of determining whether there has been recent infection in the group. A high antibody titre (>1/1000) to BVDV in maternal serum is suggestive of fetal infection and is probably due to the fetus providing the dam with an extended exposure to virus.

4.2. Persistent infections

In the past, identification of PI animals relied heavily on the use of virus isolation in cell cultures. However, antigen detection ELISAs and real-time RT-PCR assays, each with relatively high sensitivity, are widely used for the detection of viral antigens or RNA in both live and dead animals. Virus isolation aimed at the detection of non-cytopathic BVDV 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 virus isolation and IHC are labour intensive and costly and can be technically demanding. Virus isolation from blood can be confounded by the presence of maternal antibody to BVDV in calves less than 4–5 months of age (diagnostic gap). Also for antigen detection ELISAs and flow cytometry from blood or blood leukocytes, there are restrictions that limit when animals that ingested colostrum that contains antibodies against BVDV can be reliably tested. In older 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 individual milk samples have been used to monitor dairy herds for the presence of a PI animal. Antigen ELISA, real-time PCR and virus isolation have all been used. To confirm a diagnosis of persistent infection, animals should be retested after an interval of at least 3 weeks by testing of blood samples for the presence of the virus and for ~~evidence~~ absence of seroconversion. Care should be taken with retesting of skin samples as it has been shown that, in some acute cases, viral antigen may persist for many weeks in skin (Cornish *et al*., 2005).

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.

4.4. Reproductive materials

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 undergoing an acute infection and to establish their serological status. This initial testing should be carried out on whole blood or serum samples. To establish that a seropositive bull does not have a persistent testicular infection (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 also a need to submit a number of straws from each collection, or an appropriate volume of raw semen. Particular care should be taken to ensure that sample transport recommendations are adhered to and that the laboratory 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.

B. DIAGNOSTIC TECHNIQUES

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

Method	Purpose					
	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[(a)]						
Virus isolation	+	++ ~~±~~	++	++ ~~±~~	–	–
Antigen detection by ELISA	++~~+±~~	+++	+++	+++	+++	–
Antigen detection by IHC	–	–	–	++	–	–
NA detection by real-time RT-PCR	+++	+++	+++	+++	+++	–
Detection of immune response						
ELISA	+++	++	+++	~~–~~ +[(a)]	+++	+++
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 method; NA = nucleic acid; RT-PCR = reverse-
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 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 embryos). The exception is for seropositive bulls where semen must be tested rather than the donor bull. Serology only 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* hybridisation (ISH) directly on tissues are not considered to be suitable for certification for freedom from BVDV for international trade purposes. In contrast, immune-staining is an essential component of virus isolation in cell culture to detect the presence of non-cytopathic strains of BVDV which predominate in field infections.

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.

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
281 liquid nitrogen. These can then be tested over a series of passages, or seeded to other susceptible cells and checked
282 for freedom from contaminants and to evaluate their sensitivity compared to an approved batch of cells before routine
283 use. Such problems may be reduced by the use of continuous cell lines, which can be obtained BVD-free, however,
284 their BVDV-free status and susceptibility must be monitored regularly. Continuous cells should be used under a 'seed
285 lot' system where they are only used over a limited passage range, within which they have been shown to have
286 acceptable sensitivity to BVDV infection. Although particular continuous cell lines are considered to be appropriate
287 for use for BVDV isolation, there can be significant variation in batches of cells from different sources due to differing
288 passage histories so their suitability must still be confirmed before routine use.

289 Non-cytopathic BVDV is a common contaminant of bovine tissues and cell cultures must be checked for freedom
290 from adventitious virus by regular testing. Cells must be grown in proven cell culture medium components and a large
291 area of cells must be examined. It is not appropriate to screen a few wells of a 96 well plate – examining all wells of
292 a 96 well plate will be more convincing evidence of freedom. The fetal bovine serum that is selected for use in cell
293 culture must also be free not only from virus, but also and of equal or perhaps even greater importance, from BVDV
294 neutralising antibody. Heat treatment (56°C for 30–45 minutes) is inadequate for the destruction of BVDV in
295 contaminated serum; irradiation with a dose of at least 25 kiloGrays (2.5 Mrad) is more certain. Commercial batches
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
298 levels that are barely detectable but sufficient to inhibit virus isolation. To overcome this, serum can be obtained from
299 BVD virus and antibody free donor animals and used with confidence. Testing of donors for both virus and antibody
300 occurs on an individual animal basis. Although horse serum has been substituted for bovine fetal serum, it is often
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
307 have recovered from acute infection, are no longer viraemic and are now seropositive (Voges *et al.*, 1998). Virus may
308 be detected in most but not all collections of semen from these bulls. Although still considered to be uncommon, to
309 exclude the potential for a PTI it is essential to screen semen from all seropositive bulls. To be confident that a bull
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
312 extended semen, is cytotoxic and must be diluted in culture medium. For these reasons, it is important to monitor the
313 health of the cells by microscopic examination at intervals during the incubation. These problems are largely
314 overcome by the use of real-time RT-PCR which has several advantages over virus isolation, including higher
315 sensitivity and the potential to be completed within a few hours rather than weeks for virus isolation.

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
319 be regularly checked to confirm that they maintain maximum susceptibility to virus infection. Depending on the
320 specimen type and purpose for testing, virus isolation is likely to require one or more passages in cell cultures. While
321 PI animals can be readily identified by screening blood or serum with one passage, semen should be routinely
322 cultured for three passages and biological products such as fetal bovine serum up to five times (original inoculation
323 plus four passages). Conventional methods for virus isolation are used, with the addition of a final immune-staining
324 step (immunofluorescence or, more frequently, peroxidase staining) to detect growth of non-cytopathic virus. Thus
325 tube cultures should include flying cover-slips, while microplate cultures can be fixed and labelled directly in the plate.
326 Examples are given below. Alternatively, culture supernatant from the final passage can be screened by real-time
327 RT-PCR (see below).

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

330 i) 10–25 µl of the serum sample is placed into each of four wells of a 96-well tissue-culture grade
331 microplate. This is repeated for each sample. Known positive and negative controls are
332 included.

333 ii) 100 µl of a cell suspension at the appropriate concentration (usually about 150,000 cells/ml) in
334 medium without fetal calf serum (FCS) is added to all wells. *Note:* the sample itself acts as the
335 cell-growth supplement. If testing samples other than serum, use medium with 10% FCS that is
336 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 31.1.1.i to iv above.
343	vi)	The cells are then fixed and stained by one of two methods:

- **Paraformaldehyde**

345 346	a)	Add 200 µl of a 1/10 dilution of formaldehyde solution (approximately 3% concentration) to 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 µl 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 µl 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 µl 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.

- **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). *Note:* 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 µl) 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 µl 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. Ensure 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		An alternative substrate can be made, consisting of 9 mg diaminobenzidine tetrahydrochloride
387		and 6 mg sodium perborate tetrahydrate dissolved in 15 ml of PBS. Though the staining is not
388		quite so intense, these chemicals have the advantage that they can be shipped by air.

| 389 | k) | The plate is examined microscopically. Virus-positive cells show red-brown cytoplasmic |
| 390 | | staining. |

391		Alternative methods for fixation of the cells may be used and include the use of heat (see Chapter
392		3.8.3 *Classical swine fever*, Section B.2.2.1.viii). These should be first evaluated to ensure that the
393		capacity to detect viral antigen is not compromised.

1.1.2. Tube method for tissue or buffy coat suspensions

| 395 | *Note:* this method can also be conveniently adapted to 24-well plastic dishes. *Note:* a minimum of 2 and |
| 396 | preferably 3 passages (including primary inoculation) is required. |

| 397 | i) | Tissue samples are ground up and a 10% suspension in culture medium is made. This is then |
| 398 | | centrifuged to remove the debris. |

| 399 | ii) | Test tube cultures with newly confluent or subconfluent monolayers of susceptible bovine cells are |
| 400 | | inoculated with 0.1 ml of the sample. The culture is left to adsorb for 1 hour at 37°C. |

| 401 | iii) | The culture is washed with 1 ml of medium; this is then discarded and 1 ml of culture maintenance |
| 402 | | medium is added. |

| 403 | iv) | The culture is incubated for 4–5 days at 37°C, and examined microscopically for evidence of CPE |
| 404 | | or signs of cytotoxicity. |

405	v)	The culture should then be frozen and thawed for passage to fresh cultures for one or preferably
406		two more passages (including the culture inoculated for the final immunostaining). At the final
407		passage, after freeze–thaw the tissue culture fluid is harvested and passaged on to microtitre plates
408		for culture and staining by the immunoperoxidase method (see section B.31.1.1 above) or by the
409		immunofluorescent method. For immunofluorescence, cover-slips are included in the tubes and
410		used to support cultured cells. At the end of the culture period, the cover slips are removed, fixed
411		in 100% acetone and stained with an immunofluorescent conjugate to BVDV. Examine the cover
412		slips under a fluorescent microscope for diffuse, cytoplasmic fluorescence characteristic of
413		pestiviruses. Alternatively, culture supernatant from the final passage can be screened by real-time
414		RT-PCR (see below).

1.1.3. Virus isolation from semen

416	The samples used for the test are, typically, extended bovine semen or occasionally raw semen.
417	Semen samples should be transported to the laboratory in liquid nitrogen, or on dry ice. The samples
418	should be stored in liquid nitrogen or at lower than –70°C (for long-term storage) or 4°C (for short-
419	term storage of not more than 1–2 days). The receiving laboratory should document the condition
420	under which samples are received. Raw semen is generally cytotoxic and should be prediluted (e.g.
421	1/10 in BVDV free bovine serum) before being added to cell cultures. At least 0.1 ml of raw semen
422	should be tested with three passages in cell culture. Toxicity may also be encountered with extended
423	semen. For extended semen, an approximation should be made to ensure that the equivalent of a
424	minimum of 0.1 ml raw semen is examined (e.g. a minimum of 1.0 ml extended semen). If toxicity is
425	encountered, multiple diluted samples may need to be tested to reach a volume equivalent to 0.1 ml
426	raw semen (e.g. 5 × 1 ml of a sample of extended semen that has been diluted 1/5 to reduce toxicity).
427	A suggested method is as follows:

428	i)	Dilute 200 µl fresh semen in 1.8 ml bovine serum containing antibiotics. This can be the same
429		serum as is being used for supplementing the cell cultures, and must be shown to be free from
430		antibodies against BVDV.

| 431 | ii) | Mix vigorously and leave for 30 minutes at room temperature. |

| 432 | iii) | Inoculate 1 ml of the semen/serum mixture into a monolayer of susceptible cells (see virus |
| 433 | | 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 | v) | Remove the mixture, wash the monolayer several times with maintenance medium and then |
| 436 | | 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
439			positive control last.

vi)   Include BVDV negative and positive controls in the test. Special caution must be taken to avoid accidental contamination of test wells by the positive control, for example always handling the positive control last.

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 be inadvertently isolated.

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.

ix)   At the end of the second passage, the supernatant from the freeze-thaw preparation should be 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 well microplates. The sample is considered to be negative, if there is no evidence of viral antigen or BVDV RNA detected.

1.2. Nucleic acid detection

Conventional gel based RT-PCR has in the past been used for the detection of BVD viral RNA for diagnostic 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, 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 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 Chapter 2.2.3 *Development and optimisation of nucleic acid assays*). These assays have even higher sensitivity than gel based RT-PCR and can be completed in a few hours. They are in widespread use for the diagnosis of infectious diseases, allowing the direct detection of viral RNA from a wide range of specimens including serum, whole blood, tissues, milk and semen. The high analytical sensitivity allows the adoption of strategies to screen pools of individual 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 between day 7 and 40 of life, when colostrum that contains antibodies against BVDV was ingested. During this time 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 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 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-labelled DNA probes can also be used to differentiate pestiviruses (e.g. McGoldrick *et al*., 1999).

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 all BVDV types (*Pestivirus bovis, tauri* and *brazilense*), CSFV (*Pestivirus suis*), some strains of BDV (*Pestivirus ovis*) and ~~most of the~~ several 'atypical' pestiviruses (e.g. Hoffman *et al*., 2006). A sensitive broadly reactive assay is recommended for diagnostic applications because interspecies transfer of different pestiviruses is occasionally encountered. When further identification of the specific virus is required, pestivirus species-specific assays can be 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 Mg^{2+}, primers, probe and polymerase, and the cycling parameters need to be determined. However, fully formulated and optimised 'ready to use' 'mastermixes' are now available commercially and only require addition of optimised concentrations of primers and probe. Optimised cycling conditions are often recommended for a particular mastermix.

A variety of commercially available nucleic acid purification systems are available in kit form and several can be semi-automated. Systems based on the capture and purification of RNA using magnetic beads are in widespread use and allow rapid processing of large numbers of samples. Specific products should be evaluated to determine the optimal 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 PCR reaction are encountered with samples collected into heparin treated blood than EDTA. These differences are also exacerbated if the tube does not contain the recommended volume of blood, thereby increasing the concentration of anticoagulant in the sample. To identify possible false-negative results, it is recommended to spike an exogenous ('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 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

| 495 | is necessary to ensure that PCR amplification of the internal control does not compete with the diagnostic PCR and |
| 496 | thus lower the analytical sensitivity (see also chapter 1.1.6). |

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

503   1.2.1.   Real-time polymerase chain reaction for BVDV detection in semen

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

516   i)   Sample preparation, equipment and reagents

517   a)   The samples used for the test are, typically, extended bovine semen or occasionally raw
518 semen. If the samples are only being tested by real-time RT-PCR, it is acceptable for them
519 to be submitted chilled but they must still be cold when they reach the laboratory.
520 Otherwise, if a cold chain cannot be assured or if virus isolation is being undertaken, the
521 semen samples should be transported to the laboratory in liquid nitrogen or on dry ice. At
522 the laboratory, the samples should be stored in liquid nitrogen or at lower than –70°C (for
523 long-term storage) or 4°C (for short-term storage of up to 7 days). *Note:* samples for virus
524 isolation should not be stored at 4°C for more than 1–2 days.

525   b)   Due to the very high analytical sensitivity of real-time RT-PCR, much smaller volumes of
526 semen may be used. However, at least three straws (minimum 250 µl each) from each
527 collection batch of semen should be processed. The semen in the three straws should be
528 pooled and mixed thoroughly before taking a sample for nucleic acid extraction.

529   c)   A real-time PCR detection system, and the associated data analysis software, is required
530 to perform the assay. A number of real-time PCR detection systems are available from
531 various manufacturers. ̶O̶t̶h̶e̶r̶ ̶e̶q̶u̶i̶p̶m̶e̶n̶t̶ ̶r̶e̶q̶u̶i̶r̶e̶d̶ ̶f̶o̶r̶ ̶t̶h̶e̶ ̶t̶e̶s̶t̶ ̶i̶n̶c̶l̶u̶d̶e̶s̶ ̶a̶ ̶m̶i̶c̶r̶o̶-̶c̶e̶n̶t̶r̶i̶f̶u̶g̶e̶,̶
532 a̶ ̶c̶h̶i̶l̶l̶i̶n̶g̶ ̶b̶l̶o̶c̶k̶,̶ ̶a̶ ̶m̶i̶c̶r̶o̶-̶v̶o̶r̶t̶e̶x̶,̶ ̶a̶n̶d̶ ̶m̶i̶c̶r̶o̶p̶i̶p̶e̶t̶t̶e̶s̶.̶ As real-time RT-PCR assays are able
533 to detect very small amounts of target nucleic acid molecules, appropriate measures are
534 required to avoid contamination. ̶,̶ ̶i̶n̶c̶l̶u̶d̶i̶n̶g̶ ̶d̶e̶d̶i̶c̶a̶t̶e̶d̶ ̶a̶n̶d̶ ̶p̶h̶y̶s̶i̶c̶a̶l̶l̶y̶ ̶s̶e̶p̶a̶r̶a̶t̶e̶d̶ ̶'̶c̶l̶e̶a̶n̶'̶
535 a̶r̶e̶a̶s̶ ̶f̶o̶r̶ ̶r̶e̶a̶g̶e̶n̶t̶ ̶p̶r̶e̶p̶a̶r̶a̶t̶i̶o̶n̶ ̶(̶w̶h̶e̶r̶e̶ ̶n̶o̶ ̶s̶a̶m̶p̶l̶e̶s̶ ̶o̶r̶ ̶m̶a̶t̶e̶r̶i̶a̶l̶s̶ ̶u̶s̶e̶d̶ ̶f̶o̶r̶ ̶P̶C̶R̶ ̶a̶r̶e̶ ̶h̶a̶n̶d̶l̶e̶d̶)̶,̶
536 a̶ ̶d̶e̶d̶i̶c̶a̶t̶e̶d̶ ̶s̶a̶m̶p̶l̶e̶ ̶p̶r̶o̶c̶e̶s̶s̶i̶n̶g̶ ̶a̶r̶e̶a̶ ̶a̶n̶d̶ ̶a̶n̶ ̶i̶s̶o̶l̶a̶t̶e̶d̶ ̶a̶r̶e̶a̶ ̶f̶o̶r̶ ̶t̶h̶e̶ ̶P̶C̶R̶ ̶t̶h̶e̶r̶m̶o̶c̶y̶c̶l̶e̶r̶ ̶a̶n̶d̶
537 a̶s̶s̶o̶c̶i̶a̶t̶e̶d̶ ̶e̶q̶u̶i̶p̶m̶e̶n̶t̶.̶ ̶E̶a̶c̶h̶ ̶a̶r̶e̶a̶ ̶s̶h̶o̶u̶l̶d̶ ̶h̶a̶v̶e̶ ̶d̶e̶d̶i̶c̶a̶t̶e̶d̶ ̶r̶e̶a̶g̶e̶n̶t̶s̶ ̶a̶n̶d̶ ̶e̶q̶u̶i̶p̶m̶e̶n̶t̶.̶
538 Furthermore, a minimum of one negative sample should be processed in parallel to
539 monitor the possibility of low level contamination. Sources of contamination may include
540 product carry-over from positive samples or, more commonly, from cross contamination
541 by PCR products from earlier work.

542   d)   The real-time RT-PCR assay involves two separate procedures.

543   1)   Firstly, BVDV RNA is extracted from semen using an appropriate validated nucleic
544 acid extraction method. Systems using magnetic beads for the capture and
545 purification of the nucleic acid are recommended. It is also preferable that the beads
546 are handled by a semi-automated magnetic particle handling system.

547   2)   The second procedure is the RT-PCR analysis of the extracted RNA template in a
548 real-time RT-PCR system.

549   ii)   Extraction of RNA

| 550 | | | | RNA or total nucleic acid is extracted from the pooled (three straws collected at the same time from the same animal) semen sample. Use of a commercially available magnetic bead based extraction kit is recommended. However, the preferred kit should be one that has been evaluated to ensure optimal extraction of difficult samples (semen and whole blood). Some systems and kit protocols are sufficiently refined that it is not necessary to remove cells from the semen sample. Prior to extraction dilute the pooled semen sample 1/4 in ~~phosphate buffered gelatin saline (PBGS)~~ or a similar buffered solution. Complete the RNA extraction by taking 50 µl of the diluted, pooled sample and add it to the sample lysis buffer. Some commercial extraction kits may require the use of a larger volume. It has also been found that satisfactory results are obtained by adding 25 µl of undiluted pooled sample to sample lysis buffer. Complete the extraction by following the kit manufacturer's instructions. |

561   iii)   Real-time RT-PCR assay procedure

562   a)   Reaction mixture: There are a number of commercial real-time PCR amplification kits available from various sources and the particular kits selected need to be compatible with the real-time PCR platform selected. The required primers and probes can be synthesised by various commercial companies. The WOAH Reference Laboratories for BVDV can provide information on suitable suppliers.

567   b)   Supply and storage of reagents: The real-time PCR reaction mixture is normally provided as a 2 × concentration ready for use. The manufacturer's instructions should be followed for application and storage. Working stock solutions for primers and probe are made with nuclease-free water at the concentration of 20 µM and 3 µM, respectively. The stock solutions are stored at –20°C and the probe solution should be kept in the dark. Single-use or limited use aliquots can be prepared to limit freeze–thawing of primers and probes and extend their shelf life.

574   c)   Primers and probe sequences

575   Selection of the primers and probe are outlined in Hoffmann *et al*. (2006) and summarised below.

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)   Preparation of reaction mixtures

581   The PCR reaction mixtures are prepared in a separate room that is isolated from other PCR activities and sample handling. For each PCR test, appropriate controls should be included. As a minimum, a no template control (NTC), appropriate negative control (NC) <u>and</u> two positive controls (PC1, PC2) should be included. The positive and negative controls are included in all steps of the assay from extraction onwards while the NTC is added after completion of the extraction. The PCR amplifications are carried out in a volume of 25 µl. The protocol described is based on use of a 96 well microplate based system but other options using microtubes are also suitable. Each well of the PCR plate should contain 20 µl of reaction mix and 5 µl 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   <u>nuclease free</u> water

596   1 µl   25× enzyme mix

597   5 µl   sample (or controls – NTC, NC, PC1, PC2)

598		e)	Selection of controls

NTC: usually consists of <u>nuclease free water or</u> tRNA in nuclease free water that is added in place of a sample when the PCR reaction is set up.

NC: In practice, many laboratories use PBGS or a similar buffer. Ideally the controls for testing of semen samples should be negative semen, from seronegative bulls. However, as a minimum, the assay in use should have been extensively validated with negative and positive samples to confirm that it gives reliable extraction and amplification with semen.

PCs: There are two positive controls (PC1=moderate – [Ct 29-32] and PC2=weak [Ct 32–35] positive). Positive semen from naturally infected bulls is preferable as a positive 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 reliable indication of any moderate reduction in extraction or assay performance. Negative semen spiked with defined quantities of BVDV virus could be used as an alternative. If other samples are used as a routine PC, as a minimum the entire extraction process and PCR assay in use must have been extensively validated using known positive semen from bulls with a PTI or from bulls undergoing an acute infection. If these samples are not 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 of an exogenous control with each test sample will largely compensate for not using positive semen as a control and will give additional benefits by monitoring the efficiency of the assay on each individual sample. Positive control samples should be prepared carefully to avoid cross contamination from high titred virus stocks and should be prepared in advance and frozen at a 'ready to use' concentration and ideally 'single use' volume.

f) Extracted samples are added to the PCR mix in a separate room. The controls should be added last, in a consistent sequence in the following order: NTC, negative and then the two positive controls.

g) Real-time polymerase chain reaction

The PCR plate or tubes are placed in the real-time PCR detection system in a separate, designated PCR room. Some mastermixes have uniform reaction conditions that are suitable for many different assays. As an example, the PCR detection system is programmed for the test as follows:

1× 48°C 10 minutes

1× 95°C 10 minutes

45 × (95°C 15 seconds, 60°C 1 minute)

h) Analysis of real-time PCR data

The software program is usually set to automatically adjust results by compensating for any background signal and the threshold level is usually set according to the manufacturer's instructions for the selected analysis software used. In this instance, a threshold is set at 0.05.

i) Interpretation of results

a) Test controls – all controls should give the expected results with positive controls (PC1 and PC2) falling within the designated range and both the negative control (NC) and no template control (NTC) should have no Ct values.

b) Test samples

1) Positive result: Any sample that has a cycle threshold (Ct) value less than 40 is regarded as positive.

2) Negative result: Any sample that shows no Ct value is regarded as negative. However, before reporting a negative result for a sample, the performance of the exogenous internal control should be checked and shown to give a result within the accepted range for that control (for example, a Ct value no more than 2–3 Ct units higher than the NTC).

1.3. Enzyme-linked immunosorbent assay for antigen detection

Antigen detection by ELISA has become a widely adopted method for the detection of individual PI animals. These 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 manufacture. Several methods for the ELISA for antigen detection have been published and a number of commercial kits are available. Most are based on the sandwich ELISA principle, with a capture antibody bound to the solid phase, and a detector antibody conjugated to a signal system, such as peroxidase. Amplification steps such as the use of biotin and streptavidin in the detection system are sometimes used to increase assay sensitivity. Both monoclonal- and polyclonal-based systems are described. The test measures BVD antigen (NS2-3 or Eʀɴs) in lysates of peripheral blood leukocytes; the new generation of antigen-capture ELISAs (Eʀɴs capture ELISAs) are able to detect BVD 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 transient viraemia, the antigen ELISA is less useful for virus detection in acute BVD infections.

The ~~NS2-3~~ antigen detection ELISA~~s~~ may be less effective in young calves that have had colostrum due to the presence of BVDV maternal antibodies, <u>especially when blood samples or blood leucocytes are tested (Fux & Wolf, 2012). Blood or blood leucocytes should not be tested in the first month (Eʀɴs capture ELISA) or the first 3 months (NS2-3 ELISA) of life due to the inhibitory effect of maternal antibodies.</u> The real-time RT-PCR is probably the most sensitive detection method for this circumstance, but the Eʀɴs ELISA has also been shown to be a sensitive and reliable test, ~~particularly~~ when used with skin biopsy (ear-notch) samples (Cornish *et al*., 2005).

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.

2. Serological tests

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 presence of PI animals in a herd, to assist with investigation of reproductive disease and possible involvement of BVDV and to establish the serological status of bulls being used for semen collection and to identify whether there has been a 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 value<u>s</u> ~~(0.8 or more absorbance units)~~ in an unvaccinated herd indicates a high probability of the herd having been exposed to BVDV in the recent past, most likely through one or more persistently viraemic animals being present. In contrast, ~~a~~ very low or negative value<u>s</u> ~~(≤0.2)~~ indicate<u>s</u> 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 husbandry (Zimmer *et al*., 2002), recent administration of vaccine and also due to the presence of viral antigen in bulk milk, which may interfere with the antibody assay itself. Determination of the antibody status of a small number of young stock (9–18 months) has also been utilised as an indicator of recent transmission of BVDV in the herd (Houe *et al*., 1995), but this approach is also dependent on the degree of contact between different groups of animals in the herd and the potential for exposure from neighbouring herds. VN tests are more frequently used for regulatory purposes (e.g. testing of semen 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 results within predetermined limits for the test to be considered valid. In the VNT, a 'serum control' to monitor sample toxicity should also be included for each test sample.

2.1. Virus neutralisation test

Selection of the virus strain to include in a VNT is very important. No single strain is likely to be ideal for all 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 neutralisation test that uses type 1 strain of the virus, and vice versa (Fulton *et al*., 1997). It is important that BVD type 1 and BVD type 2 (*Pestivirus bovis* <u>and</u> *tauri*) be used in the test and not just the one that the diagnostician thinks is present, as this can lead to under reporting. Because it makes the test easier to read, most laboratories use highly cytopathic, laboratory-adapted strains of BVDV for VN tests. Two widely used cytopathic strains are 'Oregon C24V' and 'NADL'. However immune-labelling techniques are now available that allow simple detection of the growth

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

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.

iii) An equal volume (e.g. 50 μl) of a stock of cytopathic strain of BVDV containing 100 $TCID_{50}$ (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 $TCID_{50}$).

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 1.5×10^5/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% CO_2 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.

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 ~~genotype~~ species of BVD~~V~~ may not efficiently detect antibody induced by another ~~genotype~~ virus species. 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 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 system. The virus can be concentrated and purified by density gradient centrifugation. Alternatively, a potent antigen can be prepared by treatment of infected cell cultures with detergents, such as Nonidet P40, N-decanoyl-N-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 expressing specific viral genes in bacterial or eukaryotic systems. Such systems should be validated by testing sera specific to a wide range of different virus strains. In the future, this technology should enable the production of serological tests complementary to subunit or marker vaccines, thus enabling differentiation between vaccinated and naturally infected cattle. An example outline protocol for an indirect ELISA is given below (Edwards, 1990).

2.2.1. Test procedure

i) 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 |
| 755 | | in the test. |

756	iv)	Test sera are diluted 1/50 in serum diluent (0.5 M NaCl; 0.01 M phosphate buffer; 0.05% Tween
757		20; 0.001 M ethylene diamine tetra-acetic acid; 1% polyvinyl pyrrolidone, pH 7.2) and added to
758		virus- and control-coated wells for 1 hour at 37°C. The plates are then washed five times in
759		PBST.

| 760 | v) | Rabbit anti-bovine IgG peroxidase conjugate is added at a predetermined dilution (in serum |
| 761 | | diluent) for 1 hour at 37°C, then the plates are again washed five times in PBST. |

762	vi)	A suitable enzyme substrate is added, such as hydrogen peroxide/tetramethyl benzidine. After
763		colour development, the reaction is stopped with sulphuric acid and the absorbance is read on
764		an ELISA plate reader. The value obtained with control antigen is subtracted from the test
765		reaction to give a net absorbance value for each serum.

766	vii)	It is recommended to convert net absorbance values to sample:positive ratio (or percentage
767		positivity) by dividing net absorbance by the net absorbance on that test of a standard positive
768		serum that has a net absorbance of about 1.0. This normalisation procedure leads to more
769		consistent and reproducible results.

C. REQUIREMENTS FOR VACCINES

1. Background

770

771

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

1.1. Characteristics of a target product profile

786

787 Traditionally, BVD vaccines fall into two classes: modified live or inactivated virus vaccines. The essential requirement
788 for both types is to ~~afford~~ provide a high level of fetal ~~infection~~ protection. Many of the live vaccines have been based
789 on a cytopathic strain of the virus which is considered to be unable to cross the placenta. However, it is important to
790 ensure that the vaccine virus does not cause fetal infection. In general, vaccination of breeding animals should be
791 completed well before insemination to ensure optimal protection and avoid any risk of fetal infection. Live virus vaccine
792 may also be immunosuppressive and precipitate other infections. On the other hand, modified live virus vaccines may
793 only require a single dose. Use of a live product containing a cytopathic strain of BVDV may precipitate mucosal
794 disease by superinfection of persistently viraemic animals. Properly formulated inactivated vaccines are very safe to
795 use but, to obtain satisfactory levels of immunity, they usually require booster vaccinations, which may be
796 inconvenient. A combined vaccination protocol using inactivated followed by live vaccine may reduce the risk of
797 adverse reaction to the live strain. Whether live or inactivated, because of the propensity for antigenic variability, the
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
800 contain a suitable type 2 strain. For optimal immunity against type 1 strains (*Pestivirus bovis*), antigens from the
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.

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

2. Outline of production and minimum requirements for vaccines

2.1. Characteristics of the seed

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 (*Pestivirus tauri*) 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 purification of the two biotypes from an initial mixed culture is important to maintain the expected characteristics of the ~~seen~~ seed and depends on several cycles of a limiting dilution technique for the 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, their identity and key antigenic characteristics should be confirmed. The identity of the seed virus should be confirmed by sequencing. Where there are multiple isolates included in the vaccine, each has to be prepared separately.

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 not be transmissible to unvaccinated 'in-contact' animals and should not be able to infect the fetus. 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 cultures incorporated into the final product. Master stocks for either live or inactivated vaccines should be prepared under a seed lot system involving master and working stocks that can be used for production in such a manner that the number of passages can be limited and minimise antigenic drift. While there are no absolute criteria for this purpose, as a general guide, the seed used for production should not be passaged more than 20 times beyond the master seed and the master seed should be 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 ~~geno~~types 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.

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 complete prevention of fetal infection, a vaccine would be expected to be highly effective in other situations (for example prevention of respiratory disease). A suitable challenge system can be 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 offspring in non-immune cows. In countries where BVDV type 2 viruses (*Pestivirus tauri*) are 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.

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.

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).

Live vaccines must either be demonstrated to be safe in pregnant cattle (i.e. no transmission to the fetus), or should be licensed with a warning not to use them in pregnant animals. Live vaccines containing cytopathic strains should have an appropriate warning of the risk of inducing mucosal 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

919 2.3.1. Manufacturing process

920 For registration of a vaccine, all relevant details concerning manufacture of the vaccine and quality
921 control testing should be submitted to the relevant authorities. Unless otherwise specified by the
922 authorities, information should be provided from three consecutive vaccine batches with a volume
923 not less than 1/3 of the typical industrial batch volume.

924 There is no standard method for the manufacture of a BVD vaccine, but conventional laboratory
925 techniques with stationary, rolled or suspension (micro-carriers) cell cultures may be used.
926 Inactivated vaccines can be prepared by conventional methods, such as binary ethylenimine or beta-
927 propiolactone inactivation (Park & Bolin, 1987). A variety of adjuvants may be used.

928 2.3.2. Safety requirements

929 *In-vivo* tests should be undertaken using a single dose, overdose (for live vaccines only) and repeat
930 doses (taking into account the maximum number of doses for primary vaccination and, if appropriate,
931 the first revaccination/booster vaccination) and contain the maximum permitted antigen load and,
932 depending on the formulation of the vaccine, the maximum number of vaccine strains.

933 i) Target and non-target animal safety

934 The safety of the final product formulation of both live and inactivated vaccines should be assessed
935 in susceptible young calves that are free of maternally derived antibodies and in pregnant cattle. They
936 should be checked for any local reactions following administration, and, in pregnant cattle, for any
937 effects on the unborn calf. Live attenuated vaccines may contribute to immunosuppression that might
938 increase mortality. It may also contribute to the development of mucosal disease in PI animals that
939 is an animal welfare concern. Therefore vaccination of PI animals with live attenuated vaccines
940 containing cytopathic BVDV should be avoided. Live attenuated vaccines must not be capable of
941 being transmitted to other unvaccinated animals that are in close contact.

942 ii) Reversion-to-virulence for attenuated/live vaccines and environmental considerations

943 Virus seeds that have been passaged at least up to and preferably beyond the passage limit specified
944 for the seed should be inoculated into young calves to confirm that there is no evidence of disease.
945 If a live attenuated vaccine has been registered for use in pregnant animals, reversion to virulence
946 tests should also include pregnant animals. Live attenuated vaccines should not be transmissible to
947 unvaccinated 'in-contact' animals.

948 iii) Precautions (hazards)

949 BVDV is not considered to be a human health hazard. Standard good microbiological practice should
950 be adequate for handling the virus in the laboratory. A live virus vaccine should be identified as
951 harmless for people administering the product. However adjuvants included in either live or
952 inactivated vaccines may cause injury to people. Manufacturers should provide adequate warnings
953 that medical advice should be sought in the case of self-injection (including for adjuvants, oil-emulsion
954 vaccine, preservatives, etc.) with warnings included on the product label/leaflet so that the vaccinator
955 is aware of any danger.

956 2.3.3. Efficacy requirements

957 The potency of the vaccine should be determined by inoculation into seronegative and virus negative
958 calves, followed by monitoring of the antibody response. Antigen content can be assayed by ELISA
959 and adjusted as required to a standard level for the particular vaccine. Standardised assay protocols
960 applicable to all vaccines do not exist. Live vaccine batches may be assayed by infectivity titration.
961 Each production batch of vaccine should undergo potency and safety testing as batch release criteria.
962 BVD vaccines must be demonstrated to produce adequate immune responses, as outlined above,
963 when used in their final formulation according to the manufacturer's published instructions.

2.3.4. Vaccines permitting a DIVA strategy (detection of infection in vaccinated animals)

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.

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.

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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
1062   diagnostic tests, reagents and vaccines for bovine viral diarrhoea

1063   **NB:** FIRST ADOPTED IN 1990. MOST RECENT UPDATES ADOPTED IN 2015.

2   **MEETING OF THE WOAH BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # C H A P T E R   3 . 4 . 1 2 .

6   # LUMPY SKIN DISEASE

7   ## 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,*
10   *oedema of the skin, and sometimes death. The disease is of economic importance as it can cause a*
11   *temporary reduction in milk production, temporary or permanent sterility in bulls, damage to hides and,*
12   *occasionally, death. Various strains of capripoxvirus are responsible for the disease. These are antigenically*
13   *indistinguishable from strains causing sheep pox and goat pox yet distinct at the genetic level. LSD has a*
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*
17   *inefficient. Lumpy skin disease is endemic in* <u>most</u> ~~many~~ *African and Middle Eastern countries. Between*
18   *2012 and 2022, LSD spread into south-east Europe, the Balkans, Russia and Asia as part of the Eurasian*
19   *LSD epidemic.*

20   ***Pathology:*** *the nodules are firm, and may extend to the underlying subcutis and muscle. Acute histological*
21   *key lesions consist of epidermal vacuolar changes with intracytoplasmic inclusion bodies and dermal*
22   *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*
29   *infection and both are uncommon in cattle. LSDV will grow in tissue culture of bovine, ovine or caprine origin.*
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.*

35   *A variety of conventional and real-time PCR tests as well as isothermal amplification tests using*
36   *capripoxvirus-specific primers have been published for use on a variety of samples.*

37   ***Serological tests:*** *The virus neutralisation test (VNT) and enzyme-linked immunosorbent assays (ELISAs)*
38   *are widely used and have been validated. The agar gel immunodiffusion test and indirect immunofluorescent*
39   *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.*

42   ***Requirements for vaccines:*** *All strains of capripoxvirus examined so far, whether derived from cattle,*
43   *sheep or goats, are antigenically similar. Attenuated cattle strains, and strains derived from sheep and goats*
44   *have been used as live vaccines against LSDV.*

A. INTRODUCTION

46   Lumpy skin disease (LSD) was first seen in Zambia in 1929, spreading into Botswana by 1943 (Haig, 1957), and then into
47   South Africa the same year, where it affected over eight million cattle causing major economic loss. In 1957 it entered
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.
50   Another epizootic of LSD between 1981 and 1986 affected Tanzania, Kenya, Zimbabwe, Somalia and the Cameroon, with
51   reported mortality rates in affected cattle of 20%. The occurrence of LSD north of the Sahara desert and outside the African
52   continent was confirmed for the first time in Egypt and Israel between 1988 and 1989, and was reported again in 2006
53   (Brenner *et al.,* 2006). In the past decade, LSD occurrences have been reported in the Middle Eastern, European and
54   Asian regions (for up-to-date information, consult WOAH WAHIS interface[40]). Lumpy skin disease outbreaks tend to be
55   sporadic, depending upon animal movements, immune status, and wind and rainfall patterns affecting vector populations.
56   The principal method of transmission is thought to be mechanical by various arthropod vectors (Tuppurainen *et al*., 2015).

57   Lumpy skin disease virus (LSDV) belongs to the family *Poxviridae*, subfamily ~~*Chordopoxvirinae*~~ *Chordopoxviridae*, and
58   genus *Capripoxvirus*. In common with other poxviruses LSDV replicates in the cytoplasm of an infected cell, forming distinct
59   perinuclear viral factories. The LSD virion is large and brick-shaped measuring 293–299nm (length) and 262–273nm
60   (width). The LSDV genome structure is also similar to other poxviruses, consisting of double-stranded linear DNA that is
61   25% GC-rich, approximately 150,000 bp in length, and encodes around 156 open reading frames (ORFs). An inverted
62   terminal repeat sequence of 2200–2300 bp is found at each end of the linear genome. The linear ends of the genome are
63   joined with a hairpin loop. The central region of the LSDV genome contains ORFs predicted to encode proteins required
64   for virus replication and morphogenesis and exhibit a high degree of similarity with genomes of other mammalian
65   poxviruses. The ORFs in the outer regions of the LSDV genome have lower similarity and likely encode proteins involved
66   in viral virulence and host range determinants.

67   Phylogenetic analysis shows the majority of LSDV strains group into two monophyletic clusters (cluster 1.1 and 1.2)
68   (Biswas *et al.,* 2020; Van Schalkwyk *et al.,* 2021). Cluster 1.1 consists of LSDV Neethling vaccine strains that are based
69   on the LSDV/Neethling/LW-1959 vaccine strain (Kara *et al.,* 2003; Van Rooyen *et al.,* 1959; van Schalkwyk *et al.,* 2020)
70   and historic wild-type strains from South Africa. Cluster 1.2 consists of wild-type strains from southern Africa, Kenya, the
71   northern hemisphere, and the Kenyan KSGP O-240 commercial vaccine. In addition to these two clusters, there have
72   recently been recombinant LSDV strains isolated from clinical cases of LSD in the field in Russia and central Asia (Flannery
73   *et al.,* 2021; Sprygin *et al.,* 2018; 2020; Wang *et al.,* 2021). These recombinant viruses show unique patterns of accessory
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
76   capripoxvirus the age of the host, immunological status and breed. *Bos taurus* is generally more susceptible to clinical disease
77   than *Bos indicus;* the Asian buffalo (*Bubalus spp.*) has also been reported to be susceptible. Within *Bos taurus,* the fine-
78   skinned Channel Island breeds develop more severe disease, with lactating cows appearing to be the most at risk. However,
79   even among groups of cattle of the same breed kept together under the same conditions, there is a large variation in the
80   clinical signs presented, ranging from subclinical infection to death (Carn & Kitching, 1995). There may be failure of the virus
81   to infect the whole group, probably depending on the virulence of the virus isolate, immunological status of the host, host
82   genotype, and vector prevalence. Seroprevalence studies, experimental infections and case reports have provided indications
83   that several wildlife species (e.g. springbok, impala, giraffe, camel, banteng) are susceptible to LSDV infection (Dao *et al.,*
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
86   the role of wildlife in LSDV epidemiology. This topic deserves further study, especially given the current spread of LSDV in
87   new geographical areas where large numbers of naïve, potentially susceptible wild bovines and other ruminants are present.

88   The incubation period under field conditions has not been reported, but following experimental inoculation is 6–9 days until
89   the onset of fever. In the acutely infected animal, there is an initial pyrexia, which may exceed 41°C and persist for 1 week.
90   All the superficial lymph nodes become enlarged. In lactating cattle there is a marked reduction in milk yield. Lesions
91   develop over the body, particularly on the head, neck, udder, scrotum, vulva and perineum between 7 and 19 days after
92   virus inoculation (Coetzer, 2004). The characteristic integumentary lesions are multiple, well circumscribed to coalescing,
93   0.5–5 cm in diameter, firm, flat-topped papules and nodules. The nodules involve the dermis and epidermis, and may

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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
96 sequestrum of necrotic material/necrotic plug ("sit-fast") may appear within the nodule. The acute histological lesions
97 consist of epidermal vacuolar changes with intracytoplasmic inclusion bodies and dermal vasculitis. The inclusion bodies
98 are numerous, intracytoplasmic, eosinophilic, homogenous to occasionally granular and they may occur in endothelial
99 cells, fibroblasts, macrophages, pericytes, and keratinocytes. The dermal lesions include vasculitis with fibrinoid necrosis,
100 oedema, thrombosis, lymphangitis, dermal-epidermal separation, and mixed inflammatory infiltrate. The chronic lesions
101 are characterised by an infarcted tissue with a sequestered necrotic core, often rimmed by granulation tissue gradually
102 replaced by mature fibrosis. At the appearance of the nodules, the discharge from the eyes and nose becomes
103 mucopurulent, and keratitis may develop. Nodules may also develop in the mucous membranes of the mouth and
104 alimentary tract, particularly the abomasum and in the trachea and the lungs, resulting in primary and secondary
105 pneumonia. The nodules on the mucous membranes of the eyes, nose, mouth, rectum, udder and genitalia quickly
106 ulcerate, and by then all secretions, ocular and nasal discharge and saliva contain LSD virus (LSDV). The limbs may be
107 oedematous and the animal is reluctant to move. Pregnant cattle may abort, and there is a report of intrauterine
108 transmission (Rouby & Aboulsoudb, 2016). Bulls may become permanently or temporarily infertile and the virus can be
109 excreted in the semen for prolonged periods (Irons *et al*., 2005). Recovery from severe infection is slow; the animal is
110 emaciated, may have pneumonia and mastitis, and the necrotic plugs of skin, which may have been subject to fly strike,
111 are shed leaving deep holes in the hide (Prozesky & Barnard, 1982).

112 The main differential diagnosis is pseudo-LSD caused by bovine herpesvirus 2 (BoHV-2). This is usually a milder clinical
113 condition, characterised by superficial nodules, resembling only the early stage of LSD. Intra-nuclear inclusion bodies and
114 viral syncytia are histopathological characteristics of BoHV-2 infection not seen in LSD. Other differential diagnoses (for
115 integumentary lesions) include: dermatophilosis, dermatophytosis, bovine farcy, photosensitisation, actinomycosis,
116 actinobacillosis, urticaria, insect bites, besnoitiosis, nocardiosis, demodicosis, onchocerciasis, pseudo-cowpox, and
117 cowpox. Differential diagnoses for mucosal lesions include: foot and mouth disease, bluetongue, mucosal disease,
118 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*).

B. DIAGNOSTIC TECHNIQUES

123 *Table 1. Test methods available for the diagnosis of LSD and their purpose*

Method	Purpose					
	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	–	–	–	+	–	–
Detection of immune response						
VNT	++	++	++	++	++	++
IFAT	+	+	+	+	+	+
ELISA	++	++	++	++	++	++

124 Key: +++ = recommended for this purpose; ++ recommended but has limitations;
125 + = suitable in very limited circumstances; – = not appropriate for this purpose.
126 PCR = polymerase chain reaction; TEM = Transmission electron microscopy; VNT = virus neutralisation test;
127 IFAT = indirect fluorescent antibody test; ELISA = enzyme-linked immunosorbent assay.

1. Detection of the agent

1.1. Specimen collection, submission and preparation

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 clinical signs, before the development of neutralising antibodies (Davies, 1991; Davies *et al*., 1971), however virus can be isolated from skin nodules for at least 3–4 weeks thereafter. Samples for genome detection using conventional or real-time polymerase chain reaction (PCR) may be collected when neutralising antibody is present. Following the 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 into heparin or EDTA (ethylene diamine tetra-acetic acid) during the viraemic stage of LSD (before generalisation of lesions or within 4 days of generalisation), can also be used for virus isolation. <u>Samples for histology should include the lesion and tissue from the surrounding (non-lesion) area, be a maximum size of 2 cm^3, and be placed immediately following collection into ten times the sample volume of 10% neutral buffered formal saline.</u>

<u>Tissues in formalin have no special transportation requirements in regard to biorisks.</u> 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.

~~Samples for histology should include the lesion and tissue from the surrounding (non-lesion) area, be a maximum size of 2 cm^3, and be placed immediately following collection into ten times the sample volume of 10% neutral buffered formaldehyde. Tissues in formalin have no special transportation requirements in regard to biorisks.~~ Material for histology should be prepared using standard techniques and stained with haematoxylin and eosin (H&E) (Burdin, 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 sand and an equal volume of sterile phosphate buffered saline (PBS) or serum-free modified Eagle's medium containing sodium penicillin (1000 international units [IU]/ml), streptomycin sulphate (1 mg/ml), mycostatin (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 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 after the centrifugation step. Buffy coats may be prepared from unclotted blood using centrifugation at 600 ***g*** for 15 minutes, and the buffy coat carefully removed into 5 ml of cold double-distilled water using a sterile Pasteur pipette. After 30 seconds, 5 ml of cold double-strength growth medium is added and mixed. The mixture is centrifuged at 600 ***g*** for 15 minutes, the supernatant is discarded and the cell pellet is suspended in 5 ml growth medium, such as Glasgow's modified Eagle's medium (GMEM). After centrifugation at 600 ***g*** for a further 15 minutes, the resulting pellet is suspended in 5 ml of fresh GMEM. Alternatively, the buffy coat may be separated from a heparinised sample by using a Ficoll gradient.

1.2. Virus isolation on cell culture

LSDV will grow in tissue culture of bovine, ovine or caprine origin. MDBK (Madin–Darby bovine kidney) cells are often used, as they support good growth of the virus and are well characterised (Fay *et al.,* 2020). Primary cells, such as 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 monolayer in a 25 cm^2 culture flask at 37°C and allowed to adsorb for 1 hour. The culture is then washed with warm 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, are also infected.

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 rounding of cells and margination of the nuclear chromatin. At first only small areas of CPE can be seen, sometimes as soon as 2 days after infection; over the following 4–6 days these expand to involve the whole cell <u>monolayer</u> ~~sheet~~. If no CPE is apparent by day 14, the culture should be freeze–thawed three times, and clarified supernatant inoculated 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 being used, a cover-slip should be removed, fixed in acetone and stained using H&E. Eosinophilic intracytoplasmic inclusion bodies, which are variable in size but up to half the size of the nucleus and surrounded by a clear halo, are diagnostic for poxvirus infection. PCR may be used as an alternative to H&E for confirmation of the diagnosis. The

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.Ts) 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.

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

192　The extraction method described below can be replaced using commercially available DNA extraction
193　kits.

194　i)　Freeze and thaw 200 µl of blood in EDTA, semen or tissue culture supernatant and suspend in
195　　　100 µl of lysis buffer containing 5 M guanidine thiocyanate, 50 mM potassium chloride, 10 mM
196　　　Tris/HCl (pH 8); and 0.5 ml Tween 20.

197　ii)　Cut skin and other tissue samples into fine pieces using a sterile scalpel blade and forceps.
198　　　Grind with a pestle in a mortar. Suspend the tissue samples in 800 µl of the above mentioned
199　　　lysis buffer.

200　iii)　Add 2 µl of proteinase K (20 mg/ml) to blood samples and 10 µl of proteinase K (20 mg/ml) to
201　　　tissue samples. Incubate at 56°C for 2 hours or overnight, followed by heating at 100°C for
202　　　10 minutes. Add phenol:chloroform:isoamylalcohol (25:24:1 [v/v]) to the samples in a 1:1 ratio.
203　　　Vortex and incubate at room temperature for 10 minutes. Centrifuge the samples at 16,060 ***g***
204　　　for 15 minutes at 4°C. Carefully collect the upper, aqueous phase (up to 200 µl) and transfer
205　　　into a clean 2.0 ml tube. Add two volumes of ice cold ethanol (100%) and 1/10 volume of 3 M
206　　　sodium acetate (pH 5.3). Place the samples at –20°C for 1 hour. Centrifuge again at 16,060 ***g***
207　　　for 15 minutes at 4°C and discard the supernatant. Wash the pellets with ice cold 70% ethanol
208　　　(100 µl) and centrifuge at 16,060 ***g*** for 1 minute at 4°C. Discard the supernatant and dry the
209　　　pellets thoroughly. Suspend the pellets in 30 µl of nuclease-free water and store immediately at
210　　　–20°C (Tuppurainen *et al.*, 2005). Alternatively a column-based extraction kit may be used.

211　iv)　The primers for this PCR assay were developed from the gene encoding the viral attachment
212　　　protein. The size of the expected amplicon is 192 bp (Ireland & Binepal, 1998). The primers
213　　　have the following gene sequences:

214　　　Forward primer 5'-TCC-GAG-CTC-TTT-CCT-GAT-TTT-TCT-TAC-TAT-3'

215　　　Reverse primer 5'-TAT-GGT-ACC-TAA-ATT-ATA-TAC-GTA-AAT-AAC-3'.

216　v)　DNA amplification is carried out in a final volume of 50 µl containing: 5 µl of 10 × PCR buffer,
217　　　1.5 µl of MgCl$_2$ (50 mM), 1 µl of dNTP (10 mM), 1 µl of forward primer, 1 µl of reverse primer,
218　　　1 µl of DNA template (~10 ng), 0.5 µl of *Taq* DNA polymerase and 39 µl of nuclease-free water.
219　　　The volume of DNA template required may vary and the volume of nuclease-free water must
220　　　be adjusted to the final volume of 50 µl.

221　vi)　Run the samples in a thermal cycler as follows: 2 minutes at 95°C; then 45 seconds at 95°C,
222　　　50 seconds at 50°C and 1 minute at 72°C (34 cycles); 2 minutes at 72°C and hold at 4°C until
223　　　analysis.

224　vii)　Mix 10 µl of each sample with loading dye and load onto a 1.5% agarose gel in TAE buffer
225　　　(Tris/acetate buffer containing EDTA). Load a parallel lane with a 100 bp DNA-marker ladder.
226　　　Electrophoretically separate the products using approximately 8–10 V/cm for 40–60 minutes
227　　　and visualise with a suitable DNA stain and transilluminator.

228　Quantitative real-time PCR methods have been described that are reported to be faster and have
229　higher sensitivity than conventional PCRs (Balinsky *et al.*, 2008; Bowden *et al.*, 2008). A real-time
230　PCR method that differentiates between LSDV, sheep pox virus and goat pox virus has been
231　published (Lamien *et al.*, 2011).

232　Quantitative real-time PCR assays have been designed to differentiate between Neethling-based
233　LSDV strains, which are often used for vaccination, and wild-type LSDV strains from cluster 1.2
234　(Agianniotaki *et al.*, 2017; Pestova *et al.*, 2018; Vidanovic *et al.*, 2016). These "DIVA" assays (DIVA:
235　differentiation 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

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 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 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 buffalo pox, a disease that usually manifests as pock lesions on the teats, but may cause skin lesions 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. 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 striations over the virion. Capripoxvirus virions are also distinct from the herpesvirus that causes pseudo-LSD (also known as "Allerton" or bovine herpes mammillitis).

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).

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

Molecular tests using loop-mediated isothermal amplification to detect capripoxvirus genomes are reported to provide 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).

2. Serological tests

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.

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 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 it is more difficult to read an end-point in tubes.

2.1.1. Test procedure

i) Test sera, including a negative and a positive control, are diluted 1/5 in Eagle's/HEPES (N-2-hydroxyethylpiperazine, 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 $\log_{10} 6$ $TCID_{50}$ per ml is diluted in Eagle's/HEPES in bijoux bottles to give a log dilution series of $\log_{10}$ 5.0, 4.0, 3.5, 3.0, 2.5, 2.0, 1.5 $TCID_{50}$ per ml (equivalent to $\log_{10}$ 3.7, 2.7, 2.2, 1.7, 1.2, 0.7, 0.2 $TCID_{50}$ 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 10^5 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.

x) Antibodies to capripoxvirus can be detected from 1 to 2 days after the onset of clinical signs. These remain detectable for about 7 months.

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 anti-bovine 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.

2.4. Western blot analysis

Western blotting of test sera against capripoxvirus-infected cell lysate provides a sensitive and specific system for 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 the cellular debris pelleted using centrifugation. The supernatant should be decanted, and the proteins should be separated using SDS/PAGE (sodium dodecyl sulphate/polyacrylamide gel electrophoresis). A vertical discontinuous 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 with a glycine running buffer containing Tris (250 mM), glycine (2 M), and SDS (0.1%). Samples of supernatant should be prepared by boiling for 5 minutes with an appropriate lysis buffer prior to loading. Alternatively, purified virus or recombinant antigens may replace tissue-culture-derived antigen.

Molecular weight markers should be run concurrently with the protein samples. The separated proteins in the 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 PBS, on a rotating shaker at 4□C overnight. The NCM can then be separated into strips by employing a commercial apparatus to allow the concurrent testing of multiple serum samples, or may be cut into strips and each strip incubated 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 in blocking buffer (3% BSA and 0.05% Tween 20 in PBS; or 5% milk powder and 0.05% Tween 20 in PBS). The membrane is again thoroughly washed and incubated (in blocking buffer) with anti-species immunoglobulin 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 (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 then undertaken for approximately 3–7 minutes at room temperature on a shaker with constant observation, and the reaction is stopped by washing the NCM in PBS before excessive background colour is seen. A positive and negative 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.

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.~~

~~Capripoxvirus vaccine strains can produce a large local reaction at the site of inoculation in *Bos taurus* breeds (Davies, 1991), which some stock owners find unacceptable. This has discouraged the use of vaccine, even though the consequences of an outbreak of LSD are invariably more severe. Risk–benefit of vaccination should be assessed following stakeholder discussion.~~

388 Vaccines are a key tool to control LSD. Different types of LSD vaccines have been developed and several are commercially
389 available (Tuppurainen *et al.,* 2021).

390 Live attenuated vaccines (LAV) based on the Neethling LSDV strain (homologous LAV vaccines) have been shown to offer
391 high levels of protection against LSD under experimental conditions (Haegeman *et al.,* 2021) and have been used
392 successfully to control the disease in the field, through systematic vaccination of the entire country's cattle population for
393 a number of consecutive years (Klement *et al.,* 2020). Homologous vaccines may induce fever, produce a local reaction
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
396 within a few days and are largely outweighed by the overall benefits of vaccination with homologous vaccines. The duration
397 of immunity induced by good quality live attenuated LSDV vaccines was shown to be at least 18 months (Haegeman *et*
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
401 vaccines usually contain higher doses of virus than when administered to sheep. Although safe, their effectiveness in
402 protecting cattle against LSD is inferior compared to homologous vaccines (Ben-Gera *et al*., 2015; Zhugunissov *et al*.,
403 2020). Heterologous vaccines containing goat pox virus strains for use in cattle against LSD have been developed more
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.

409 In addition, homologous inactivated vaccines against LSD have been developed and tested (Haegeman *et al*., 2023; Hamdi
410 *et al*., 2020; Wolff *et al*., 2022). These vaccines are reported to be safe and efficacious. They however require a booster
411 vaccination one month after primo-vaccination and then every 6 months thereafter, based on the fact that the duration of
412 immunity is shorter than 1 year (Haegeman *et al*., 2023).

413 None of the commercial vaccines currently available has practical DIVA capacity. This problem may be resolved in the
414 future by introducing new types of vaccines (e.g. vector-vaccines, subunit vaccines, mRNA vaccines) that are at various
415 stages of development and evaluation.

2. Outline of production of LSD vaccines ~~and minimum requirements for conventional vaccines~~

416
417

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 The production of vaccines, including LSD vaccines, starts within research and development (R&D) facilities where vaccine
423 candidates are produced and tested in preclinical studies to demonstrate the quality, safety and efficacy of the product.

424 Minimum requirements for different production stages of veterinary vaccines are available in different chapters of the
425 *Terrestrial Manual.* These are intended to be used in combination with country-specific regulatory requirements for vaccine
426 production and release. Here we outline the most important requirements for the production of live and inactivated LSD
427 vaccines. Full requirements are available in Chapter 1.1.8 *Principles of veterinary vaccine production,* Chapter 2.3.3
428 *Minimum requirements for the organisation and management of a vaccine manufacturing facility* and Chapter 2.3.4
429 *Minimum requirements for the production and quality control of vaccine,* and other regulatory documentation.

2.1. Quality assurance

430

431 Facilities for manufacturing LSD vaccines should operate in line with the concepts of good laboratory practice (GLP)
432 and good manufacturing practice (GMP) to produce high quality products. Quality risk management and quality
433 control with adequate documentation management, as an integral part of the production process, have to be in place.
434 In case some activities of the production process are outsourced, those should also be appropriately defined,
435 recorded and controlled.

436 The vaccine production process (Outline of Production) should be documented in a series of standard operating
437 procedures (SOPs), or other documents describing the manufacturing of each batch and the final product (including
438 starting materials to be used, manufacturing steps, in-process controls and controls on the final product). Detailed
439 requirements for documentation management in the process of vaccine production are available in Chapter 2.3.3.

440 A completed Outline of Production is to be enclosed in a vaccine candidate dossier and used for the evaluation of
441 the production process and product by regulatory bodies.

2.2.   Process validation

443 The dossier with the enclosed Outline of Production for the vaccine candidate has to be submitted for regulatory
444 approval, so it can be assessed and authorised by the competent authority to ensure compliance with local regulatory
445 requirements. Among others, data on quality, safety, and efficacy will be assessed. The procedures necessary to
446 obtain these data are described in the subsequent sections.

447 National regulatory authorities might also require official control authority re-testing (check testing) of final products
448 and batches in government laboratories or an independent batch quality control by a third party.

3.   Requirements for LSD vaccine candidates and batch production

3.1.   Requirements for starting materials

451 Live attenuated vaccines (LAV) and inactivated vaccines (IV) for LSD are produced using the system of limited and
452 controlled passages of master seed and working seed virus and cell banks with a specified maximum. This approach
453 aims to prevent possible and unwanted drift of properties of seed virus and cells that might arise from repeated
454 passaging.

3.1.1.   Characteristics of the seed virus

456 Each seed strain of capripoxvirus used for vaccine production must be accompanied by records
457 clearly and accurately describing its origin, isolation and tissue culture or animal passage history.
458 Preferably, the species and strain of capripoxvirus are characterised using PCR or DNA sequencing
459 techniques.

460 A quantity of master seed vaccine virus should be prepared, frozen or desiccated and stored at low
461 temperatures such as −80°C and used to produce a consistent working seed for regular vaccine
462 production.

463 Each master seed strain must be non-transmissible, remain attenuated after further tissue culture
464 passage, and provide complete protection against challenge with virulent field strains for a minimum
465 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*.

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

468 Each master seed must be tested to ensure its identity and shown to be free from adventitious
469 viruses, in particular pestiviruses, such as border disease and bovine viral diarrhoea virus, and free
470 from contamination with bacteria, fungi or mycoplasmas.

471 The general procedures for sterility or purity tests are described in Chapter 1.1.9 *Tests for sterility*
472 *and freedom from contamination of biological materials intended for veterinary use*.

473 Master seed virus is a quantity of virus of uniform composition derived from an original isolate,
474 passaged for a documented number of times and distributed into containers at one time and stored
475 adequately to ensure stability (via freezing or lyophilisation). Selection of master seed viruses (MSVs)
476 should ideally be based on their ease of growth in cell culture, virus yield, and in accordance with the
477 regional epidemiological importance. Also, measures to minimise transmissible spongiform
478 encephalopathies (TSE) contamination should be taken into account (see Section C.3.5.1 *Purity*
479 *tests*).

480 For each seed strain selected for LSD vaccine production, the following information should be
481 provided:

482 -   Historical record: geographical origin, animal species from which the virus was recovered,
483 isolation procedure, tissue culture or animal passage history

484 -   Identity: species and strain identification using DNA sequencing

| 485 | - | Purity: the absence of bacteria, fungi, mycoplasma, and other viruses (see Chapter 1.1.9 *Tests* |
| 486 | | *for sterility and freedom from contamination of biological materials intended for veterinary use*) |

| 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 - Stability

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 limited) cell culture passages from the master seed stock and used to produce vaccine batches. This
498 approach and limitation of seed virus passaging will assist in maintaining uniformity and consistency
499 in production.

3.1.2. 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 has an inherently higher risk of introducing extraneous agents compared with the use of established
505 (well characterised) cell lines and should be avoided where alternative methods of producing effective
506 vaccines exist. For each MCS, manufacturers should demonstrate:

507 - MCS identity

508 - genetic stability by subculturing from the lowest to the highest passage used for production

509 - stable MCS karyotype with a low level of polyploidy

510 - freedom from oncogenicity or tumorigenicity by using *in-vivo* studies using the highest cell
511   passage that may be used for production

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*).

3.2. Method of <u>vaccine</u> manufactur<u>ing</u>

515 ~~The method of manufacture should be documented as the Outline of Production.~~

~~2.2.1. Procedure~~

3.2.1. LSD vaccine batch production

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 <u>production of LAV and IV against</u>
521 <u>LSD starts with the inoculation of the</u> required number of <u>working</u> vials of seed virus ~~is~~ reconstituted
522 ~~with GMEM or other~~ <u>in</u> appropriate medium ~~and inoculated~~ onto a <u>suitable primary or continuous cell</u>
523 <u>line grown in suspension or</u> monolayer. ~~Cells should be harvested after 4–8 days when they exhibit~~
524 ~~50–70% CPE for maximum~~ <u>in the exponential growth phase. At the time highest</u> viral ~~infectivity, or~~
525 ~~earlier if CPE is extensive and cells appear ready to detach.~~ Techniques such as <u>loads are present,</u>
526 sonication or repeated freeze–thawing ~~are~~ <u>is</u> used to release the intracellular virus from the
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. <u>For LAV, the virus-containing</u>
531 suspension <u>is diluted to attain the dose at which the vaccine candidate will be evaluated or to at least</u>
532 <u>the determined protective dose for approved vaccines and is</u> then mixed with a suitable protectant
533 ~~such as an equal volume of sterile, chilled 5% lactalbumin hydrolysate and 10% sucrose (dissolved~~

534		in double-distilled water or appropriate balanced salt solution), and transferred to individually
535		~~numbered~~ labelled bottles or bags for storage at low temperatures such as –80°C, or for freeze–
536		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		~~The specification and source of all ingredients used in the manufacturing procedure should be~~
539		~~documented and the freedom of extraneous agents (bacteria, fungi, mycoplasma and viruses) should~~
540		~~be tested. The detailed testing procedure is described in Chapter 1.1.9. The use of antibiotics must~~
541		~~meet the requirements of the licensing authority.~~

542	~~2.2.3.~~	~~In-process control~~

543	~~i)~~	~~Cells~~

544	~~ii)~~	~~Records of the source of the master cell stocks should be maintained. The highest and lowest~~
545		~~passage numbers of the cells that can be used for vaccine production must be indicated in the~~
546		~~Outline of the Production. The use of a continuous cell line (such as MDBK, etc.) is strongly~~
547		~~recommended, unless the virus strain only grows on primary cells. The key advantage of~~
548		~~continuous over primary cell lines is that there is less risk of introduction of extraneous agents.~~

549	~~iii)~~	~~Serum~~

550	~~iv)~~	~~Serum used in the growth or maintenance medium must be free from antibodies to capripoxvirus~~
551		~~and free from contamination with pestivirus or other viruses, extraneous bacteria, mycoplasma~~
552		~~or fungi.~~

553	~~v)~~	~~Medium~~

554	~~vi)~~	~~Media must be sterile before use.~~

555	~~vii)~~	~~Virus~~

556	~~viii)~~	~~Seed virus and final vaccine must be titrated and pass the minimum release titre set by the~~
557		~~manufacturer. For example, the minimum recommended field dose of the South African~~
558		~~Neethling strain vaccines (Mathijs et al., 2016) is log_{10} 3.5 TCID_{50}, although the minimum~~
559		~~protective dose is log_{10} 2.0 TCID_{50}. Capripoxvirus is highly susceptible to inactivation by sunlight~~
560		~~and allowance should be made for loss of activity in the field.~~

561	~~ix)~~	~~The recommended field dose of the Romanian sheep pox vaccine for cattle is log_{10} 2.5 sheep~~
562		~~infective doses (SID_{50}), and the recommended dose for cattle of the RM65 adapted strain of~~
563		~~Romanian sheep pox vaccine is log_{10} 3 TCID_{50} (Coakley & Capstick, 1961).~~

564	**3.2.2.**	**Inactivation process for inactivated LSD vaccines**

565		Unlike LAV, inactivated LSD vaccines contain inactivated antigens in combination with adjuvants to
566		strengthen the induced immune response after administration. The vaccine evaluation process
567		described below needs to show the amount of antigen necessary to elicit a protective immune
568		response. Currently, literature data indicate that an inactivated vaccine originating from an LSDV
569		virus stock with titre 10^4 cell culture infectious dose_{50} (CCID_{50})/ml before inactivation can be sufficient
570		to induce an efficient immune response to prevent clinical disease, viremia and virus shedding after
571		challenge of young cattle (Wolf *et al.,* 2022)

572		To monitor the inactivation process and the level of antigen inactivation, samples are taken at regular
573		intervals during inactivation and titrated. Inactivation conditions and the length of initial and repeated
574		exposure should be documented in detail since one or more factors during the process could
575		influence the outcomes. The inactivation kinetics should reach a predefined target e.g. one remaining
576		infectious unit per million doses (1×10^{-6} infectious units/dose) as suggested by APHIS (2013). The
577		confirmatory testing of inactivation is performed on each vaccine lot and represents an important part
578		of the inactivation process monitoring. In addition to all the procedures mentioned above, the
579		inactivation procedure and tests demonstrating that antigen inactivation is complete and consistent
580		must additionally be documented in the Outline of Production.

3.3. Vaccine safety

During the vaccine development process, vaccine safety must be evaluated in the target animal (target animal batch safety test –TABST) to demonstrate the safety of the dose intended for registration. The animals used in the safety testing should be representative (species, age and category [calves, heifers, bulls, cows.]) for all the animals for which 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 moribund animals.

Essential parameters to be evaluated in safety studies are local and systemic reactions to vaccination, including local reactions at the site of administration, fever, effect on milk production, and induction of a 'Neethling' response. The effect of the vaccine on reproduction needs to be evaluated where applicable.

A part of the safety evaluation of LAV and IV can be performed during the efficacy trials (see Section C.3.4 *Vaccine 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 LAV and IV (EMEA, 2009). Safety aspects of LAV and IV against LSD to be evaluated are:

3.3.1. Overdose test for LAV

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).

3.3.2. 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.

3.3.3. Reversion 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. 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~~

ii) ~~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.~~

iii) ~~Safety and efficacy~~

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.~~

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~~
663 ~~evidence of protection.~~

3.4.   Vaccine efficacy

665 Data enclosed in the vaccine candidate dossier should support the efficacy of the vaccine in each animal species for
666 each vaccination regimen that is described in the product label recommendation. This includes studies regarding the
667 onset of protection when claims for onset are made and for the duration of immunity. Efficacy studies should be
668 conducted with the vaccine candidate that has been produced at the highest passage level permitted for vaccine
669 production as specified in the Outline of Production.

670 Efficacy (and safety) should be demonstrated in vaccination–challenge studies using representative (by species, age
671 and category) seronegative healthy animals for which the vaccine is intended and which are tested negative for
672 standard viral pathogens.

673 An example of a vaccination-challenge test set-up is outlined here. The group numbers mentioned can be varied if
674 statistically justified. Thirteen animals are placed in a high containment large animal unit and are divided into two
675 groups:

676 -   single/repeated dose test group (*n*=8) – animals inoculated with the vaccine dose and route intended for
677   registration (in case of an IV against LSD, a booster dose should follow primary vaccination after minimum 14
678   days).

679 -   control group (*n*=5) – non-vaccinated animals

680 Throughout the *in-vivo* study, all animals are clinically examined and rectal temperatures recorded. Blood, serum and
681 swab samples are regularly collected and subjected to laboratory testing. On day 21 after the vaccination with a LAV
682 or after the booster vaccination for an IV, the animals in both groups are challenged with a known virulent LSDV

683 strain. The challenge virus solution should be of known titre and tested free from extraneous viruses. Experience
684 obtained from previous animal experiments indicates that a dose of challenge virus between $10^{4.0}$ and $10^{6.5}$ TCID$_{50}$
685 produces clinical disease in about half of the susceptible experimental cattle (;.Tuppurainen *et al.*, 2021).

686 The clinical response following challenge is recorded over a period of 14 days. No clinical signs should occur in the
687 vaccinates, other than a local reaction at the site of inoculation. At least 1 animal in the unvaccinated control group
688 should develop the typical clinical signs of LSD. Although a generalised disease with skin nodules may not be seen
689 in all the unvaccinated control animals based on the knowledge that the outcome of a LSDV infection can range from
690 inapparent to severe, at the very least a large local reaction is to be expected.

691 Clinical and laboratory results will enable assessment of the safety and efficacy of the LSD vaccine candidate and
692 the induced immune responses. Serum samples collected at different time points during the trial can be examined to
693 study seroconversion against selected viral diseases that could have contaminated the vaccine.

694 ## 2.3. Requirements for regulatory approval

695 ### 2.3.1. Safety requirements

696   i)   Target and non-target animal safety

697   ii)   The vaccine must be safe to use in all breeds of cattle for which it is intended, including young
698   and pregnant animals. It must also be non-transmissible and remain attenuated after further
699   tissue culture passage.

700   iii)   Safety tests should be carried out on the final product of each batch as described in Section
701   C.2.2.4.

702   iv)   Reversion to virulence for attenuated/live vaccines

703   v)   The selected final vaccine should not revert to virulence during further passages in target
704   animals.

705   vi)   Environmental consideration

706   vii)   Attenuated vaccine should not be able to perpetuate autonomously in a cattle population.
707   Strains of LSDV are not a hazard to human health.

708 ### 2.3.2. Efficacy requirements

709   i)   For animal production

710   The efficacy of the vaccine must be demonstrated in statistically valid vaccination challenge
711   experiments under laboratory conditions. The group numbers recommended here can be varied if
712   statistically justified. Fifteen cattle are placed in a high containment level large animal unit and serum
713   samples are collected. Five randomly chosen vials of the freeze-dried vaccine are reconstituted in
714   sterile PBS and pooled. Two cattle are inoculated with 10 times the field dose of the vaccine, eight
715   cattle are inoculated with the recommended field dose. The remaining five cattle are unvaccinated
716   control animals. The animals are clinically examined daily and rectal temperatures are recorded. On
717   day 21 after vaccination, the animals are again serum sampled and challenged with a known virulent
718   capripoxvirus strain using intravenous and intradermal inoculation (the challenge virus solution
719   should also be tested and shown to be free from extraneous viruses). The clinical response is
720   recorded during the following 14 days. Animals in the unvaccinated control group should develop the
721   typical clinical signs of LSD, whereas there should be no local or systemic reaction in the vaccinates
722   other than a raised area in the skin at the site of vaccination which should disappear after 4 days.
723   Serum samples are again collected on day 30 after vaccination. The day 21 serum samples are
724   examined for seroconversion to selected viral diseases that could have contaminated the vaccine,
725   and the days 0 and 30 samples are compared to confirm the absence of antibody to pestivirus.
726   Because of the variable response in cattle to challenge with LSDV, generalised disease may not be
727   seen in all of the unvaccinated control animals, although there should be a large local reaction.

728   Once the potency of the particular strain being used for vaccine production has been determined in
729   terms of minimum dose required to provide immunity, it is not necessary to repeat this on the final
730   product of each batch, provided the titre of virus present has been ascertained.

731   ii)   For control and eradication

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.

2.3.3.   Stability

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.

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.

3.5.1.   Purity test

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 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

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).

3.5.3.   Potency tests

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.

3.5.3.1.   Live vaccines

774   The potency of LAV against LSD can be measured by means of virus titration. The virus titre
775   must, as a rule, be sufficiently greater than that shown to be protective in the efficacy test
776   for the vaccine candidate. This will ensure that at any time prior to the expiry date, the titre
777   will be at least equal to the evaluated protective titre. The titres of currently available

	commercial homologous LSD vaccines range between 10^3 and 10^4 infectious units/dose
778	commercial homologous LSD vaccines range between 10^3 and 10^4 infectious units/dose
779	(Tuppurainen *et al.*, 2021).

3.5.3.2. Inactivated LSD vaccines

For inactivated LSD vaccines, potency tests are performed using vaccination–challenge efficacy studies in animal hosts (see Section C.3.4. *Vaccine efficacy*).

3.5.4. Safety/efficacy

Safety and efficacy testing is undertaken during the evaluation process of the vaccine candidate, and also needs to be performed on a number of vaccine batches until robust data are generated in line with international and national regulations. Afterwards, when using a seed lot system in combination with strict implementation of GMP standards and depending on local regulations, TABST could be waived as described in VICH50 and VICH55, providing the titer has been ascertained using potency testing. Batches or serials are considered satisfactory if local and systemic reactions to vaccination are in line with those described in the dossier of the vaccine candidate and product literature.

3.5.4.1. Field safety/efficacy tests

Field testing of two or more batches should be performed on all animal categories for which the product is indicated before release of the product for general use (see chapter 1.1.8). The aim of these studies is to demonstrate the safety and efficacy of the product under normal field conditions of animal care and use in different geographical locations where different factors may influence product performance. A protocol for safety/efficacy testing in the field has to be developed with defined observation and recording procedures. However, it is generally more difficult to obtain statistically significant data to demonstrate efficacy under field conditions. Even when properly designed, field efficacy studies may be inconclusive due to uncontrollable outside influences.

3.5.4.2. Duration of Immunity

The duration of immunity (DOI) following vaccination should be demonstrated via challenge or the use of a validated serology test. Efficacy testing at the end of the claimed period of protection should be conducted in each species for which the vaccine is indicated or the manufacturer should indicate that the DOI for that species is not known. Likewise, the manufacturer should demonstrate the effectiveness of the recommended booster regime in line with these guidelines, usually by measuring the magnitude and kinetics of the serological response observed.

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).~~

4. Post-market studies

4.1. Stability

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.

4.2. Post-marketing surveillance

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.

827   First, a reliable reporting system should be in place to collect consumer complaints and notifications of adverse
828   reactions. Secondly, post-marketing surveillance should be established to investigate whether the reported
829   observations are related to the use of the product and to identify, at the earliest stage, any serious problem that may
830   be encountered from its use and that may affect its future uptake. Vaccinovigilance should be an on-going and integral
831   part of all regulatory programmes for LSD vaccines, especially for live vaccines.

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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
986 diagnostic tests, reagents and vaccines for lumpy skin disease

987 **NB:** FIRST ADOPTED IN 1989. MOST RECENT UPDATES ADOPTED IN 2021.

2   **MEETING OF THE BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5

6   # C H A P T E R   3 . 6 . 9 .

7   # EQUINE RHINOPNEUMONITIS (INFECTION WITH
8   # EQUID HERPESVIRUS-1 ~~AND -4~~)

9   **SUMMARY**

10   *Equine rhinopneumonitis (ER) is a collective term for any one of several contagious, clinical disease entities*
11   *of equids that may occur as a result of infection by either of two closely related herpesviruses, equid*
12   *<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</u>*
13   *<u>focus of this chapter.</u> EHV-1 <u>is</u> ~~and EHV-4 are~~ endemic in most domestic equine populations worldwide.*

14   *Primary infection by ~~either~~ EHV-1 ~~or EHV-4~~ is characterised by upper respiratory tract disease of varying*
15   *severity that is related to the age and immunological status of the infected animal. <u>Following viraemia</u> EHV-*
16   *1 also causes the more serious complications of abortion, perinatal foal death, or paralytic neurological*
17   *disease (equine herpesvirus myeloencephalopathy). ~~EHV-4 has been associated with sporadic cases of~~*
18   *~~abortion, but rarely multiple abortions and not the large outbreaks associated with EHV-1.~~ Like other*
19   *herpesviruses, EHV-1 ~~and 4~~ induce<u>s</u> long-lasting latent infections and can be reactivated following stress ~~or~~*
20   *~~pregnancy~~. Furthermore, most horses are likely to be re-infected multiple times during their lifetime, often*
21   *~~mildly or~~ subclinically. ~~Detection of viral DNA or anti-EHV antibodies should therefore be interpreted with~~*
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~~.*

26   *Positive identification of viral isolates as EHV-1 ~~or EHV-4~~ can be achieved by type-specific PCR <u>or</u>*
27   *<u>sequencing</u>. Viruses can be isolated in ~~equine~~ cell culture from nasal or nasopharyngeal swab extracts taken*
28   *from horses ~~during the febrile stage of~~ <u>with acute</u> respiratory tract infection, from the placenta<u>,</u> <u>from</u> ~~and~~ liver,*
29   *lung, spleen, <u>adrenal glands</u> or thymus of aborted fetuses and early foal deaths, and from the leukocyte*
30   *fraction of the blood of animals ~~with acute~~ <u>during the febrile stage of</u> EHV-1 infection. ~~Unlike EHV-4, EHV-1~~*
31   *~~will also grow in various non-equine cell types such as the RK-13 cell line and this property can be used to~~*
32   *~~distinguish between the two viruses.~~*

33   *A rapid presumptive diagnosis of abortion induced by EHV-1 ~~or (infrequently) EHV-4~~ can be achieved by*
34   *direct immunofluorescent detection of viral antigen in cryostat sections of placenta and tissues from aborted*
35   *fetuses, using a conjugated polyclonal antiserum.*

36   *Post-mortem demonstration of histopathological lesions of EHV-1 in placenta and tissues from aborted*
37   *fetuses, cases of perinatal foal death or in the central nervous system of neurologically affected animals*
38   *complements <u>other diagnostic techniques</u> ~~the laboratory diagnosis~~.*

39   ***Serological tests:*** *Most horses possess some level of antibody to EHV-1/4, the demonstration of specific*
40   *antibody in the serum collected from a single blood sample is therefore not confirmation of a positive*
41   *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.*

A. INTRODUCTION

60   Equine rhinopneumonitis (ER) is a historically-derived term that describes a constellation of several disease entities of
61   horses that may include respiratory disease, abortion, neonatal foal pneumonitis, or myeloencephalopathy (Allen & Bryans,
62   1986; Allen *et al*., 1999; Bryans & Allen, 1988; Crabb & Studdert, 1995). The disease has been is recognised for over 60
63   years as a threat to the international horse industry, and is caused by either of two members of the *Herpesviridae* family,
64   equid alphaherpesvirus-1 and -4 (EHV-1 and EHV-4). EHV-1 and EHV-4 are closely related alphaherpesviruses of horses
65   with nucleotide sequence identity within individual homologous genes ranging from 55% to 84%, and amino acid sequence
66   identity from 55% to 96% (Telford *et al*., 1992; 1998). The two herpesviruses With the exception of EHV-1 in Iceland
67   (Thorsteinsdóttir *et al*., 2021), the two herpesviruses are considered endemic enzootic in all countries in which large
68   populations of horses are maintained as part of the cultural tradition or agricultural economy. There is no recorded evidence
69   that the two herpesviruses of ER pose any health risks to humans working with the agents. Infection with EHV-1 is listed
70   by WOAH and is therefore the focus of this chapter.

71   Viral transmission to cohort animals occurs by inhalation of aerosols of virus-laden respiratory secretions. Morbidity tends
72   to be highest in young horses sharing the same air space. Aborted tissues and placental fluids from infected mares can
73   contain extremely high levels of live virus and represent a major source of infection. Extensive use of vaccines has not
74   eliminated EHV-1 infections, and the world-wide annual financial impact from this these equine pathogens is immense
75   considerable.

76   In horses under 3 years of age, clinical ER usually takes the form of an acute, febrile respiratory illness that spreads rapidly
77   through the group of animals. The viruses infects and multiplies multiply in epithelial cells of the respiratory mucosa. Signs
78   of infection become apparent 2–8 days after exposure to virus, and are characterised by fever, inappetence, depression,
79   and nasal discharge. The severity of respiratory disease varies with the age of the horse and the level of immunity resulting
80   from previous vaccination or natural exposure. Bi-phasic fever, viraemia and complications are more likely with EHV-1 than
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
83   cause of interrupted schedules among horses assembled for training, racing, or other equestrian events. Fully protective
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
86   disease, the risks of subsequent abortion or neurological disease remain. Like other herpesviruses, EHV-1/4 causes long-
87   lasting latent infections and latently infected horses represent a potential infection risk for other horses. Virus can be
88   reactivated as a result of stress or pregnancy. The greatest clinical threats to individual breeding, racing, or pleasure horse
89   operations posed by ER are the potential abortigenic and neurological *sequelae* of EHV-1 respiratory infection. ER
90   abortions occur annually in horse populations worldwide and may be sporadic or multiple. Foals infected *in utero* may be
91   born alive and die within a few days of birth. EHV-1 neurological disease is less common than abortions but has been
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*.,
97   2007; Nugent *et al*., 2006). Strain typing techniques have been employed to identify viruses carrying the neuropathic

98 ~~marker, and it can be helpful to be aware of an increased risk of neurological complications. However, for practical purposes~~
99 ~~strain typing is not relevant for agent identification, or international trade. Strain typing may be beneficial for implementation~~
100 ~~of biosecurity measures in the management of outbreaks of equine herpesvirus myeloencephalopathy.~~

101 Strain typing has been shown to be not reliable for predicting the clinical outcome of EHV-1 infection but can be useful in
102 epidemiological investigations (Garvey *et al.*, 2019; Nugent *et al.*, 2006; Sutton *et al.*, 2019).

103 # B. DIAGNOSTIC TECHNIQUES

104 ~~Both~~ EHV-1 ~~and EHV-4~~ is transmitted by the respiratory route and has ~~have~~ the potential to be highly contagious ~~viruses~~
105 particularly where large numbers of horses are housed in the same air space. EHV1 ~~and the former~~ can cause explosive
106 outbreaks of abortion or neurological disease. Rapid diagnostic methods are therefore essential ~~useful~~ for managing the
107 disease. Real-time polymerase chain reaction (PCR) assays are ~~widely~~ routinely used by diagnostic laboratories worldwide
108 and are both rapid and sensitive. Real-time PCR assays that allow ~~simultaneous~~ testing for EHV-1 ~~and EHV 4~~ and
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~~
111 ~~true of~~ for EHV-1 associated abortions and neonatal foal deaths, when the high level of virus in the tissues usually produces
112 a cytopathic effect in 1–3 days. Immunohistochemical or immunofluorescent approaches are employed in some
113 laboratories ~~can be extremely useful~~ for rapid diagnosis of EHV-induced abortion from fresh or embedded tissue ~~and are~~
114 ~~relatively straightforward. Several other techniques based on enzyme-linked immunosorbent assay (ELISA) or nucleic acid~~
115 ~~hybridisation probes have also been described, however their use is often restricted to specialised laboratories and they~~
116 ~~are not included here.~~ Virus neutralisation (VN) and complement fixation (CF) are the most frequently used serological
117 tests, and seroconversion in paired samples is considered indicative of exposure to virus by natural infection or by
118 vaccination.

119 *Table 1.* Test methods available for the diagnosis of equine rhinopneumonitis and their purpose

Method	Purpose					
	Population freedom from infection	Individual animal freedom from infection prior to movement	Contribute to eradication policies[a]	Confirmation of clinical cases	Prevalence of infection - surveillance	Immune status in individual animals or populations post-vaccination
Identification of the agent[b]						
Virus isolation	–	++ ~~+~~	–	+++	–	–
PCR	–	+++	–	+++	–	–
Direct immunofluorescence	=	=	=	++	=	=
Detection of immune response						
VN	~~+~~ ±	~~+~~ ±	= ~~+~~	+++	+++	+++
ELISA	+	= ~~+~~	= ~~+~~	+ ~~+~~	++ ~~+~~	+
CFT	–	~~–~~ ++	–	+++	–	~~–~~ +++

120 Key: +++ = recommended for this purpose; ++ recommended but has limitations;
121 + = suitable in very limited circumstances; – = not appropriate for this purpose.
122 PCR = polymerase chain reaction; VN = virus neutralisation;
123 ELISA = enzyme-linked immunosorbent assay; CFT = complement fixation test.
124 [a]No eradication policies exist for equine rhinopneumonitis.
125 [b]A combination of agent identification methods applied on the same clinical sample is recommended.

1. ~~Identification~~ <u>Detection</u> **of the agent**

1.1. Collection and preparation of specimens

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 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~~ <u>acute stage</u> of the respiratory disease, and are collected via the nares by sampling the area with a swab of an appropriate size and length for horses. After collection, the swab should be 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 infectivity can be prolonged by the addition of bovine serum albumin, fetal calf serum or gelatine to 0.1% (w/v).

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 EHV-1 abortion is most successful when performed on aseptically collected samples of placenta, liver, lung, thymus<u>,</u> <u>adrenal glands</u> and spleen. Virus may be isolated from post-mortem cases of EHV-1 neurological disease by culture of samples of brain and spinal cord but such attempts to isolate virus are often unsuccessful; however, they may be 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 at –70°C.

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.

<u>*Cerebrospinal fluid:* the detection of EHV-1 DNA in cerebrospinal fluid has been reported in cases of neurological disease.</u>

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-embedded archival tissue, or inoculated cell cultures (Borchers & Slater, 1993; Lawrence *et al.*, 1994; O'Keefe *et al.*, 1994; Varrasso *et al.*, 2001). A variety of type-specific PCR primers have been designed to distinguish between the 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 of infectious virus in the clinical sample. ~~For diagnosis of active infection by EHV,~~ PCR methods are <u>routinely used to detect EHV-1 DNA in nasopharyngeal swabs and tissue samples</u> ~~most reliable with tissue samples from aborted fetuses and placental tissue or from nasopharyngeal swabs of foals and yearlings~~. They are particularly useful in explosive outbreaks of abortion, respiratory or neurological disease in which ~~a~~ rapid identification <u>and monitoring</u> of the virus <u>spread</u> 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 diagnosis on a horse with neurological signs<u> (Pronost *et al.*, 2012)</u>.

~~Several PCR assays have been published. A nested PCR procedure can be used to distinguish between EHV-1 and EHV-4. A sensitive protocol suitable for clinical or pathological specimens (nasal secretions, blood leukocytes, brain and spinal cord, fetal tissues, etc.) has been described by Borchers & Slater (1993). However, nested PCR methods have a high risk of laboratory cross-contamination, and sensitive rapid one-step PCR tests to detect EHV-1 and EHV-4 (e.g. Lawrence *et al.*, 1994) are preferred. The WOAH Reference Laboratories use quantitative real-time PCR assays such as those targeting heterologous sequences of major glycoprotein genes to distinguish between EHV-1 and EHV-4. A multiplex real-time PCR targeting glycoprotein B gene of EHV-1 and EHV-4 was described by Diallo *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 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 employ in-house methods for strain typing, however these protocols have not yet been validated between different laboratories at an international level.~~

Real-time (or quantitative) PCR has become the method of choice for ~~many~~ <u>the majority of</u> diagnostic ~~tests~~ <u>laboratories</u> and provides rapid and sensitive detection of viral DNA. Equine post-mortem tissues from newborn and adult animals or equine fetal tissue from abortions (tissues containing lung, liver, spleen, thymus, adrenal gland and placental tissues) can be used. For respiratory samples, equine nasopharyngeal swabs or deep nasal swabs

| 180
181 | (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. |

<u>There is no internationally standardised real-time PCR method for EHV-1 but Table 2 summarises the primer and probe sequences for some of the most widely used assays. Type-specific PCR primers have been designed to distinguish between the presence of EHV-1 and EHV-4. The optimised thermocycler times and temperatures are documented in the publications cited.</u>

Table 2. *Primer and probe sequences for EHV1/4 detection by real-time PCR*

Primer	Primer sequence (5' to 3')	Target	Reference
<u>Forward</u>	<u>CAT-GTC-AAC-GCA-CTC-CCA</u>		
<u>Reverse</u>	<u>GGG-TCG-GGC-GTT-TCT-GT</u>	<u>EHV-1 gB</u>	<u>Diallo *et al.,* 2006</u>
<u>Probe</u>	FAM-CCC-TAC-GCT-GCT-CC-MGB-NFQ		
<u>Forward</u>	<u>CAT-ACG-TCC-CTG-TCC-GAC-AGA-T</u>		
<u>Reverse</u>	<u>GGTACTCGGCCTTTGACGAA</u>	<u>EHV-1 gB</u>	<u>Hussey *et al.,* 2006</u>
<u>Probe</u>	FAM-TGA-GAC-CGA-AGA-TCT-CCT-CCA-CCG-A-BHQ1		
<u>Forward</u>	<u>TAT-ACT-CGC-TGA-GGA-TGG-AGA-CTT-T</u>		
Reverse	TTG-GGG-CAA-GTT-CTA-GGT-GGT-T	<u>EHV-1 gB</u>	<u>Pusterla *et al.,* 2009</u>
Probe	<u>6FAM-ACA-CCT-GCC-CAC-CGC-CTA-CCG</u>		
<u>Forward</u>	GCG-GGC-TCT-GAC-AAC-ACA-A		ISO 17025 accredited for the detection of EHV-1 at WOAH Reference Laboratory
<u>Reverse</u>	TTG-TGG-TTT-CAT-GGG-AGT-GTG-TA	<u>EHV-1 gC</u>	
<u>Probe</u>	<u>FAM-TAA-CGC-AAA-CGG-TAC-AGA-A-BHQ1</u>		

| 187
188
189
190
191
192
193
194 | ~~*This multiplex real-time PCR test has been validated to ISO 17025 and is designed for use in a 96-well format. This 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.* (2006) and Lawrence *et al.* (1994). To establish such a real-time PCR assay for diagnostic purposes, validation 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 Laboratories.~~ Reference material and sample panels <u>for real-time PCR</u> can be obtained from the WOAH Reference Laboratories. |

- **Point of care (POC) molecular tests**

<u>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.</u>

- **Molecular characterisation**

<u>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).</u>

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 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~~
214 ~~dermal (E Derm) or lung tissue. EHV-1 can be isolated on other cell types, as will be discussed later. The~~
215 ~~nasopharyngeal swab and its accompanying transport medium are transferred into the barrel of a sterile 10 ml~~
216 ~~syringe. Using the syringe plunger, the fluid is squeezed from the swab into a sterile tube. A portion of the expressed~~
217 ~~fluid can be filtered through a sterile, 0.45 µm membrane syringe filter unit into a second sterile tube if heavy bacterial~~
218 ~~contamination is expected, but this may also lower virus titre. Recently prepared cell monolayers in tissue culture~~
219 ~~flasks are inoculated with the filtered, as well as the unfiltered, nasopharyngeal swab extract. Cell monolayers in~~
220 ~~multiwell plates incubated in a 5% CO2 environment may also be used. Virus is allowed to attach by incubating the~~
221 ~~inoculated monolayers at 37°C. Monolayers of uninoculated control cells should be incubated in parallel.~~

222 ~~At~~ Recently prepared cell monolayers in tissue culture flasks or plates are inoculated with nasopharyngeal swab
223 extract or homogenised tissue: approximately 10% (w/v) pooled tissue homogenates of liver, lung, thymus, adrenal
224 and spleen (from aborted fetuses/neonatal foals) or of brain and spinal cord (from cases of neurological disease).
225 Virus is allowed to attach by incubating ~~the end of the attachment period,~~ inoculated monolayers at 37°C for 1 hour
226 after which the inocula are removed and the monolayers are rinsed twice with PBS ~~to remove virus-neutralising~~
227 ~~antibody that may~~ or maintenance medium. Monolayers of uninoculated control cells should be ~~present in the~~
228 ~~nasopharyngeal secretions~~ incubated in parallel. After addition of supplemented maintenance medium (MEM
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% CO_2 environment.

231 The use of a positive control ~~virus samples~~ of relatively low titre may be used to validate the isolation procedure
232 ~~carries the risk that this may lead~~ but should be processed separately to ~~eventual~~ avoid contamination of diagnostic
233 specimens. ~~This risk can be minimised by using routine precautions and good laboratory technique, including the use~~
234 ~~of biosafety cabinets, inoculating positive controls after the diagnostic specimens, decontaminating the surfaces in~~
235 ~~the hood while the inoculum is adsorbing and using a positive control of relatively low titre.~~ Inoculated flasks should
236 be inspected daily by microscopy for the appearance of characteristic herpesvirus cytopathic effect (CPE) (focal
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
239 and cells as the inoculum. Further blind passage is usually not productive.

240 ~~It can be useful to inoculate samples into both non-equine and equine cells in parallel to distinguish between EHV-1~~
241 ~~and EHV-4, since EHV-4 can cause sporadic cases of abortion. Around 10% (w/v) pooled tissue homogenates of~~
242 ~~liver, lung, thymus, and spleen (from aborted fetuses) or of central nervous system tissue (from cases of neurological~~
243 ~~disease) are used for virus isolation. These are prepared by first mincing small samples of tissue into 1 mm cubes in~~
244 ~~a sterile Petri dish with dissecting scissors, followed by macerating the tissue cubes further in serum free culture~~
245 ~~medium with antibiotics using a homogeniser or mechanical tissue grinder. After centrifugation at 1200 **g** for~~
246 ~~10 minutes, the supernatant is removed and 0.5 ml is inoculated into duplicate cell monolayers in tissue culture flasks.~~
247 ~~Following incubation of the inoculated cells at 37°C for 1.5–2 hours, the inocula are removed and the monolayers are~~
248 ~~rinsed twice with PBS or maintenance medium. After addition of 5 ml of supplemented maintenance medium, the~~
249 ~~flasks are incubated at 37°C for up to 1 week or until viral CPE is observed. Cultures exhibiting no evidence of viral~~
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
253 unclotted (heparinised) blood by centrifugation at ~~600~~ 525 **g** for ~~15~~ 5 minutes~~, and.~~ The buffy coat is taken after the
254 plasma has been carefully removed. ~~The buffy coat is then layered onto a PBMC separating solution (Ficoll; density~~
255 ~~1077 g/ml, commercially available) and centrifuged at 400 **g** for 20 minutes. The PBMC interface (without most~~
256 ~~granulocytes) is~~ and washed ~~twice in PBS (300 **g** for 10 minutes) and resuspended in 1 ml~~ three times in 3 ml MEM
257 containing 2% FCS. ~~As a quicker alternative method, PBMC may be collected by centrifugation directly from plasma.~~
258 (525 **g** for 5 minutes). Following the third wash, the buffy coat is harvested and resuspended in 2.5 ml MEM containing
259 2% FCS. An aliquot of the rinsed cell suspension ~~is added to each of the duplicate monolayers of equine fibroblast,~~
260 ~~equine fetal or RK-13 cell monolayers in 25 cm² flasks containing 8–10 ml freshly added maintenance medium. The~~
261 ~~flasks~~ can be used for DNA extraction. For virus isolation, the resuspended cells (1 ml) are co-cultivated with freshly
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% CO_2 environment for 3 days or until the cells have reached 90%
264 confluence. The monolayers are then rinsed three times with 1 × PBS and supplemented with 5 ml MEM containing
265 2% FCS. They are incubated ~~at 37°C for 7 days; either with or without removal of the inoculum. If PBMCs are not~~
266 ~~removed prior to incubation, CPE may be difficult to detect in the presence of the massive inoculum of leukocytes:~~
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~~
269 ~~monolayers that are just subconfluent. These are incubated~~ and observed daily for viral CPE ~~for at least 5–6 days.~~
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.

272　Virus identity may be confirmed by PCR or by immunofluorescence with specific antisera. Virus isolates from
273　positive cultures should be submitted to a WOAH Reference Laboratory <u>for strain characterisation and</u> to maintain
274　a geographically diverse archive. ~~Further strain characterisation for surveillance purposes or detection of the~~
275　~~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.

280　In the United States of America (USA), potent polyclonal antiserum to EHV-1, prepared in swine and conjugated with
281　FITC, is available to veterinary diagnostic laboratories for this purpose from the National Veterinary Services
282　Laboratories of the United States Department of Agriculture (USDA). The antiserum cross-reacts with EHV-4 and
283　hence is not useful for serotyping, however virus typing can be conducted on any virus positive specimens by PCR.

284　Freshly dissected samples (5 × 5 mm pieces) of fetal tissue (lung, liver, thymus, and spleen) are frozen, sectioned
285　on a cryostat at –20°C, mounted on to microscope slides, and fixed with 100% acetone. After air-drying, the sections
286　are incubated at 37°C in a humid atmosphere for 30 minutes with an appropriate dilution of the conjugated swine
287　antibody to EHV-1. Unreacted antibody is removed by two washes in PBS, and the tissue sections are then covered
288　with aqueous mounting medium and a cover-slip, and examined for fluorescent cells indicating the presence of EHV
289　antigen. Each test should include a positive and negative control consisting of sections from known EHV-1 infected
290　and uninfected fetal tissue.

1.5.　Virus detection by immunoperoxidase staining

292　Immunohistochemical (IH) staining methods, such as immunoperoxidase, have been developed for detecting EHV-1
293　antigen in fixed tissues of aborted equine fetuses, placental tissues or neurologically affected horses (Schultheiss *et*
294　*al*., 1993; Whitwell *et al*., 1992). Such techniques can be used as an alternative to immunofluorescence described
295　above and can also be readily applied to archival <u>frozen or fixed</u> tissue samples. Immunohistochemical staining for
296　EHV-1 is particularly useful for the simultaneous evaluation of morphological lesions and the identification of the virus.
297　Immunoperoxidase staining for EHV-1<u>/</u>4 may also be carried out on infected cell monolayers ~~(van Maanen *et al*.,~~
298　~~2000)~~. Adequate controls must be included with each immunoperoxidase test run for evaluation of both the method
299　specificity and antibody specificity. ~~In one WOAH Reference Laboratory, this~~ method is used ~~routinely for frozen or~~
300　~~fixed tissue, using~~ <u>If non-specific</u> ~~rabbit~~ polyclonal sera <u>is used</u> ~~raised against EHV-1. This staining method is not~~
301　~~type-specific and therefore~~ <u>the staining method</u> needs to be combined with virus isolation or PCR to discriminate
302　between EHV-1 and EHV-4~~, however it provides a useful method for rapid diagnosis of EHV-induced abortion~~.

1.6.　Histopathology

304　Histopathological examination of sections of fixed placenta and lung, liver, spleen, adrenal and thymus from aborted
305　fetuses and brain and spinal cord from neurologically affected horses should be carried out. In aborted fetuses,
306　eosinophilic intranuclear inclusion bodies present within bronchiolar epithelium or in cells at the periphery of areas of
307　hepatic necrosis are consistent with a diagnosis of herpesvirus infection. The characteristic microscopic lesion
308　associated with EHV-1 neuropathy is a degenerative thrombotic vasculitis of small blood vessels in the brain or spinal
309　cord (perivascular cuffing and infiltration by inflammatory cells, endothelial proliferation and necrosis, and thrombus
310　formation).

2.　Serological tests

312　EHV-1 and EHV-4 are endemic in most parts of the world and seroprevalence is high, however serological testing of paired
313　sera can be useful for diagnosis of ER in horses. A positive diagnosis is based on the demonstration of significant increases
314　(four-fold or greater) in antibody titres in paired sera taken during the acute and convalescent stages of the disease. The
315　results of tests performed on sera from a single collection date are, in most cases, impossible to interpret with any degree
316　of confidence. The initial (acute phase) serum sample should be taken as soon as possible after the onset of clinical signs,
317　and the second (convalescent phase) serum sample should be taken 2–4 weeks later.

318　'Acute phase' sera from mares after abortion or from horses with EHV-1 neurological disease may already contain maximal
319　titres of EHV-1 antibody, with no increase in titres detectable in sera collected at later dates. In such cases, serological
320　testing of paired serum samples from clinically unaffected cohort members of the herd may prove useful for retrospective
321　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.~~

325 Serum antibody levels to EHV-1/4 may be determined by virus neutralisation (VN) (Thomson *et al*., 1976), complement
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.~~
329 ~~Furthermore,~~ The CF and VN tests detect antibodies that are cross-reactive between EHV-1 and EHV-4. Nonetheless, the
330 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
331 serological confirmation of recent infection with one of the viruses. <u>Commercial ELISAs that distinguish EHV-1 and EHV-</u>
332 <u>4 antibodies are available but less widely used than the CF and VN tests. Unlike other alphaherpesviruses, DIVA[41] ELISAs,</u>
333 <u>which have been very useful in eradication programmes for bovine rhinotracheitis and pseudorabies (Aujeszky's disease),</u>
334 <u>have not been developed for EHV-1/4 .</u>

335 The ~~microneutralisation test is a~~ <u>VN and the CF tests are</u> widely used and sensitive serological assay<u>s</u> for detecting EHV-
336 1/4 antibody and will thus be described here.

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
339 dose of virus and doubling dilutions of equine test sera. At least ~~two~~ <u>three</u> replicate wells for each serum dilution are
340 required. <u>Heat-inactivated maintenance medium with a concentration of 2% FCS (HIMM)</u> ~~Serum-free MEM~~ is used
341 throughout as a diluent. Virus stocks of known titre are diluted just before use to contain 100 TCID$_{50}$ (50% tissue
342 culture infective dose) in 25 µl. Monolayers of ~~E-Derm or~~ RK-13 cells are <u>prepared</u> ~~monodispersed with EDTA/trypsin~~
343 ~~and resuspended at a concentration of 5 × 10⁵/ml. Note that RK-13 cells can be used with EHV-1 but do not show~~
344 ~~CPE with EHV 4.~~ Antibody positive and negative control equine sera and controls for cell viability, virus infectivity,
345 and test serum cytotoxicity, must be included in each assay. End-point VN titres of antibody are calculated by
346 determining the reciprocal of the highest serum dilution that protects <u>≥75%</u> ~~100%~~ of the cell monolayer from virus
347 destruction in ~~both of~~ the replicate wells.

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.

2.1.1. Test procedure

352 A suitable test procedure is as follows:

353   i)   <u>Prepare semi-confluent monolayers in tissue culture microtitre plates.</u>

354   ii)   Inactivate test and control sera for 30 minutes in a water bath at 56°C.

355   iii)   Add <u>40</u> ~~25~~ µl of <u>HIMM</u> ~~serum-free MEM~~ to all wells of the microtitre assay plates.

356   iv)   <u>For test sample titration,</u> pipette <u>25</u>~~40~~ µl of each test serum into ~~duplicate~~ <u>triplicate</u> wells of ~~both~~
357   ~~rows A and B of~~ the plate. The first <u>two</u> rows serve as the dilution of the test serum and the third
358   row serves as the serum toxicity control ~~and the second row as the first dilution of the test~~. Make
359   doubling dilutions of each serum ~~starting with row B and proceeding to the bottom of the plate~~
360   by sequential mixing and transfer of ~~25~~<u>40</u> µl to each subsequent row of wells. ~~Six sera can be~~
361   ~~assayed in each plate.~~ <u>Add 40µl of HIMM to the serum control rows.</u>

362   v)   Add <u>40</u> ~~25~~ µl of the appropriately diluted EHV-1 ~~or EHV-4~~ virus stock to ~~each~~ <u>all</u> well<u>s</u>
363   (100 TCID$_{50}$/well) <u>of the test plate</u> except those of ~~row A, which are~~ the serum control<u>s</u> ~~wells~~.
364   Note that the final serum dilutions, after addition of virus, run from <u>a starting dilution of 1/4</u> ~~to~~
365   ~~1/256~~. A separate control plate should include titration of both a negative and positive <u>(high and</u>
366   <u>low)</u> horse ~~serum~~<u>sera</u> of known titre, cell control (no virus), <u>and a back titration of</u> virus ~~control~~
367   ~~(no serum), and a virus titration~~ <u>using six wells per log dilution (100 TCID$_{50}$ to 0.01 TCID$_{50}$/well)</u>
368   ~~calculate the actual amount of virus used in the test~~

369   vi)   Incubate the plates for 1 hour at 37°C in 5% CO$_2$ atmosphere. ~~Add 50 µl of the prepared E-~~
370   ~~Derm or RK-13 cell suspension (5 × 10⁵ cells/ml) in MEM/10% FCS to each well.~~

371   vii)   <u>Transfer 50 µl from each well of the test and control plates to the tissue culture microtitre plates.</u>

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41   DIVA: detection of infection in vaccinated animals

372	viii)	Incubate the plates for 2̲-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. Confirm the validity of the test by establishing that the working dilution of stock virus is at 100 $TCID_{50}$/well, that the (high and low) positive control sera are within one well of their pre-determined titre and that the negative control serum is negative at a 1/4 dilution. This takes approximately 72 hours. If at this stage the antigen is too weak the virus concentration may be increased by extending the incubation period up to 5 days. If the antigen is too strong the test must be repeated.

Wells are scored as positive for neutralisation of virus if ≥ 75% of the cell monolayer remains intact. The highest dilution of serum resulting in ≥ 75% neutralisation of virus (<25% CPE) in replicate wells is the end-point titre for that serum. Examine the plates microscopically for CPE and record the results on a worksheet.

383 x) Alternatively, the cell monolayers can be scored for CPE after fixing and staining as follows: after removal of the culture fluid, immerse the plates for 15 minutes in a solution containing 2 mg/ml crystal violet, 10% formalin, 45% methanol, and 45% water. Then, rinse the plates vigorously under a stream of running tap water. Wells containing intact cell monolayers stain blue, while monolayers destroyed by virus do not stain. ~~Verify that the cell control, positive serum control, and serum cytotoxicity control wells stain blue, that the virus control and negative serum control wells are not stained, and that the actual amount of virus added to each well is 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 neutralisation of virus (no CPE) in both duplicate wells is the end-point titre for that serum.~~

393 xi) Calculate the neutralisation titre for each test serum, and compare acute and convalescent phase serum titres from each animal for a four-fold or greater increase.

2.2. Complement fixation test

The CF test can be used for the detection and quantification of antibodies against EHV-1. The test determines whether an antigen and an antibody are capable of forming a complex. The presence of an immune complex is revealed by the detector system, which consists of guinea-pig complement and sensitised sheep red blood cells (SRBCs) coated with rabbit haemolytic serum (haemolysin). In the absence of antibodies against equine herpesvirus, no antibody/antigen complex is formed, the complement remains free in the solution and the sensitised SRBCs become lysed. In the presence of antibodies against equine herpesvirus, an antibody/antigen complex is formed, the complement becomes fixed and is therefore unable to lyse the SRBCs. They subsequently form a button at the bottom of the test well.

Guinea-pig complement, rabbit haemolytic serum, complement fixation diluent (CFD) and bovine serum albumin (BSA) can be obtained commercially. The dilution of guinea-pig complement that has activity at 3 HD (haemolytic dose) in the presence of sensitised SRBCs should be optimised. The recommended dilution of rabbit haemolytic serum (or the working dilution) is sometimes provided by the supplier. However, the optimal dilution of haemolysin should be determined with the in use reagents (complement etc.) so that the test can be performed reproducibly. The optimum concentration of antigen to be used in the test should be determined using an antigen versus antibody chequerboard technique and by testing a panel of known positive sera.

The test is performed in U bottomed microtitre plates. Paired sera should be assayed on the same plate. An antibody positive serum should be included as a control on each plate. All sera are tested on a second plate containing all components except virus to check for anti-complementary activity. A back titration of the working dilution (3 HD) of complement to 2 HD, 1 HD, 0.5 HD is set up in duplicate wells on the complement control plate (eight wells in total). An SRBC control is set up in eight wells.

2.2.1. Preparation of samples

417 i) Samples and controls are prepared by adding 4 volumes (600 µl) of CFD to 1 volume (150 µl) 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 complement.

2.2.2. Test procedure

422 i) Prepare the test plate and anti-complementary plate by adding 25 µl 0.05% BSA/CFD to all wells except the first column (H).

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.
426 427 428	iv)	Add 50 µl of the diluted inactivated test serum and controls to the first well of each row on both 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.
433 434	viii)	Add 25 µl of guinea-pig complement diluted in 0.05% BSA/CFD to 3 HD to all wells except the 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.

2.2.3. Preparation 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.
442 443 444	iii)	Mix the 2% SRBCs with an equal volume of BSA/CFD solution containing haemolysin at its 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.
446 447	v)	The following day, incubate the 1% sensitised SRBCs at 37°C for 30 minutes. During the final 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.

2.2.4. Reading results

455 456 457	i)	Confirm the validity of the test by establishing that the working dilution of complement is at 3 HD: 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.
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.
460 461 462 463	iii)	Confirm that there are no buttons visible on the anti-complementary plates. Buttoning indicates 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 retested and treated as described below.
464 465	iv)	In the test wells, buttoning indicates the presence of antibodies in the serum. The antibody titre is the dilution at which there is 50% buttoning and 50% lysis observed.

2.2.5. Treatment 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

1. Background

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 permutations of EHV-1 and EHV-4 and some also include equine influenza virus.

Clinical experience has demonstrated that vaccination can be useful for reducing clinical signs of respiratory disease and incidence of abortion, however none of the vaccines protects against neurological disease. Multiple doses repeated annually, of each of the currently marketed ER vaccines are recommended by their respective manufacturers. Vaccination schedules vary with a particular vaccine.

The indications stated on the product label for use of several available vaccines for ER are either as a preventative of herpesvirus-associated respiratory disease, or as an aid in the prevention of abortion, or both. A minority of ~~Only four~~ 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 have been demonstrated to prevent the occurrence of neurological disease sometimes associated with EHV-1 infection.

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.

2. Outline of production and minimum requirements for vaccines

2.1. Characteristics of the seed

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 line approved for equine vaccine production by the appropriate regulatory agency. A complete record of original source (including isolate number, location, year of isolation), passage history, medium used for propagation, etc., shall be kept for the master seed preparations of both the virus(es) and cell stock(s) intended for use in vaccine production.

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 experimental test vaccine prepared from the highest passage level of the MSV allowed for use in vaccine production. The test for MSV immunogenicity consists of vaccination of horses with low antibody titres to EHV-1/4, with doses of the test vaccine that will be recommended on the final product label. Second serum samples should be obtained and tested for significant increases in neutralising antibody titre against the virus, 21 days after the final dose.

Samples of each lot of MSV to be used for preparation of live attenuated ER vaccines must be tested for safety in horses determined to be susceptible to the virulent wild-type virus, including pregnant

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.

2.2. Method of manufacture

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

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

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 vaccine must demonstrate complete inactivation of virus (for inactivated vaccines) as well as a level of residual virus-killing agent that does not exceed the maximal allowable limit (e.g. 0.2% for formaldehyde).

iv) Batch potency

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

2.3.1. Manufacturing process

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 |
| 570 | typical industrial batch volume. |

2.3.2 Safety requirements

| 572 | Vaccine safety should be evaluated in vaccinated animals using different assays (see Section |
| 573 | 2.2.3.iii). |

2.3.3 Efficacy requirements

| 575 | Vaccine efficacy (protection) is estimated in vaccinated animals directly by evaluating their resistance |
| 576 | to live pathogen challenge. |

2.3.4 Duration of immunity

578	As part of the licensing or marketing authorisation procedure, the manufacturer may be required to
579	demonstrate the duration of immunity (DOI) of a given vaccine by either challenge or alternative test
580	at the end of the claimed period of protection.

581	Tests to establish the duration of immunity to EHV-1/4 or EHV1/4 achieved by immunisation with
582	each batch of vaccine are not required. The results of many reported observations indicate that
583	immunity induced by vaccination against EHV-1 or EHV induced immunity to EHV-1/4 is not more
584	than a few months in duration; these observations are reflected in the frequency of revaccination
585	recommended on ER vaccine product labels.

2.3.5 Stability

587	As part of the licensing or marketing authorisation procedure, the manufacturer will be required to
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
590	freezing or ambient temperature.

591	At least three production batches of vaccine should be tested for shelf life before reaching a
592	conclusion on the vaccine's stability. When stored at 4°C, inactivated vaccine products generally
593	maintain their original antigenic potency for at least 1 year. Lyophilised preparations of the live virus
594	vaccine are also stable during storage for 1 year at 4°C. Following reconstitution, live virus vaccine
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.

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745 * *

746 **NB:** There are WOAH Reference Laboratories for equine rhinopneumonitis (please consult the WOAH Web site:
747 https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).
748 Please contact the WOAH Reference Laboratories for any further information on

749   diagnostic tests, reagents and vaccines for equine rhinopneumonitis
750   and to submit strains for further characterisation.

751   **NB:** First adopted in 1990. Most recent updates adopted in 2017.

2　　　　**MEETING OF THE BIOLOGICAL STANDARDS COMMISSION**

3　　　　**Paris, 4–8 September 2023**

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5　　　　**S E C T I O N　3 . 8 .**

6　　　　~~**O V I D A E　A N D**~~ **C A P R I N A E**

7　　　　**C H A P T E R　3 . 8 . 1 .**

8　　　　**B O R D E R　D I S E A S E**

9　　　　**SUMMARY**

10　　*Border disease (BD) is a viral disease of sheep and goats first reported in sheep in 1959 from the border*
11　　*region of England and Wales, and since recorded world-wide. Prevalence rates in sheep vary from 5% to*
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)<u>. Consequently,</u> the disease*
15　　*has <u>sometimes</u> been referred to as 'hairy shaker disease'. Vertical transmission plays an important role in*
16　　*the epidemiology of the disease. Infection of fetuses can result in the birth of persistently infected (PI) lambs.*
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*
21　　*there is close contact between sheep or goats and cattle, the same clinical signs may be caused by infection*
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*
24　　*germplasm for international movement. It is important to identify the viraemic PI animals so that they will not*
25　　*be used for breeding or trading purposes. Serological testing is insufficient. However, it is generally*
26　　*considered that serologically positive, nonviraemic sheep ~~are 'safe',~~ <u>do not present a risk</u> as latent infections*
27　　*are not known to occur in recovered animals.<u> Pregnant seropositive, nonviraemic animals may, however,</u>*
28　　*<u>present a risk by carrying a PI fetus that cannot be detected until after parturition.</u>*

29　　***Identification of the agent:*** *BDV is a <u>species of</u>* Pestivirus <u>(Pestivirus ovis)</u> *in the family* Flaviviridae *and is*
30　　*closely related to classical swine fever virus* (Pestivirus suis) *and BVDV <u>viruses, which are classified in the</u>*
31　　*<u>distinct species</u>* Pestivirus bovis *(<u>commonly known as BVDV type 1</u>),* Pestivirus tauri *(<u>formerly BVDV type 2</u>)*
32　　*<u>and</u>* Pestivirus brazilense *(<u>BVDV type 3 or Hobi-like pestivirus</u>). Nearly all isolates of BDV are*
33　　*noncytopathogenic in cell culture. There are no defined serotypes but virus isolates exhibit considerable*
34　　*antigenic diversity. A number of separate genotypes, have been identified.*

35　　*Apparently healthy PI sheep resulting from congenital infection can be identified by direct detection of virus*
36　　*or nucleic acid in blood or tissues or by virus isolation in cell culture followed by immunostaining to detect*
37　　*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 <u>reverse transcriptase</u> polymerase chain reaction*
42   *methods that are able to detect residual nucleic acid. However, tissues and blood from PI sheep more than*
43   *a few months old contain high levels of virus, which can be easily identified by isolation and direct methods*
44   *to detect antigens or nucleic acids. As sheep may be infected with BVDV, it is preferable to use diagnostic*
45   *assays that are 'pan-pestivirus' reactive and will readily detect all strains of BDV and BVDV.*

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

60 Border disease virus (BDV) is a *Pestivirus* of the family *Flaviviridae* and is closely related to classical swine fever virus
61 (CSFV) and bovine viral diarrhoea virus (BVDV). There are ~~four~~<u>a number of</u> officially recognised species, namely – <u>BDV</u>
62 <u>(*Pestivirus ovis*)</u> CSFV (*Pestivirus suis*), BVDV types 1 and 2 <u>(taxonomically known as *Pestivirus bovis* and *Pestivirus tauri*,</u>
63 <u>respectively)</u> and ~~BDV (ICTV, 2016)~~ BVDV 3 or Hobi-like pestivirus (*Pestivirus brazilense*), but a number of other
64 pestiviruses that are considered to be distinct species have been reported. While CSF viruses are predominantly restricted
65 to pigs, ~~examples of~~ <u>there are situations where</u> the other ~~three~~ species have ~~all~~ been recovered from sheep. While the
66 majority of isolates have been identified as BD viruses in areas where sheep or goats are raised in isolation from other
67 species (Vilcek *et al*., 1997), in regions where there is close contact between small ruminants and cattle, BVDV may be
68 frequently identified (Carlsson, 1991). Nearly all virus isolates of BDV are noncytopathogenic, although occasional
69 cytopathic viruses have been isolated (Vantsis *et al*., 1976). BDV spreads naturally among sheep by the oro-nasal route
70 and by vertical transmission. It is principally a cause of congenital disease in sheep and goats, but can also cause acute
71 and persistent infections. Infection is less common in goats, in which persistent infection is rare as abortion is the main
72 presenting sign. Pigs may also be infected by pestiviruses other than CSFV and antibodies to BDV in pigs may interfere
73 with tests for the diagnosis of CSF (Oguzoglu *et al*., 2001). Several genotypes of BD viruses from sheep, goats and
74 Pyrenean chamois (*Rupicapra pyrenaica pyrenaica*) have been described. Phylogenetic analysis using computer-assisted
75 nucleotide sequence analysis suggests that genetic variability among BD viruses is greater than within each of the other
76 *Pestivirus* species. Four distinguishable genogroups of BDV have been described as well as putative novel *Pestivirus*
77 genotypes from Tunisian sheep and a goat (Becher *et al*., 2003; Vilcek & Nettleton, 2006). The chamois BD virus is similar
78 to isolates from sheep in the Iberian Peninsula (Valdazo-Gonzalez *et al*., 2007). This chapter describes BDV infection in
79 sheep. Chapter 3.4.7 *Bovine viral diarrhoea* should also be consulted for related diagnostic methods.

1. Acute infections

81 Healthy newborn and adult sheep exposed to BDV usually experience only mild or inapparent disease. Slight fever and a
82 mild leukopenia are associated with a short-lived viraemia detectable between days 4 and 11 post-infection, after which
83 virus neutralising antibody appears in the serum (Thabti *et al*., 2002).

84 Acute infections are best diagnosed serologically using paired sera from a representative number of sheep. Occasional
85 BDV isolates have been shown to produce high fever, profound and prolonged leukopenia, anorexia, conjunctivitis, nasal
86 discharge, dyspnoea and diarrhoea, and 50% mortality in young lambs. One such isolate was recovered from a severe
87 epidemic of BD among dairy sheep in 1984 (Chappuis *et al*., 1986). A second such isolate was a BDV contaminant of a
88 live CSFV vaccine (Wensvoort & Terpstra, 1988).

2. Fetal infection

The main clinical signs of BD are seen following the infection of pregnant ewes. While the initial maternal infection is subclinical or mild, the consequences for the fetus are serious. Fetal death may occur at any stage of pregnancy, but is 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 larger fetuses, stillbirths and the premature births of small, weak lambs will be seen. Confirmation that an abortion or stillbirth is due to BDV is often difficult to establish, but virus may be isolated from fetal tissues in some cases. The use of an appropriate real-time reverse-transcription polymerase chain reaction (RT-PCR) assay may give a higher level of 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). Samples of fetal fluids or serum should be tested for BDV antibody.

During lambing, an excessive number of barren ewes will become apparent, but it is the diseased live lambs that present the main clinical features characteristic of BD. The clinical signs exhibited by BD lambs are very variable and depend on 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 nervous signs of BD are its most characteristic feature. The tremor can vary from violent rhythmic contractions of the muscles of the hindlegs and back, to barely detectable fine trembling of the head, ears, and tail. Fleece abnormalities are 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 BDV or antibody should be collected into anticoagulant from suspect lambs before they have received colostrum. Once lambs have ingested colostrum, it is difficult to isolate virus until they are 2 months old and maternal antibody levels have 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 detection of the Erns antigen appear to be less prone to interference by maternal antibodies and can often be used to detect antigen in serum.

With careful nursing, a proportion of BD lambs can be reared, although deaths may occur at any age. The nervous signs gradually decline and can disappear by 3–6 months of age. Weakness, and swaying of the hind-quarters, together with fine trembling of the head, may reappear at times of stress. Affected lambs often grow slowly and under normal field conditions many will die before or around weaning time. In cases where losses at lambing time have been low and no 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).

3. Persistent viraemia

When fetuses survive an infection that occurs before the onset of immune competence, they are born with a persistent viraemia. The ovine fetus can first respond to an antigenic stimulus between approximately 60 and 85 days of its 150-day gestation period. In fetuses infected before the onset of immune competence, viral replication is uncontrolled and 50% fetal death is common. In lambs surviving infection in early gestation, virus is widespread in all organs. Such lambs appear 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 in the central nervous system (CNS) and skin. Throughout the CNS, there is a deficiency of myelin, and this causes the 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.

Persistently viraemic sheep can be identified by the detection of viral antigens, nucleic acids or infectious virus in a blood sample. Viraemia is readily detectable by testing of serum at any time except within the first 2 months of life, when virus may be masked by colostral antibody and, possibly, in animals older than 4 years, some of which develop low levels of anti-BDV antibody (Nettleton *et al*., 1992). Methods other than virus isolation may be preferred to avoid interference from antibodies. When the presence of colostral antibodies is suspected, the virus may be detected in washed leukocytes and 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 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.

144 Some viraemic sheep survive to sexual maturity and are used for breeding. Lambs born to these infected dams are always
145 persistently viraemic. Persistently viraemic sheep are a continual source of infectious virus to other animals and their
146 identification is a major factor in any control programme. Sheep being traded should be screened for the absence of BDV
147 viraemia.

148 Usually persistently infected (PI) rams have poor quality, highly infective semen and reduced fertility. All rams used for
149 breeding should be screened for persistent BDV infection on a blood sample. Semen samples can also be screened for
150 virus, but virus isolation is much less satisfactory than from blood because of the toxicity of semen for cell cultures. Real-
151 time RT-PCR for detection of pestivirus nucleic acid would usually overcome toxicity problems, and thus this assay should
152 be useful for testing semen from rams.

4. Late-onset disease in persistently viraemic sheep

154 Some PI sheep housed apart from other animals spontaneously develop intractable diarrhoea, wasting, excessive ocular
155 and nasal discharges, sometimes with respiratory distress. At necropsy such sheep have gross thickening of the distal
156 ileum, caecum and colon resulting from focal hyperplastic enteropathy. Cytopathic BDV can be recovered from the gut of
157 these lambs. With no obvious outside source of cytopathic virus, it is most likely that such virus originates from the lamb's
158 own virus pool, similar to what occurs with BVDV. Other PI sheep in the group do may not develop the disease. This
159 syndrome, which has been produced experimentally and recognised in occasional field outbreaks of BD, has several
160 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*

Method	Purpose					
	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	+++	+++	++	+++	+++	+++

163 Key: +++ = recommended for this purpose; ++ recommended but has limitations;
164 + = suitable in very limited circumstances; – = not appropriate for this purpose.
165 ELISA = enzyme-linked immunosorbent assay; IHC = immunohistochemistry; NA = nucleic acid; RT-PCR = reverse-transcription
166 polymerase chain reaction; ISH = *in-situ* hybridisation; VN = virus neutralisation.
167 [a]A combination of agent identification methods applied on the same clinical sample is recommended.

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 advice[42]. 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.

1.1. Virus isolation

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 that a laboratory quality assurance programme be in place. Chapter 3.4.7 provides detailed methods for virus isolation 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 cultures, medium components and reagents are equally relevant to this chapter. Provided proven pan-pestivirus reactive reagents (e.g. monoclonal antibodies [MAbs], primers and probes for real-time RT-PCR) are used for antigen or nucleic acid detection, the principal difference is the selection of appropriate cell cultures.

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 *et al*., 2002) or sheep choroid plexus can be useful, but different lines vary considerably in their susceptibility to the virus. Ovine cells have been used successfully for the isolation and growth of BD viruses and BVDV types 1 and 2 from sheep. In regions where sheep may become infected with BVD viruses from cattle, a virus isolation system using both ovine and bovine cells could be optimal. However, bovine cells have lower sensitivity for the primary isolation and growth of some BD viruses, so reliance on bovine cells alone is inadvisable. Details of suitable bovine cell cultures are provided chapter 3.4.7. The precautions outlined in that chapter for the establishment of cells and medium 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.

From live animals, serum is the most frequently used sample to be tested for the presence of infectious virus. However, for difficult cases, the most sensitive way to confirm pestivirus viraemia is to wash leukocytes repeatedly (at least three times) in culture medium before co-cultivating them with susceptible cells in either cell culture tubes or 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 immunocytochemistry. Staining for noncytopathic pestiviruses will usually detect virus at the end of the primary 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 medium because some high titred field isolates will replicate poorly if passaged undiluted (i.e. at high multiplicity of infection – moi).

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 at least 1/10 in culture medium. As the major threat of BDV-infected semen is from PI rams, blood is a more reliable clinical sample than semen for identifying such animals. There are many variations in virus isolation procedures. All should be optimised for maximum sensitivity using a standard reference virus preparation and, whenever possible, recent BDV field isolates. Most of the limitations of virus isolation for the detection of BDV in serum or blood, tissues 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 strains for future reference or research purposes.

For specific technical details of virus isolation procedures, including immunoperoxidase staining, refer to chapter 3.4.7.

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42 Please consult the WOAH Web site: https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3

1.2. Nucleic acid detection methods

The complete genomic sequences of three BD viruses have been determined and compared with those of other pestiviruses (Becher *et al*., 1998; Ridpath & Bolin, 1997). Phylogenetic analysis shows BD viruses to be more closely related to CSFV than to BVDV (Becher *et al*., 2003; Van Rijn *et al*., 1997; Vilcek & Nettleton, 2006; Vilcek *et al*., 1997). Real-time RT-PCR for diagnosing pestivirus infection is now used widely and a number of formats have been described. Real-time RT-PCR assays have the advantages of being able to detect both infectious virus and residual nucleic acid, the latter being of value for investigating abortions and lamb deaths. Furthermore, the presence of virus-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, are less affected by components of the semen compared with virus isolation. Because of the potential for small ruminants to be infected with genetically different strains of BDV or with strains of BVDV, a ~~proven~~ pan-pestivirus reactive real-time RT-PCR <u>with proven high sensitivity</u> should be used. <u>To ensure that the genetic spectrum of BDV strains is sufficiently covered, it may be necessary to apply a broadly reactive BDV specific real time RT-PCR in parallel to maximise diagnostic sensitivity.</u> 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 closely.

After testing samples in a pan-pestivirus reactive assay, samples giving ~~positive results can~~ <u>any level of reactivity should</u> be investigated further by the application of a BDV-specific real-time RT-PCR (Willoughby *et al*., 2006). It is important to note however that different genotypes of BDV may be circulating in some populations, especially wild ruminants such as chamois and deer, and may be transferred to sheep. An assay that is specific for the detection of 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. <u>Nevertheless, there are also situations where a pan-pestivirus reactive real-time RT-PCR may have lower analytical sensitivity. Consequently, in any situation where BDV infection is suspected, the application of several diagnostic methods is recommended. Maternal serology can also play an important role as negative results should exclude the potential involvement of a pestivirus.</u>

1.3. Enzyme-linked immunosorbent assay for antigen detection

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 detection was described for detecting viraemic sheep and was later modified into a double MAb capture ELISA for use in sheep and cattle (Entrican *et al.,* 1994). The test is most commonly employed to identify PI viraemic sheep 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 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 detect acute BDV infections on blood samples. As well as for testing leukocytes, the antigen ELISA can also be used 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 are at present no commercially available kits that have been fully validated for detecting BDV. Prior to use for regulatory purposes, these kits should be validated in the region where they are to be used to ensure that a wide range of field strains of BDV can be detected and that they are suitable for the sample types to be tested.

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. Pan-pestivirus 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.

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.

2.1. Virus neutralisation test

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, to achieve optimal analytical sensitivity, 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 $TCID_{50}$ (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 $TCID_{50}$).

iv) The plate is incubated for 1 hour at 37°C.

v) A flask of suitable cells (e.g. ovine testis or kidney cells) is trypsinised and the cell concentration is adjusted to 2×10^5/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% CO_2 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 of serum (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.

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 cell-culture grown antigen. The results correlate qualitatively with the VN test (Fenton *et al*., 1991).

2.2.1. Antigen preparation

Use eight 225 cm^2 flasks of newly confluent FLM cells; four flasks will be controls and four will be infected. Wash the flasks and infect four with a 0.01–0.1 m.o.i. of Moredun cytopathic BDV. Allow the virus to adsorb for 2 hours at 37°C. Add maintenance media containing 2% FBS (free from BDV antibody), and incubate cultures for 4–5 days until CPE is obvious. Pool four control flask supernatants and separately pool four infected flask supernatants. Centrifuge at 3000 ***g*** for 15 minutes to pellet cells. Discard the supernatants. Retain the cell pellets. Wash the flasks with

324 50 ml of PBS and repeat the centrifugation step as above. Pool all the control cell pellets in 8 ml PBS
325 containing 1% Nonidet P40 and return 2 ml to each control flask to lyse the remaining attached cells.
326 Repeat for infected cells. Keep the flasks at 4°C for at least 2 hours agitating the small volume of fluid
327 on the cells vigorously every 30 minutes to ensure total cell detachment. Centrifuge the control and
328 infected antigen at 12,000 **g** for 5 minutes to remove the cell debris. Supernatant antigens are stored
329 at –70°C in small aliquots.

2.2.2. Test procedure

331 i)   The two MAbs are diluted to a predetermined dilution in 0.05 M bicarbonate buffer, pH 9.6. All
332   wells of a suitable ELISA-grade microtitre plate (e.g. Nunc maxisorb, Greiner 129b) are coated
333   overnight at 4°C.

334 ii) After washing three times in PBST, a blocking solution of PBST containing 10% horse serum
335   (PBSTH) is added to all wells, which are incubated at 37°C for 1 hour.

336 iii) The antigen is diluted to a predetermined dilution in PBSTH and alternate rows of wells are
337   coated with virus and control antigens for 1 hour at 37°C. The plates are then washed three
338   times in PBST before addition of test sera.

339 iv) Test sera are diluted 1/50 in PBSTH and added to duplicate virus and duplicate control wells
340   for 1 hour at 37°C. The plates are then washed three times in PBST.

341 v)   Anti-ovine IgG peroxidase conjugate is diluted to a predetermined dilution in PBSTH and added
342   to all wells for 1 hour at 37°C. The plates are washed three times in PBST.

343 vi) A suitable activated enzyme substrate/chromogen, such as ortho-phenylene diamine (OPD) or
344   tetramethyl benzidine (TMB), is added). After colour development, the reaction is stopped with
345   sulphuric acid and the absorbance read on an ELISA plate reader. The mean value of the two
346   control wells is subtracted from the mean value of the two virus wells to give the corrected
347   absorbance for each serum. Results are expressed as corrected absorbance with reference to
348   the corrected absorbance of known positive and negative sera. Alternatively, ELISA titres can
349   be extrapolated from a standard curve of a dilution series of a known positive reference serum.

350 vii) If antigens of sufficient potency can be produced the MAb capture stage can be omitted. In this
351   case alternate rows of wells are coated with virus and control antigen diluted to a predetermined
352   dilution in 0.05 M bicarbonate buffer, pH 9.6, overnight at +4°C. The plates are washed and
353   blocked as in step ii above. After washing, diluted test sera are added and the test proceeds
354   from step iv as above.

C. REQUIREMENTS FOR VACCINES

1. Background

357 To be useful, a BDV vaccine should be effective when administered to female sheep before breeding to prevent
358 transplacental infection. Experimental and commercial inactivated whole virus BDV vaccines have been produced in
359 Europe (Brun *et al*., 1993; Vantsis *et al*., 1980). Unlike vaccines for BVDV, there is limited demand for vaccines against
360 BDV and those produced have only been inactivated products. No live attenuated or recombinant subunit vaccines for
361 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
363 use in pigs, cattle, sheep and goats. Contaminated vaccines have included those used for the control of Aujesky's disease,
364 CSF, rotavirus, coronavirus, rinderpest, sheep pox and contagious pustular dermatitis. The insidious ability of pestiviruses
365 to cross the placenta, and thus establish PI animals, gives them the potential to contaminate vaccines through cells, serum
366 used as medium supplement, or seed stock virus. As nearly all isolates of pestiviruses are noncytopathic, they will remain
367 undetected unless specific tests are carried out. Although such contamination should be less likely to be a problem with
368 an inactivated vaccine, nevertheless steps should be taken to ensure that materials used in production are not
369 contaminated.

1.1. Characteristics of a target product profile

371 Traditionally, pestivirus vaccines fall into two classes: modified live or inactivated virus vaccines. The essential
372 requirement for both types is to afford a high level of fetal infection. Only inactivated vaccines have been produced
373 for BDV. Properly formulated inactivated vaccines are very safe to use but, to obtain satisfactory levels of immunity,
374 they usually require booster vaccinations, which may be inconvenient. Because of the propensity for antigenic
375 variability, the vaccine should contain strains of BDV that are closely matched to viruses found in the area in which

376 they are used. This may present particular challenges with BDV in regions where several antigenic types have been
377 found. Due to the need to customise vaccines for the most commonly encountered strains within a country or region,
378 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.

2. Outline of production and minimum requirements for vaccines

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
385 may be challenging however, because of the range of pestiviruses with which sheep can be infected. There is
386 considerable antigenic variation across these viruses – both between viruses that have been classified in the BDV
387 genogroup as well as between viruses in the BVDV1 and BVDV2 genotypes (Becher *et al*., 2003; Vilcek & Nettleton,
388 2006; Wensvoort *et al*., 1989). Infection of sheep with the putative BVDV-3 genotype has also been described (Decaro
389 *et al.,* 2012). It is likely that the antigenic composition of a vaccine will vary from region to region to provide an
390 adequate antigenic match with dominant virus strains. Cross-neutralisation studies are required to establish optimal
391 combinations. Nevertheless it would appear that any BDV vaccine should contain at least a representative of the BDV
392 and BVDV (type 1) groups. Characterisation of the biologically cloned vaccine viruses should include typing with
393 MAbs and genotyping (Paton *et al*., 1995).

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

395 It is crucial to ensure that all materials used in the preparation of the bulk antigens have been
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
398 viruses and particularly BVDV can readily infect small ruminants such as sheep. Therefore, it is
399 particularly important to ensure that any serum used that is of bovine origin is free of both adventitious
400 BVDV and antibodies against BVDV strains because low levels of either virus or antibody can mask
401 the presence of the other. Materials and vaccine seeds should be tested for sterility and freedom
402 from contamination with other agents, especially viruses as described in the chapter 1.1.8 and
403 Chapter 1.1.9 *Tests for sterility and freedom from contamination of biological materials intended for*
404 *veterinary use*.

405 If a vaccine passes basic tests, the efficacy of vaccination should ultimately be measured by the
406 capacity to prevent transplacental transmission. Effective challenge of vaccinated pregnant ewes at
407 50–60 days gestation has been achieved by intranasal installation of virus or by mixing with PI sheep
408 (Brun *et al.,* 1993). Usually this reliably produces persistently viraemic offspring in non-immune ewes.
409 In regions where multiple genotypes of BDV viruses are commonly encountered, efficacy in protecting
410 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
414 rolled cell cultures. Inactivants have included formalin and beta-propriolactone. Adjuvants have
415 included aluminium hydroxide and oil (Brun *et al*., 1993; Vantsis *et al*., 1980). Optimal yields depend
416 on the cell type and isolate used. A commercial BDV vaccine containing two strains of virus has been
417 prepared on ovine cell lines (Brun *et al*., 1993). Cells must be produced according to a seed-lot
418 system from a master cell seed (MCS) that has been shown to be free from all contaminating
419 microorganisms. Vaccine should only be produced in cells fewer than 20 passages from the MCS.
420 Control cells from every passage should be checked for pestivirus contamination. Standard
421 procedures may be used, with the expectation for harvesting noncytopathic virus on days 4–7 after
422 inoculation of cultures. The optimal yield of infectious virus will depend on several factors, including
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 large-
425 scale virus production. Whether a live or inactivated vaccine, the essential aim will be to produce a
426 high-titred virus stock. This bulk antigen preparation can subsequently be prepared according to the
427 type of vaccine being considered.

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.

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.

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.

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		*in-vivo* potency results. It should be demonstrated that the lowest recommended dose of vaccine can
482		prevent transplacental transmission of BDV in pregnant sheep.

2.3. Requirements for authorisation/registration/licensing

2.3.1. Manufacturing process

485	For registration of a vaccine, all relevant details concerning manufacture of the vaccine and quality
486	control testing should be submitted to the relevant authorities. Unless otherwise specified by the
487	authorities, information should be provided from three consecutive vaccine batches with a volume
488	not less than 1/3 of the typical industrial batch volume.

489	There is no standard method for the manufacture of a BDV vaccine, but conventional laboratory
490	techniques with stationary, rolled or suspension (micro-carriers) cell cultures may be used.
491	Inactivated vaccines can be prepared by conventional methods, such as binary ethylenimine, formalin
492	or beta-propiolactone inactivation (Park & Bolin, 1987). A variety of adjuvants may be used.

2.3.2. Safety requirements

494	*In-vivo* tests should be undertaken using repeat doses (taking into account the maximum number of
495	doses for primary vaccination and, if appropriate, the first revaccination/booster vaccination) and
496	contain the maximum permitted antigen load and, depending on the formulation of the vaccine, the
497	maximum number of vaccine strains.

498	i) Target and non-target animal safety

499	The safety of the final product formulation of inactivated vaccines should be assessed in susceptible
500	young sheep that are free of maternally derived antibodies and in pregnant ewes. They should be
501	checked for any local reactions following administration, and, in pregnant ewes, for any effects on
502	the unborn lamb.

503	ii) Reversion-to-virulence for attenuated/live vaccines and environmental considerations

504	In the event that a live virus vaccine was developed for BDV, virus seeds that have been passaged
505	at least up to and preferably beyond the passage limit specified for the seed should be inoculated
506	into young lambs to confirm that there is no evidence of disease. If a live attenuated vaccine has
507	been registered for use in pregnant animals, reversion to virulence tests should also include pregnant
508	animals. Live attenuated vaccines should not be transmissible to unvaccinated 'in-contact' animals.

509	iii) Precautions (hazards)

510	BDV is not considered to be a human health hazard. Standard good microbiological practice should
511	be adequate for handling the virus in the laboratory. While the inactivated virus in a vaccine should
512	be identified as harmless for people administering the product, adjuvants included in the vaccine may
513	cause injury to people. Manufacturers should provide adequate warnings that medical advice should
514	be sought in the case of self-injection (including for adjuvants, oil-emulsion vaccine, preservatives,
515	etc.) with warnings included on the product label/leaflet so that the vaccinator is aware of any danger.

2.3.3. Efficacy requirements

517	The potency of the vaccine should be determined by inoculation into seronegative and virus negative
518	lambs, followed by monitoring of the antibody response. Antigen content can be assayed by infectivity
519	titration prior to inactivation and subsequently by ELISA and adjusted as required to a standard level
520	for the particular vaccine. Standardised assay protocols applicable to all vaccines do not exist. Live
521	vaccine batches may be assayed by infectivity titration. Each production batch of vaccine should
522	undergo potency and safety testing as batch release criteria. BVD vaccines must be demonstrated
523	to produce adequate immune responses, as outlined above, when used in their final formulation
524	according to the manufacturer's published instructions.

2.3.4. Vaccines permitting a DIVA strategy (detection of infection in vaccinated animals)

526	To date, there are no commercially available vaccines for BDV that support use of a true DIVA
527	strategy.

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.

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.

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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.

2   **MEETING OF THE BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # C H A P T E R   3 . 8 . 1 2 .

6   # SHEEP POX AND GOAT POX

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.*

13 *Sheeppox virus (SPPV) and goatpox virus (GTPV) are the causative agents of sheep pox and goat pox, and*
14 *with lumpy skin disease virus <u>(LSDV)</u> make up the genus* Capripoxvirus *in the family* Poxviridae. *Sheep pox*
15 *and goat pox are endemic in Africa north of the Equator, the Middle East and Asia, while some parts of*
16 *Europe have experienced outbreaks recently. <u>See WAHIS (https://wahis.woah.org/#/home) for recent</u>*
17 *<u>information on distribution at the country level.</u> ~~Countries that reported outbreaks of the disease between~~*
18 *~~2010 and 2015 include Bulgaria, Chinese Taipei, Israel, Kazakhstan, Kyrgyzstan, Mongolia, Morocco,~~*
19 *~~Greece and Russia, with Greece, Israel and Russia having experienced recurring incidences.~~*

20 ***Identification of the agent:*** *Laboratory confirmation of capripox<u>virus</u> is most rapid using the polymerase*
21 *chain reaction (PCR) method in combination with a clinical history consistent with generalised capripox<u>virus</u>*
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*
25 *haematoxylin and eosin staining. The antigen can also be detected in tissue culture using specific sera and*
26 *immunoperoxidase or immunofluorescence techniques. Capripoxvirus antigen and inclusion bodies may be*
27 *seen in stained cryostat or paraffin sections of biopsy or post-mortem lesion material.*

28 *~~An antigen-detection enzyme-linked immunosorbent assay (ELISA) using a polyclonal detection serum~~*
29 *~~raised against a recombinant immunodominant antigen of capripoxvirus has been developed.~~*

30 ***Serological tests:*** *The virus neutralisation test is the most specific serological test. The indirect*
31 *immunofluorescence test is less specific due to cross-reactions with antibody to other poxviruses. Western*
32 *blotting using the reaction between the P32 antigen of capripoxvirus with test sera is both sensitive and*
33 *specific, but is expensive and difficult to carry out. <u>An enzyme-linked immunosorbent assay (ELISA) has</u>*
34 *<u>been developed and validated to detect antibodies to capripoxviruses, however it cannot differentiate</u>*
35 *<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 capripoxviruses. 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.*

A. INTRODUCTION

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 (GTPPV), which cause sheep pox and goat pox, respectively. Sheep pox and goat pox are characterised by disseminated cutaneous nodules and up to 100% mortality in fully susceptible breeds of sheep and goats. In indigenous animals, generalised disease and mortality are less common, although they are seen where disease has been absent from an area or village for a period of time, when intensive husbandry methods are introduced, or in association with other disease agents, such as peste des petits ruminants virus or foot and mouth disease virus. Sheep pox and goat pox are major constraints to the introduction of exotic breeds of sheep and goats to endemic areas, and to the development of intensive livestock production.

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, non-endemic 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 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 development of skin lesions. In other breeds there is an initial rise in rectal temperature to above 40°C, followed in 2–5 days by the development of, at first, macules – small circumscribed areas of hyperaemia, which are most obvious on 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 researchers have distinguished between a vesicular and nodular form of sheep pox and goat pox (Zro *et al*., 2014b).

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

Table 1. Test methods available for diagnosis of sheep pox and goat pox and their purpose

Method	Purpose					
	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~~	++	++	++	++	++	–
IFAT	±	±	±	++	±	=
IHC	±	±	±	++	±	=
PCR	++	+++	++	+++	++	–
Detection of immune response						
VNT	++	++	++	++	++	++
IFAT	+	+	+	+	+	+
ELISA	++	++	++	++	++	++

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;
VNT = virus neutralisation; ELISA = enzyme-linked immunosorbent assay.
[a]~~A combination of agent identification methods applied on the same clinical sample is recommended.~~

1. Identification of the agent

1.1. Specimen collection and submission

Material for virus isolation and antigen detection should be collected by biopsy or at post-mortem from skin papules, lung lesions or lymph nodes. Samples for virus isolation ~~and antigen detection enzyme-linked immunosorbent assay (ELISA)~~ should be collected within the first week of the occurrence of clinical signs, before the development of neutralising antibodies. Samples for genome detection by polymerase chain reaction (PCR) may be collected before or after the development of neutralising antibody responses. Buffy coat from blood collected into EDTA (ethylene diamine tetra-acetic acid) during the viraemic stage of capripoxvirus infection (before generalisation of lesions or 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.

1.2. Virus isolation

Lesion material for virus isolation and antigen detection is homogenised. The following is an example of one technique for homogenisation: The tissue is minced using sterile scissors and forceps, and then macerated in a steel ball bearing mixer mill or ground with a sterile pestle in a mortar with sterile sand and an equal volume of sterile phosphate buffered saline (PBS) or serum-free Modified Eagle's Medium (MEM) containing sodium penicillin (1000 international units [IU]/ml), streptomycin sulphate (1 mg/ml), mycostatin (100 IU/ml) or fungizone (2.5 µg/ml) and neomycin (200 IU/ml). The homogenised suspension is freeze–thawed three times and then partially clarified by centrifugation 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 after the centrifugation step, however, the amount of virus in the supernatant might be reduced. Buffy coats may be prepared from 5–8 ml unclotted blood by centrifugation at 600 *g* for 15 minutes; the buffy coat is carefully removed into 5 ml of cold double-distilled water using a sterile Pasteur pipette. After 30 seconds, 5 ml of cold double-strength growth medium is added and mixed. The mixture is centrifuged at 600 *g* for 15 minutes, the supernatant is discarded and the cell pellet is suspended in 5 ml of growth medium, such as Glasgow's modified Eagle's medium (GMEM). After centrifugation at 600 *g* for a further 15 minutes, the resulting pellet is suspended in 5 ml of fresh GMEM. Alternatively, the buffy coat may be separated from a heparinised sample using a density gradient.

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 they are not contaminated with viruses such as bovine viral diarrhoea virus, ~~particularly those derived from a wool sheep breed~~ (see chapter 1.1.9). Madin–Darby bovine kidney (MDBK) cells have been shown to be suitable for capripoxvirus isolation (Fay *et al.,* 2020). The following is an example of an isolation technique: either 1 ml of buffy coat cell suspension or 1 ml of clarified biopsy preparation supernatant is inoculated on to a 25 cm^2 tissue culture flask of appropriate cells at 90% ~~confluent LT or LK cells~~ confluence, and the supernatant is allowed to adsorb for 1 hour at 37°C. The culture is then washed with warm 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 LT or LK cells and a~~, flying cover-slip~~s~~, or tissue culture microscope slides, ~~are~~ can also infected.

The flasks should be examined daily for 7–14 days for evidence of cytopathic effect (CPE). Contaminated flasks should be discarded. Infected cells develop a characteristic CPE consisting of retraction of the cell membrane from surrounding cells, and eventually rounding of cells and margination of the nuclear chromatin. At first only small areas of CPE can be seen, sometimes as soon as 4 days after infection; over the following 4–6 days these expand to involve the whole cell sheet. If no CPE is apparent by day 7, the culture should be freeze–thawed three times, and clarified supernatant inoculated on to fresh ~~LT or LK~~ cell culture~~s~~. At the first sign of CPE in the flasks, or earlier if a number of infected cover-slips are being used, a cover-slip should be removed, fixed in acetone and stained using H&E. Eosinophilic intracytoplasmic inclusion bodies, which are variable in size but up to half the size of the nucleus 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 specific anti-capripoxvirus serum in the medium; this provides a presumptive identification of the agent. Some strains of capripoxvirus have been adapted to grow on African green monkey kidney (Vero) cells, but these cells are not recommended for primary isolation.

1.3. Electron microscopy

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 is given below:

Material from the original tissue suspension is prepared for transmission electron microscope examination, prior to centrifugation, by floating a 400-mesh hexagon electron microscope grid, with piloform-carbon substrate activated by glow discharge in pentylamine vapour, on to 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 capripoxvirus 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 virions of capripoxvirus are indistinguishable from those of orthopoxvirus, but, apart from *Vaccinia* virus, no orthopoxvirus causes lesions in sheep and goats. However, capripoxvirus is distinguishable from the virions of parapoxvirus, that cause contagious pustular dermatitis, as they are smaller, oval in shape, and each is covered in a single continuous tubular element, which appears as striations over the virion.

1.4. Histopathology

Material for histopathology should be prepared by standard techniques. Following preparation, staining with haematoxylin and eosin (H&E), and mounting of the formalin-fixed biopsy material, a number of sections should be examined by light microscopy. On histological examination, the most striking aspects of acute-stage skin lesions are a massive cellular infiltrate, vasculitis and oedema. Early lesions are characterised by marked perivascular cuffing. Initially infiltration is by macrophages, neutrophils and occasionally eosinophils, and as the lesion progresses, by more macrophages, lymphocytes and plasma cells. A characteristic feature of all capripoxvirus infections is the 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, poorly defined intracytoplasmic inclusions and vacuolated nuclei. Vasculitis is accompanied by thrombosis and 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 the upper respiratory tract are characterised by ulceration.

1.5. Immunological methods

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.

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 (LAMP). Nucleic acid recognition methods can be used to detect the *Capripoxvirus* genome in biopsy, swab or tissue culture samples.

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). A conventional PCR assay that differentiates GTPV and LSDV from SPPV has been described (Lamien *et al.,* 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).

The conventional gel-based PCR method described below is a simple, fast and sensitive method for the detection of capripoxvirus genome in EDTA blood, semen or tissue culture samples (Tuppurainen *et al.,* 2005).

Test procedure

The extraction method described below can be replaced using commercially available DNA extraction kits.

i)   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.

ii) 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*

220		for 15 minutes at 4°C. Carefully collect the upper, aqueous phase (up to 200 µl) and transfer
221		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 µl) 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   be adjusted to the final volume of 50 µl.

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   Electrophoretically separate the products using approximately 8–10 V/cm for 40–60 minutes
243   and visualise with a suitable DNA stain and transilluminator.

1.6.2. Real-time PCR methods

245 Several highly sensitive and specific fluorescent detection-based real-time PCR methods have been
246 developed and validated (Balinsky *et al*., 2008; Bowden *et al.,* 2008; Das *et al.,* 2012; Stubbs *et al.,*
247 2012). Each test detects a small conserved genetic locus within the capripoxvirus genome, but these
248 methods do not discriminate between SPPV, GTPV or LSDV. Real-time PCR methods for direct
249 capripoxvirus genotyping species differentiation without the need for gene sequencing have been
250 described (Gelaye *et al.,* 2013; Lamien *et al.,* 2011b; Wolff *et al., 2021*).

251 The real-time PCR method described below is a rapid, sensitive and specific method for the detection
252 of DNA from SPPV, GTPV or LSDV. This assay will not differentiate between capripoxvirus species.

DNA extraction from blood and tissue

254 A number of DNA extraction kits are commercially available for the isolation of template DNA for real-
255 time PCR. Manufacturer's instructions should always be consulted for guidance on the appropriate
256 method for the sample type being extracted. WOAH Reference Laboratories can be contacted for
257 advice on suitable commercial kits.

Real-time PCR

259 i) The real-time PCR method outlined below uses the primers and probe described by Bowden *et*
260   *al.* (2008). and further validated by Stubbs *et al.* (2012). Cycling conditions and reagent
261   concentrations can be altered to ensure optimal performance in individual laboratories.

262 ii) Forward and reverse primers should be prepared at concentrations of 20 µM. A minor grove
263   binder (MGB) TaqMan probe should be prepared at a concentration of 10 µM.

264 iii) Forward primer: 5'-AAA-ACG-GTA-TAT-GGA-ATA-GAG-TTG-GAA-3'

265 iv) Reverse primer: 5'-AAA-TGA-AAC-CAA-TGG-ATG-GGA-TA-3'

266 v) Probe: 5'-FAM-TGG-CTC-ATA-GAT-TTC-CT-MGB-3'

267 vi) Mastermix is prepared by combining 10 µl of 2 × real-time PCR mastermix with 0.4 µl of forward
268   primer, 0.4 µl of reverse primer, 0.5 µl of probe and 6.7 µl of RNase free water per reaction.

269 vii) Add 2 µl of extracted DNA to 18 µl of mastermix in a 96-well PCR plate or PCR strip and perform
270   real-time PCR according to the example given below or similar method:

	viii)	95°C for 10 minutes followed by 45 cycles of 95°C for 15 seconds, 60°C for 60 seconds. Fluorescence detection should be performed at the end of each cycle.

	ix)	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 negative.

1.6.3. Isothermal genome amplification

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).

2. Serological tests

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 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 difficulty of ensuring the use of 100 TCID$_{50}$, the neutralisation index is the preferred method, although it does require 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 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).

2.1.1. Test procedure

	i)	Test sera including a negative and a positive control are diluted 1/5 in Eagle's/HEPES (N-2-hydroxyethylpiperazine, N-2-ethanesulphonic acid) 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 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 log$_{10}$ 6 TCID$_{50}$ per ml is diluted in Eagle's/HEPES in bijoux bottles to give a log dilution series of log$_{10}$ 5.0; 4.0; 3.5; 3.0; 2.5; 2.0; 1.5 TCID$_{50}$ per ml (equivalent to log$_{10}$ 3.7; 2.7; 2.2; 1.7; 1.2; 0.7; 0.2 TCID$_{50}$ 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 An appropriate cell suspension (such as MDBK cells) is prepared from pregrown monolayers as a suspension of 10^5 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		can be made more sensitive if serum from the same animal is examined before and after
325		infection. Because immunity to capripoxvirus is predominantly cell mediated, a negative result,
326		particularly following vaccination in which the response is necessarily mild, does not imply that
327		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
330 capripoxvirus than LT cells, the problem of virus 'breakthrough' is overcome.

331 ### 2.2. Indirect fluorescent antibody test

332 Capripoxvirus-infected tissue culture grown on flying cover-slips or tissue culture microscope slides can be used for
333 the indirect fluorescent antibody test. Uninfected tissue culture control, and positive and negative control sera, should
334 be included in the test. The infected and control cultures are fixed in acetone at –20°C for 10 minutes and stored at
335 4°C. Dilutions of test sera are made in PBS, starting at 1/5, and positives are identified using an anti-sheep gamma-
336 globulin conjugated with fluorescein isothiocyanate (Davies & Otema, 1978). Cross-reactions can occur with orf,
337 bovine papular stomatitis virus and perhaps other poxviruses.

338 ### 2.3. Western blot analysis

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.

344 Both in-house and commercial enzyme-linked immunosorbent assay (ELISAs) are available, but these tests cannot
345 discriminate 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

350 A variety of attenuated live and inactivated capripoxvirus vaccines has been used to provide protection against
351 sheeppox and goatpox. All strains of capripoxvirus of ovine, caprine or bovine origin examined so far share a major
352 neutralising site, so that animals recovered from infection with one strain are resistant to infection with any other strain
353 (Capstick, 1961). Consequently, it is possible to use a single strain of capripoxvirus to protect both sheep and goats
354 against all field strains of virus, regardless of whether their origin was in Asia or Africa (Kitching *et al.*, 1986; Kitching
355 & Taylor, 1985). However, field evidence suggests some strains are quite host-specific and are used only in sheep
356 against SPPV and only in goat against GTPV.

357 A number of strains of capripoxvirus have had widespread use as live vaccines (Davies & Mbugwa, 1985), for
358 example the Romanian and RM-65 strains used mainly in sheep and the Mysore and Gorgan strains used in goats.
359 The real identity of the commonly used Kenyan sheep and goat pox vaccine virus (KSGP) 0240 was recently shown
360 to be actually LSDV (Tuppurainen *et al.,* 2014). Virus strain identity and attenuation properties must be ascertained
361 and taken into consideration when selecting vaccine strains for use in cattle, sheep and goats. The protective dose
362 depends on the vaccine strain used. Immunity in sheep and goats against capripoxvirus following vaccination with
363 the 0240 strain lasts over a year and the Romanian strain gave protection for at least 30 months.

364 Killed vaccines produced from tissue culture contain only the intracellular mature virion form of the virus, and lack the
365 less robust but biologically crucial extracellular enveloped virion form. As a result, the vaccine does not stimulate
366 immunity against the extracellular enveloped virion, resulting in poor protection. Killed capripoxvirus vaccines provide,
367 at best, only temporary protection.

2. Outline of production and minimum requirements for conventional vaccines

General requirements set for the facilities used for the production of vaccines and for the documentation and record keeping 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 for the testing of cells and reagents used in the process, each batches and the final product.

2.1. Characteristics of the seed

2.1.1. Biological characteristics

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 for which it is intended, including pregnant and young animals. It 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. A quantity of master seed vaccine virus should be 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

The method of manufacture should be documented as the Outline of Production.

2.2.1. Procedure

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.

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.

The procedure is repeated and the harvests from individually numbered flasks are each mixed separately with an equal volume of sterile and chilled 5% lactalbumin hydrolysate and 10% sucrose, and transferred to individually numbered bottles for storage at –20°C. Prior to storage, 0.2 ml is removed from each bottle for sterility control. An additional 0.2 ml is removed for virus titration; 2 ml pools composed of 0.2 ml samples taken from ten bottles are used. A written record of all the procedures must be kept for all vaccine batches.

Inactivated vaccines are produced, usually from unattenuated field strains of capripoxvirus, grown in tissue culture as described above, inactivated with 0.03% formaldehyde, and mixed with an equal volume of alhydrogel as adjuvant. Formaldehyde is no longer considered to be a suitable inactivant 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.

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.

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 2×10^7 cells/ml in sterile 10% DMSO (dimethyl sulphoxide) and 90% FBS (fetal bovine serum) solution stored in liquid nitrogen.

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.

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 $\log_{10}$ 4.5 $TCID_{50}$ per ml after freeze-drying, equivalent to a field dose of $\log_{10}$ 2.5 $TCID_{50}$. A further titration is carried out on five randomly chosen vials of the freeze-dried preparation to confirm the titre.

2.2.4. Final product batch tests

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.

iii) Potency

Potency tests must be undertaken if the minimum immunising dose of the virus strain is not known. This is usually carried out by comparing the titre of a virulent challenge virus on the flanks of vaccinated and control animals. Following vaccination, the flanks of at least three animals and three controls are shaved of wool or hair. $\log_{10}$ dilutions of the challenge virus are prepared in sterile PBS and six dilutions are inoculated intradermally (0.1 ml per inoculum) along the length of the flank; four

466	replicates of each dilution are inoculated down the flank. An oedematous swelling will develop at
467	possibly all 24 inoculation sites on the control animals, although preferably there will be little or no
468	reaction at the four sites of the most dilute inocula. The vaccinated animals should develop an initial
469	hypersensitivity reaction at sites of inoculation within 24 hours, which should quickly subside. Small
470	areas of necrosis may develop at the inoculation site of the most concentrated challenge virus. The
471	macule/papule is measured at between 8 and 10 days post-challenge. The titre of the challenge virus
472	is calculated for the vaccinated and control animals; a difference of $\log_{10}$ titre > 2.5 is taken as
473	evidence of protection.

2.3. Requirements for authorisation

2.3.1. Safety requirements

i) Target and non-target animal safety

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.

Safety tests should be carried out on the final product of each batch as described in Section C.2.2.4.

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.

ii) Reversion-to-virulence for attenuated/live vaccines

The selected final vaccine should not revert to virulence during a further passages in target animals.

iii) Environmental consideration

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 *Bos taurus* 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.

2.3.2. Efficacy requirements

i) For animal production

The efficacy of the vaccine must be demonstrated in vaccination challenge experiment under laboratory conditions. As described in Section C.2.2.4.

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.

ii) For control and eradication

Vaccination is the only effective way to control the sheep pox and goat pox outbreaks in endemic countries. Unfortunately, currently no marker vaccines allowing the differentiation of infected from vaccinated animals are available.

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 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 there is no possibility of field strains of capripoxvirus confusing the results. The inactivated vaccines 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.

2.3.3. Stability

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.

513   Properly freeze-dried preparations of capripox vaccine, particularly those that include a protectant,
514   such as sucrose and lactalbumin hydrolysate, are stable for over 25 years when stored at –20°C and
515   for 2–4 years when stored at 4°C. There is evidence that they are stable at higher temperatures, but
516   no long-term controlled experiments have been reported. The inactivated vaccines must be stored at
517   4°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.

3. Vaccines based on biotechnology

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).

3.2. Special requirements for biotechological vaccines, if any

526 Not applicable.

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590 * *

591 **NB:** There are WOAH Reference Laboratories for sheep pox and goat pox (please consult the WOAH Web site:
592 https://www.woah.org/en/what-we-offer/expertise-network/reference-laboratories/#ui-id-3).
593 Please contact the WOAH Reference Laboratories for any further information on
594 diagnostic tests, reagents and vaccines for sheep pox and goat pox

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

2   **MEETING OF THE BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # S E C T I O N   3 . 9 .

6   # S U I D A E

7   ## C H A P T E R   3 . 9 . 1 .

8
AFRICAN SWINE FEVER
9   # (INFECTION WITH AFRICAN SWINE FEVER VIRUS)

10   ## SUMMARY

11   *African swine fever (ASF) is an infectious disease of domestic and wild pigs of all breeds and ages, caused*
12   *by ASF virus (ASFV). The clinical syndromes vary from peracute, acute, subacute to chronic, depending on*
13   *the virulence of the virus. Acute disease is characterised by high fever, haemorrhages in the*
14   *reticuloendothelial system, and a high mortality rate. Soft ticks of the* Ornithodoros *genus, especially* O.
15   moubata *and* O. erraticus*, have been shown to be both reservoirs and transmission vectors of ASFV. The*
16   *virus is present in tick salivary glands and passed to new hosts (domestic or wild suids) when feeding. It can*
17   *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*.*

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.*

22   ***Identification of the agent:*** *Laboratory diagnosis must be directed towards isolation of the virus by*
23   *inoculation of pig leukocyte or bone marrow cultures, the detection of antigen in smears or cryostat sections*
24   *of tissues by fluorescent antibody test and/or the detection of genomic DNA by the polymerase chain reaction*
25   *(PCR) or real-time PCR. The PCRs are excellent, highly sensitive, specific and rapid techniques for ASFV*
26   *detection and are very useful under a wide range of circumstances. They are especially useful if the tissues*
27   *are unsuitable for virus isolation and antigen detection. In doubtful cases, the material is passaged in*
28   *leukocyte cell cultures and the procedures described above are repeated.*

29   ***Serological tests:*** *Pigs that survive natural infection usually develop antibodies against ASFV from 7–10*
30   *days post-infection and these antibodies persist for long periods of time. Where the disease is endemic, or*
31   *where a primary outbreak is caused by a strain of low or moderate virulence, the investigation of new*
32   *outbreaks should include the detection of specific antibodies in serum or extracts of the tissues submitted.*
33   *A variety of methods such as the enzyme-linked immunosorbent assay (ELISA), the indirect fluorescent*
34   *antibody test (IFAT), the indirect immunoperoxidase test (IPT), and the immunoblotting test (IBT) is available*
35   *for antibody detection.*

36   ***Requirements for vaccines***: ~~At present, there is no vaccine for ASF.~~ <u>Commercially produced</u> <u>modified live</u>
37   <u>virus vaccines are available and licenced in some countries.</u>

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.

ASF virus (ASFV) is a complex large, enveloped DNA virus with icosahedral morphology. It is currently classified as the only member of the *Asfaviridae* family, genus *Asfivirus* (Dixon *et al.*, 2005). More than 60 structural proteins have been 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 pigs (Sánchez-Vizcaíno & Arias, 2012). The ASFV double-stranded linear DNA genome comprises between 170 and 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 genome. The complete genomes of several ASFV strains have been sequenced (Bishop *et al.,* 2015; Chapman *et al.,* 2011; de Villiers *et al.,* 2010; Portugal *et al.,* 2015). Different strains of ASFV vary in their ability to cause disease, but at present there is only one recognised serotype of the virus detectable by antibody tests.

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*., 2017; Boshoff *et al.,* 2007; Quembo *et al.* 2018). To distinguish subgroups among closely related ASFV, sequence analysis 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 end of the genome (Gallardo *et al.,* 2014), is undertaken. Several other gene regions such as the E183L encoding p54 protein, the CP204L encoding p30 protein, and the protein encoded by the EP402R gene (CD2v), have been proved as useful tools to analyse ASFVs from different locations and hence track virus spread.

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).

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, sometimes even before the first clinical signs are observed. Case fatality rates may be as high as 100%. Less virulent 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 variable disease forms, ranging from acute to subacute. Low virulence, non-haemadsorbing strains can produce subclinical non-haemorrhagic infection and seroconversion, but some animals may develop discrete lesions in the lungs or on the skin in areas over bony protrusions and other areas subject to trauma. Animals that have recovered from either acute, 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 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 ASFV 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
100　live virus (MLVs) vaccines are based on the live virus that have been naturally attenuated or attenuated by targeted genetic
101　recombination through cell cultures (Gladue & Borca, 2022). MLV production is based on a seed-lot system consistent with
102　the *European Pharmacopoeia* (11th edition) and that has been validated with respect to virus identity, sterility, purity,
103　potency, safety, non-transmissibility, stability and immunogenicity. ASF MLV first generation vaccines – defined as those
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.

108　ASF MLV first generation vaccines allowing the differentiation of infected animals from vaccinated animals (DIVA) by
109　suitable methods (e.g. serology-based tests) are preferred. Demonstration of MLV safety and efficacy in breeding-age
110　boars, gilts and pregnant sows, and onset and duration of protective immunity, are also preferred but are not required to
111　meet the minimum standard.

112　ASF epidemiology is complex with different epidemiological patterns of infection occurring in Africa and Europe. ASF
113　occurs through transmission cycles involving domestic pigs, wild boar, wild African suids, and soft ticks (Sánchez-Vizcaíno
114　*et al.,* 2015). In regions where *Ornithodoros* soft-bodied ticks are present, the detection of ASFV in these reservoirs of
115　infection contributes to a better understanding of the epidemiology of the disease. This is of major importance in
116　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　　　　　　　　　　　　　　　　　　　　　. . .

C. REQUIREMENTS FOR VACCINES

123　~~At present there is no commercially available vaccine for ASF.~~

1.　Background

125　The ASF p72 genotype II strain (ASFV Georgia 2007/1 lineage) (NCBI, 2020) is recognised to be the current highest global
126　threat for domestic pig production worldwide (Penrith *et al.,* 2022).

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.

130　Wherever live, virulent ASFV or ASF MLVs are stored, handled and disposed, the appropriate biosecurity level, procedures
131　and practices should be used. The ASF MLV vaccine production facility should meet the requirements for containment
132　outlined in Chapter 1.1.4 *Biosafety and biosecurity: Standard for managing biological risk in the veterinary laboratory and*
133　*animal facilities.*

134　An optimal ASF MLV first generation vaccine for the target host should have the following general characteristics (minimum
135　standards):

136 · Safe: demonstrate absence of fever and clinical signs of acute or chronic ASF in vaccinated and in-contact animals,
137 minimal and ideally no vaccine virus transmission, and absence of an increase in virulence (genetic and phenotypic
138 stability);

139 · Efficacious: protects against mortality, reduces acute disease (fever accompanied by the appearance of clinical signs
140 caused by ASF) and reduces vertical (boar semen and placental) and horizontal disease transmission;

141 · Quality – purity: free from wild-type ASFV and extraneous microorganisms that could adversely affect the safety,
142 potency or efficacy of the product;

143 · Quality – potent: the $\log_{10}$ virus titre maintained throughout the vaccine shelf life that guarantees the efficacy
144 demonstrated by the established minimum immunising (protective) dose.

145 · Identity: based on the capacity to protect against the ASFV B646L (p72) genotype II pandemic strain or other p72
146 genotypes of recognised epidemiologic importance.

147 Vaccine production should be carried out using a validated, controlled and consistent manufacturing process.

148 ASF MLV first generation vaccines must be safe (i.e. an acceptable safety profile) for non-target species and the
149 environment in general.

150 Ideally, ASF MLV first generation vaccines that meet the minimum standards should also fulfil the following additional
151 general characteristics: i) prevents acute and persistent (carrier state) disease; ii) prevents horizontal and vertical disease
152 transmission; iii) induces rapid protective immunity (e.g. < 2 weeks); and iv) confers stable, life-long immunity.

153 Furthermore, ASF MLV second and future generation vaccines should meet the minimum safety and efficacy standards
154 as ASF MLV first generation vaccines, and ideally provide additional product profile benefits, including but not limited to: i)
155 contain a negative marker allowing the differentiation of infected from vaccinated animals (DIVA) by reliable discriminatory
156 tests such as serology-based tests; and ii) confer broad range of protection against other p72 genotype field strains of
157 varying virulence (low, moderate, and high).

158 The majority of ASF global vaccine research groups and companies are currently focused on ASF MLV first generation
159 vaccine candidates that are safe and efficacious against ASF viruses belonging to the ASFV p72 genotype II pandemic
160 strain (ASFV Georgia 2007/1 lineage) (NCBI, 2020).

161 Currently, two gene deleted MLV recombinant vaccines (ASFV-G-ΔI177L and ASFV-G-ΔMGF) have been licenced for
162 field use in Vietnam following supervised field testing to evaluate the safety and effectiveness of several vaccine batches.

163 There are numerous, promising ASF MLV vaccine candidates targeting the p72 genotype II pandemic strain under
164 development, including:

165 · A naturally attenuated field strain (Lv17/WB/Rei1) (Barasona *et al.,* 2019) being developed as an oral bait vaccine for
166 wild boars;

167 · A laboratory thermo-attenuated field strain (ASFV-989) (Bourry *et al.,* 2022);

168 · Single gene-deleted, recombinant viruses (e.g. SY18ΔI226R, ASFV-G-ΔA137R) (Gladue *et al.,* 2021; Zhang *et al.,*
169 2021);

170 · Double gene-deleted, recombinant viruses (e.g. ASFV-G-Δ9GL/ΔUK; ASFV-SY18-ΔCD2v/UK; Arm-ΔCD2v-ΔA238L)
171 (O'Donnell *et al.,* 2016; Pérez-Núñez *et al.,* 2022; Teklue *et al.,* 2020);

172 · Multiple gene-deleted, recombinant viruses (ASFV-G-ΔI177L/ΔLVR; ASFV-G-ΔMGF; BA71ΔCD2; HLJ/18-7GD;
173 ASFVGZΔI177LΔCD2vΔMGF) (Borca *et al.,* 2021; Chen *et al.,* 2020; Liu *et al.,* 2023; Monteagudo *et al.,* 2017;
174 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.

179 Inactivated (non-replicating) whole virus vaccines are not presently available and may be difficult to develop to meet
180 minimum efficacy standards. Recombinant vectored, subunit vaccine candidates that can be produced in scalable vaccine
181 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 *Genomics* (ASFV Genomics, 2022[43]) that provides the structural protein predictions for all 193 ASFV proteins may help
184 accelerate ASF first and second generation vaccine research and development.

185 Fit-for-purpose vaccine use scenarios matched to the intended use in a domestic pig specific type of production system
186 may require different vaccine product profiles or may influence the focus of essential versus ideal vaccine requirements.
187 As with any MLV vaccine, all ASF MLV vaccines should be used according to the label instructions, under the strict control
188 of the country's Regulatory Authority.

189 The minimum standards given here and in chapter 1.1.8 are intended to be general in nature and may be supplemented
190 by national, regional, and veterinary international medicinal product harmonised requirements. Minimum data requirements
191 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.

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 **2.1.2.   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).

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 yields (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.

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43 http://asfvgenomics.com. Accessed 4/4/2023.

2.1.3. Validation as a vaccine strain

The vaccine derived from the MSV must be shown to be satisfactory with respect to safety and efficacy.

Even if pigs are not known for susceptibility to transmissible spongiform encephalopathy (TSE) agents, consideration should also be given to minimising the risk of TSE transmission by ensuring 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.

Ideally, the vaccine virus in the final product should generally not differ by more than five passages from the master seed lot.

ASF vaccine should be presented in a suitable pharmaceutical form (e.g. lyophilisate or liquid form).

2.2. Method of manufacture

2.2.1. Procedure

The MLV virus is used to infect swine primary cell cultures obtained from specific-pathogen free pigs, the requirements for which are defined in specific monographs (Chapter 2.3.3 *Minimum requirements for the organisation and management of a vaccine manufacturing facility*, Section 2.4.2). Compared with primary cell cultures, use of a continuous cell line generally allows for more consistency, higher serial volumes in manufacturing and aligns better with a seed lot system. Thus, preferably a master cell bank based established, continuous cell line shown to support genetically stable ASFV replication and acceptable titres over several passages should be used.

Cell cultures shall comply with the requirements for cell cultures for production of veterinary vaccines in chapter 1.1.8. Regardless of the production method, the substrate should be harvested under aseptic conditions and may be subjected to appropriate methods to release cell-associated virus (e.g. freeze–thaw cycles, detergent lysis). The harvest can be further processed by filtration and other purification methods. A stabiliser or other excipients may be added as appropriate. The vaccine is homogenised to ensure a uniform batch/serial.

2.2.2. Requirements for ingredients

All ingredients used for vaccine production should be in line with requirements in chapter 1.1.8.

2.2.3. In-process controls

In-process controls will depend on the protocol of production: they include virus titration of bulk antigen and sterility tests.

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.

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

Batch safety testing is to be carried out unless consistent safety of the product is demonstrated and approved in the registration dossier and the production process is approved for consistency in accordance with the standard requirements referred to in chapter 1.1.8.

274   v) Batch/serial potency

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) Residual humidity/residual moisture

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.

2.3.   Requirements for authorisation/registration/licensing

2.3.1.   Manufacturing process

284 For regulatory approval of a vaccine, all relevant details concerning history of the pre-MSV,
285 preparation of MSV, manufacture of the vaccine and quality control testing (Sections C.2.1
286 *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 a volume not less than 1/3 of the typical industrial batch volume. The in-process controls are part of
290 the manufacturing process.

2.3.2.   Safety requirements

292 For the purpose of gaining regulatory approval, the following safety tests should be performed
293 satisfactorily.

294 As a minimum standard, vaccines should be tested for any pathogenic effects on healthy domestic
295 pigs of the target age intended for use. Additional demonstration of MLV safety in breeding age gilts
296 and pregnant sows is preferred but not required as a minimum standard.

297   i)   Safety in young animals

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$_{50}$], quantitative 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 cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive
315 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

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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:

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
324   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   •   No vaccinated pigs show notable signs of disease by gross pathology

327   ii)   Safety test in pregnant sows and test for transplacental transmission

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   GL44 (*Guidelines on Target Animal Safety for Veterinary Live and Inactivated Vaccines, Section 2.2.*
332   *Reproductive Safety Test*, 2009[46]) should be completed.

333   iii)   Horizontal transmission

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   days, preferably 60 days.

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.

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 (HAD_{50}/ml or $TCID_{50}$/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.

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46 https://www.ema.europa.eu/en/documents/scientific-guideline/vich-gl44-target-animal-safety-veterinary-live-inactived-vaccines-step-7_en.pdf.

The vacine complies with the test if:

- 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 attributable to the vaccine;

- The average body temperature increase for all naïve, contact 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;

- No naïve, contact piglet shows notable signs of disease by gross pathology and no virus is detected in their blood or tissue samples

- No naïve contact pigs test positive for antibodies to the vaccine virus.

iv)   Post-vaccination kinetics of viral replication (MLV blood and tissue dissemination) study

Prior to the reversion to virulence study (Section C2.3.2.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.

The test consists of the administration of the vaccine virus from the master seed lot to no fewer than eight healthy piglets, and preferably ten healthy piglets, a minimum of 6-weeks old and not older than 10-weeks old and of the same origin, that do not have antibodies against ASFV, and blood samples are negative on real-time PCR.

Administer to each piglet, using the recommended route of administration most likely to result in spread (such as the intramuscular route or intranasal route), a quantity of the master seed vaccine virus equivalent to not less than the maximum virus titre (maximum release dose) likely to be contained in 1 dose of the vaccine.

Record daily body temperatures and observe inoculated animals daily for clinical disease for at least 45 days, preferably 60 days.

Carry out the daily observations for signs of acute and chronic clinical disease using a quantitative clinical scoring system adding the values for multiple clinical signs (e.g. Gallardo *et al*. (2015a). These clinical signs should include fever, anorexia, recumbency, skin haemorrhage or cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive findings.

Collect blood samples from all the piglets at least two times per week from 3 days post-vaccination for the first 2 weeks, then weekly for the duration of the test. Determine vaccine virus titres by quantitative virus isolation (HAD_{50}/ml or $TCID_{50}$/ml) and using a real-time PCR test. If the vaccine virus is non-haemadsorbing or does not cause cytopathic effects, a real-time PCR test only may be used.

Determine which blood timepoint(s) should be used in the design of the reversion to virulence study (Section C2.3.2.v. below).Collect oral, nasal and faecal swab samples (preferably devoid of blood to minimise assay interference) at least two times per week from 3-days post-vaccination for the first 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) and using a real-time PCR test. If the vaccine 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.

Euthanise at least two piglets on days 7, 14, 21, and preferably on day 28 (±2 days at each timepoint) and collect spleen, lung, tonsil, kidney tissue samples and at least three different lymph nodes (which should include lymph node closest to site of inoculation, gastrohepatic and submandibular nodes). Determine virus titres in all collected samples by quantitative virus isolation (HAD_{50}/mg or $TCID_{50}$/mg) and using real-time PCR test. If the vaccine 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.

Determine which tissue(s) and timepoint(s) should be used to aid in the design of the reversion to virulence study (Section C.2.3.2.v), for example, specific tissues at specific timepoints which show the highest titres should be considered for selection and use in the reversion to virulence study.

| 417 | | v) | Reversion to virulence |

The test should be carried out consistent with VICH GL41 (Examination of live veterinary vaccines in target animals for absence of reversion to virulence, 2008[47]).

The test for increase in virulence consists of the administration of the vaccine master seed virus to healthy piglets of an age (e.g. between 6-weeks and 10-weeks old) suitable for recovery of the strain and of the same origin, that do not have antibodies against ASFV, and blood samples that are negative on real-time PCR. This protocol is typically repeated five times.

First pass (p1)

Administer to no fewer than two piglets, and preferably no fewer than four piglets using the intended route of administration for the final product, a quantity of the master seed vaccine virus equivalent to not less than the maximum virus titre (maximum release dose) likely to be contained in 1 dose of the vaccine. Observe inoculated animals daily for the appearance of at least two and preferably at least three clinical signs and record daily body temperatures.

Based on results from at least one completed vaccine shed and spread (virus blood and tissue dissemination study, Section C.2.3.2.iv above) collect an appropriate quantity of blood from each piglet on the predetermined single timepoint (day 5–13). Determine virus titres in individual blood samples by quantitative virus isolation (HAD_{50}/ml or $TCID_{50}$/ml) and by real-time PCR. If the vaccine 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 blood sample(s) with the highest infectious titre and reserve for the subsequent *in-vivo* passage (second pass, p2).

Based on results from at least one completed vaccine virus blood and tissue distribution dissemination study, Section C.2.3.2.iv above) euthanise piglets on the predetermined timepoint (i.e. day 7, 14, 21, or 28). Determine infectious virus titres in individual tissue samples by quantitative virus isolation (HAD_{50}/ml or $TCID_{50}$/ml). If the vaccine 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 individual tissue sample(s) with the highest infectious titre. Pool the tissues from different organs from all animals with the highest titres and prepare at least a 10% suspension in PBS, pH 7.2 kept at 4°C or at –70°C for longer storage. Test each blood and tissue pool used for inoculation by PCR to confirm the absence of potential viral agent contaminants (i.e. CSFV, FMDV, PRRS, PCV2). Blood and pooled tissue (p1) are used to inoculate 2 ml of positive material using the intended route of administration for the final product to each of least two and ideally at least four further pigs of the same age and origin.

Second pass (p2)

If no virus is found (p1), repeat the administration by the intended route once again with the same pooled material (blood and pooled tissue, p1) in another ten healthy piglets of the same age and origin.

If no virus is found at this point, end the process here. If, however, virus is found, carry out a second series of passages by administering 2 ml of positive material using the intended route of administration for the final product to each of no fewer than two piglets, and preferably no fewer than four piglets of the same age and origin. Observe inoculated animals daily for the appearance of at least two and preferably at least three clinical signs and record daily body temperatures.

Third and fourth pass (p3 and p4)

If no virus (p2), repeat the intramuscular administration once again with the same pooled material (blood and pooled tissue, p2) in another eight healthy piglets of the same age and origin.

If no virus is found at this point, end the process here. If, however, virus is found, carry out this passage operation no fewer than two additional times (p3 and p4) (to each of no fewer than two piglets, and preferably no fewer than four piglets of the same age and origin) and verifying the presence of the virus at each passage in blood and tissues. Observe inoculated animals daily for the

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47 https://www.ema.europa.eu/en/documents/scientific-guideline/vich-gl41-target-animal-safety-examination-live-veterinary-vaccines-target-animals-absence-reversion_en.pdf.

| 466 | appearance of at least two and preferably at least three clinical signs and record daily body |
| 467 | temperatures. |

| 468 | *Fifth pass (p5)* |

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 | • No piglet shows abnormal local or systemic reaction, reaches the pre-determined humane end |
| 474 | 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	accompanied by clinical sign observations) of the maximally passaged virus compared with the
477	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	mean) for the observation period does not exceed 1.5°C above baseline; and no individual piglet
481	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	minimal chronic clinical signs (defined as mild swollen joints with a low clinical score that resolve
484	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 | • Absence of an increase in virulence (genetic and phenotypic stability) (complies with the reversion |
| 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 | ### 2.3.3. Efficacy requirements |

| 498 | i) Protective dose |

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	time PCR. The test is conducted using no fewer than 15 and preferably no fewer than 24 vaccinated
503	pigs, and no fewer than five 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 administration of the challenge virus, at the time of challenge, 4 hours after challenge, and then daily
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 (HAD_{50}/ml or $TCID_{50}$/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.

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
554 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 Re=R0 × (S/N) (S= susceptible pigs; N= total number of pigs in a given population) is greater than
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.

566 To evaluate ASF vaccine impact on ASF disease transmission, the test consists of a
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.

569 The test is conducted using no fewer than 15 healthy piglets at a ratio comprising twice the number of vaccinated piglets to naïve piglets (e.g. ten vaccinated and five naïve). Use vaccine containing 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 virus titre (minimum dose) likely to be contained in one dose of the vaccine as stated on the label. 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 vaccine then 28 days [± 2 days] following the second injection), temporarily separate [into different pen(s) or room(s)] all vaccinated piglets from naïve piglets. Challenge all vaccinated piglets by the intramuscular or other previously verified route. Carry out the challenge using an ASFV representative strain of the epidemiologically relevant field strain(s) where the vaccine is intended for 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 challenge virus type is available, then carry out the test with the parental, virulent virus used to generate the MLV recombinant virus. Use a 10e3–10e4 HAD_{50} (or $TCID_{50}$ for non-HAD viruses challenge dose sufficient to cause death or met the humane endpoint in 100% of the nonvaccinated piglets in less than 21 days. Higher or lower challenge doses can be considered if appropriately justified.

587 Approximately 18-24 hours later, re-introduce naïve piglets to vaccinated, challenged piglets and allow for direct nose to nose contact exposure with vaccinated, challenged piglets. Allow for continuous contact exposure by co-mingling both groups through the end of the study. If more than one pen or room is used for co-housing, following reintroduction initially maintain a ratio of 2:1 of 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 administration of the challenge virus to vaccinated pigs, immediately prior to direct contact exposure, 4 hours post-contact exposure, and then daily for at least 28, preferably 35 days. Observe all contact 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 using a quantitative clinical scoring system adding the values for multiple clinical signs (e.g. Gallardo _et al._, 2015a). These clinical signs should include fever, anorexia, recumbency, skin haemorrhage or cyanosis, joint swelling and necrotic lesions around the joints, respiratory distress and digestive findings.

601 In addition, blood should be taken from the naïve contact piglets at least twice a week from 3 days post-contact exposure for the duration of the test period. From the blood samples, determine infectious challenge virus titres by quantitative virus isolation (HAD_{50}/ml or $TCID_{50}$/ml) and using a real-time PCR test. If the vaccine virus is non-haemadsorbing or does not cause cytopathic effects, 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), and at the end of the test period, and carry out an appropriate test to detect vaccine virus antibodies.

608 Collect oral, nasal and faecal swab samples (preferably devoid of blood to minimise assay interference) from all contact-exposed naïve piglets at least two times per week from 3-days post-contact exposure for the first 2 weeks, then weekly for the duration of the test and test swabs for the presence of challenge virus. Determine virus titres in all collected samples by quantitative virus isolation (HAD_{50}/ml or $TCID_{50}$/ml) and using a real-time PCR test. If the vaccine 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.

615 At the end of the test period, humanely euthanise all contact piglets. Conduct gross pathology on spleen, lung, tonsil, and kidney tissue samples and at least three different lymph nodes. (which should include lymph node closest to site of inoculation, gastrohepatic and submandibular nodes). Determine virus titres in all collected samples by quantitative virus isolation (HAD_{50}/mg or $TCID_{50}$/mg) and real-time PCR (see Section B.1. Identification of the agent). If the vaccine 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.

| 622 | The test is invalid if the vaccine fails to comply with the compliance criteria described for the protected |
| 623 | 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 • No naïve, contact exposed piglet shows abnormal (local or systemic) reactions, reaches the
626 defined humane endpoint or dies from causes attributable to ASF;

627 • No naïve, contact exposed piglet displays fever accompanied by typical signs of disease,
628 including gross pathology.

629 • Naïve contact pigs show a reduction or absence of challenge virus levels in blood and tissues.

630 • None of or a reduced number of naïve contact exposed pigs test positive for antibodies to the
631 challenge virus.

632 At a minimum, an efficacious MLV vaccine shall demonstrate ALL the following features (minimal
633 standards):

634 • Protects against mortality;

635 • Reduces acute disease (fever accompanied by a reduction of typical clinical and pathological
636 signs of acute disease)

637 • Reduces horizontal disease transmission (no naïve, contact exposed piglet shows abnormal
638 [local or systemic] reactions, reaches the humane endpoint or dies from causes attributable to
639 ASF, and displays fever accompanied by typical acute disease signs caused by ASF)

640 • Reduces levels of viral shedding and viraemia.

641 In addition, the vaccines in their commercial presentation before being authorised for general use
642 should be tested for efficacy in the field (see chapter 1.1.8 Section 7.2.3). Additional field efficacy
643 evaluation studies may include but are not limited to: onset of immunity, duration of immunity, and
644 impact on disease transmission.

2.3.4. Duration of immunity

646 Although not included in the guidance for ASF MLV first generation vaccines, manufacturers are
647 encouraged as part of the authorisation procedure, to demonstrate the duration of immunity of a given
648 vaccine by evaluation of potency at the end of the claimed period of protection.

2.3.5. Stability

650 Stability of the vaccine should be demonstrated over the shelf life recommended for the product.
651 Although not included in the standards for first generation MLV ASF vaccines, manufacturers are
652 encouraged, as part of the authorisation procedure, to generate data supporting the period of validity
653 of a lyophilised or other pharmaceutical form of the ASF vaccine as part of the authorisation
654 procedure.

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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.

823 **Appendix 1 to Annex 16 of the Report of the Biological Standards Commission meeting**
824 **4–8 September 2023**
825
826 **Draft Standards for African Swine Fever (ASF) Modified Live Virus (MLV)**
827 **Vaccines for Domestic and Wild Pigs**

Annex16_3.09.01_ap
pendix1_GL_ASF_MLV

828

2   **MEETING OF THE BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   ——————————

5   # C H A P T E R 3 . 1 0 . 4 .

6   # INFECTION WITH
7   # *CAMPYLOBACTER JEJUNI* AND *C. COLI*

8   ## SUMMARY

9   ***Description of the disease:*** Campylobacter jejuni (C. jejuni) *and* Campylobacter coli (C. coli) *can colonise*
10   *the intestinal tract of most mammals and birds and are the most frequently isolated* Campylobacter *species*
11   *in humans with gastroenteritis. Although poultry is the main reservoir of* Campylobacter, *transmission to*
12   *humans is only partly through handling and consumption of poultry meat; other transmission routes are also*
13   *considered to be important. This chapter focuses on* C. jejuni *and* C. coli *in primary livestock production with*
14   *regard to food safety.*

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*
22   *reaction (*PCR*).* Campylobacter jejuni *and* C. coli *are thermophilic, Gram-negative, highly motile bacteria*
23   *that, for optimal growth, require microaerobic environment and incubation temperatures of 37–42°C. Agar*
24   *media containing selective antibiotics are required to isolate these bacteria from faecal/intestinal samples.*
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*
28   *phase are short and S-shaped in appearance, while coccoid forms predominate in older cultures. Under*
29   *phase-contrast microscopy the organisms have a characteristic rapid corkscrew-like motility. Phenotypic*
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*
33   *for the direct detection of* C. jejuni *and* C. coli*.*

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

1. Disease

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, but the organisms are also found in healthy animals. One specific *C. jejuni* clone has been associated with abortion in 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 agent of spotty liver disease in layers (Crawshaw *et al.,* 2015).~~ *Campylobacter jejuni* and *C. coli* are of interest mainly from 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* together account for more than 90% of all human campylobacteriosis cases. Over 80% of cases are caused by *C. jejuni* and about 10% of cases are caused by *C. coli*. In humans, *C. jejuni/C. coli* infection is associated with acute enteritis and 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 shown ~~that~~ the majority of campylobacteriosis cases in humans can be attributed to poultry and a smaller fraction to cattle (Mughini-Gras *et al.,* 2012) ~~is the main reservoir of *Campylobacter* and responsible for between 50 and 80% of the human infections~~. In the European Union (EU), an estimated ~~30~~ 20–40% of the human infections are associated with handling and 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, e.g. via ~~environmental contamination~~ surface water (~~EFSA, 2010b,~~ Mulder *et al.,* 2020). Contact with pets and livestock, the consumption of contaminated water or raw milk and travelling ~~in high prevalence areas~~ are also considered risks factors in human disease (Domingues *et al.,* 2012; Mughini-Gras *et al.,* 2021). The control of *Campylobacter* in the food chain has ~~now~~ become a major target of agencies responsible for food safety world-wide.

Laboratory manipulations should be performed 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 diagnostic laboratory and animal facilities*).

2. Taxonomy

There are currently ~~34~~ 43 *Campylobacter* species recognised (July 2023), but with the improved diagnostic techniques and genomic analysis, this number is expected to increase over time (~~cf~~ List of prokaryotic names with standing in nomenclature: (https://lpsn.dsmz.de/genus/campylobacter ~~http://www.bacterio.net/index.html~~). Members of the genus *Campylobacter* are typically Gram-negative, non-spore-forming, S-shaped or spiral shaped bacteria (0.2–~~0.8~~ 0.5 µm wide and 0.5–~~5~~ 8 µm long), with single polar flagella at one or both ends, conferring a characteristic corkscrew-like motility. ~~These bacteria~~ *Campylobacter* require~~s~~ microaerobic conditions, but some strains also grow aerobically or anaerobically. They neither ferment nor oxidise carbohydrates. Some species, particularly *C. jejuni*, *C. coli* and *C. lari*, are thermophilic, 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 be discriminated on the basis of several phenotypic tests, but this subspeciation has no added value for epidemiological 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*.~~

B. DIAGNOSTIC TECHNIQUES

Table 1. Test methods available for the diagnosis of Campylobacter jejuni and C. coli and their purpose

Method	Purpose[a]					
	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
Agent identification[b]						
Isolation	+++	–	= +++	+++	+++	–
~~**MALDI-TOF**~~	+++	–	+++	+++	+++	–

Method	Purpose[a]					
	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
Antigen detection	++	–	= ~~++~~	–	++~~+~~	–
~~16S rRNA sequencing~~	++	–	++	++	++	–
Real-time PCR	++~~+~~	–	= ~~++~~	++	++~~+~~	–
Detection of immune response: n/a for *Campylobacter jejuni* and *C. coli*						

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.

1. Isolation and identification of the agent

Two ISO (International Organization for Standardization) procedures for detection of *Campylobacter* exist. ISO 10272 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 (ISO 10272-1:2017) and part 2 colony count technique (ISO 10272-2:2017). ~~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

Poultry is frequently colonised with *C. jejuni* (65–95%), less often with *C. coli* and rarely with other *Campylobacter* species (~~Newell & Wagenaar, 2000~~ Wagenaar *et al.*, 2023). Colonisation rates in broiler chickens are age-related. Most flocks are negative until 2 weeks of age. Once *Campylobacter* colonisation occurs in a broiler flock, transmission, via exposure to faecal contamination, is extremely rapid and up to 100% of birds within a flock can become colonised within a few days. Samples from live birds, destined for the food chain, should therefore be taken as close to slaughter as possible (~~Newell & Wagenaar, 2000~~ Wagenaar *et al.*, 2023). The majority of birds shed large numbers of organisms (>10^6 colony-forming units/g faeces). Campylobacters can be isolated from fresh faeces/caecal droppings or cloacal swabs. For reliable detection of *Campylobacter* by culture, freshly voided faeces (preferably without traces of urine) should be collected. **Such samples must be prevented from drying out before culturing.** When swabs are used, a transport medium such as Cary Blair, Amies, or Stuart must be used. Sampling strategy in primary poultry has been reviewed (Vidal *et al.*, 2013) and is normally based on boot-swab samples, faecal/caecal droppings or cloacal swabs.

1.1.2. Cattle, sheep and pigs at the farm

Campylobacters are frequent colonisers of the intestine of livestock such as cattle, sheep and pigs; data have been reviewed by Newell *et al.,* (~~in press~~ 2017). Cattle and sheep are found to be colonised mainly with *C. jejuni*, *C. coli*, *C. hyointestinalis*, and *C. fetus*, whereas pigs are predominantly colonised by *C. coli*. In young animals, the numbers are higher than in older animals. In older animals, the organisms can be intermittently detected in faeces, probably due to low numbers or due to

118	intermittent shedding. Fresh samples have to be taken (rectal samples if possible) and **they should**
119	**be prevented from drying out**. When swabs are used, a transport medium (like Cary Blair, Amies,
120	or Stuart) must be used.

1.1.3. At slaughter

122	In poultry, the caecal <u>contents</u> are usually used for the detection of *Campylobacter*. ~~They~~ <u>Caeca</u> can
123	be cut with sterile scissors from the remaining part of the intestines and submitted intact to the
124	laboratory in a suitable container.

| 125 | Samples from cattle, sheep and pigs are collected from the intestines by aseptically opening the gut |
| 126 | wall or by taking guarded rectal swabs. |

127	<u>At all stages from collecting the samples until they are processed in the laboratory, utmost attention</u>
128	<u>should be given to make sure that campylobacters do not die. Follow the instructions for</u>
129	<u>transportation and shipment carefully.</u>

1.2. Transportation and treatment of specimens

1.2.1. Transport

132	Campylobacters are sensitive to environmental conditions, including dehydration, atmospheric
133	oxygen, sunlight and elevated temperature. Transport to the laboratory and subsequent processing
134	should therefore be as rapid as possible preferably the same day<u>.</u> ~~, but~~ <u>It is recommended to process</u>
135	<u>the samples within 72 hours, but if not possible, storage of samples is accepted up to 96 hours (Tast-</u>
136	<u>Lahti *et al.*, 2022)</u> ~~within at least 3 days~~. The samples must be protected from light, extreme
137	temperatures and desiccation.

138	No recommendation on the ideal temperature for transportation can be made, but it is clear that
139	freezing or high temperatures can reduce viability. <u>If possible, samples should be maintained at a</u>
140	<u>temperature of 4°C (±2°C).</u> ~~High temperatures (>20°C), low temperatures (<0°C) and fluctuations in~~
141	~~temperature must be avoided. When the time between sampling and processing is longer than 48~~
142	~~hours, storage at 4°C (±2°C) is advised.~~

1.2.2. Transport media

144	*Swabs:* When samples are collected on boot-swabs or rectal swabs, the use of commercially
145	available transport tubes, containing a medium, such as Cary Blair or Amies, is recommended. This
146	medium may be plain agar or charcoal-based. The function of the medium is not for growth of
147	*Campylobacter* spp., but to protect the swab contents from drying and the toxic effects of oxygen.

148	When only small amounts of faecal/caecal samples can be collected and transport tubes are not
149	available, shipment of the specimen in transport medium is recommended. Several transport media
150	have been described: <u>Amies,</u> Cary-Blair, modified Cary-Blair, modified Stuart medium, Campy<u>-</u>
151	thioglycolate medium, alkaline peptone water and semisolid motility test medium. ~~Good recovery~~
152	~~results have been reported using Cary-Blair (Luechtefeld *et al.*, 1981; Sjogren *et al.*, 1987).~~

1.2.3. Maintenance of samples

154	On arrival at the laboratory, samples should be processed as soon as possible, preferably on the day
155	of arrival<u>. It is recommended to process the samples within 72 hours, but if not possible, storage of</u>
156	<u>samples is accepted up to 96 hours, whereby *C. coli* is more sensitive for long storage times than *C.*</u>
157	<u>*jejuni*</u> ~~but no longer than 3 days after collecting the samples~~ <u>(Tast-Lahti *et al.*, 2022)</u>. To avoid
158	temperature variation, samples should only be refrigerated when they cannot be processed on the
159	same day, otherwise they should be kept at room temperature <u>when processed the same day</u>. When
160	samples are submitted or kept in the laboratory at 4°C, they should be allowed to equilibrate to room
161	temperature before processing to avoid temperature shock.

1.3. Isolation of *Campylobacter*

163	For the isolation of *Campylobacter* from faecal/caecal or intestinal samples, no pre-treatment is needed;
164	samples can be plated on selective medium or the filtration method on non-selective agar can be used. In the
165	case of caecal samples, caeca are aseptically opened by cutting the end with a sterile scissors and squeezing
166	out the material to be processed. Enrichment ~~is recommended~~ <u>can be considered</u> to enhance the culture
167	sensitivity ~~of potentially environmentally stressed organisms or in the case~~ of low levels of organisms in faeces

168	(ISO, 2017), for example from cattle, sheep or pigs. However, enrichment of faecal samples is usually subject
169	to overgrowth by competing bacteria and is not carried out routinely. There is no need to use enrichment
170	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
iii) 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).

1.3.2 Enrichment

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 10–15 drops 100 µl of this suspension are then carefully layered on to a 0.45 or 0.65 µm sterile cellulose acetate 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 37°C or 42°C.

217 ### 1.3.4. Incubation

218   i) Atmosphere

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
221 variety of methods. In some laboratories, (repeated) gas jar evacuations followed by atmosphere
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
226 growth of contaminants and to select for optimal growth of *C. jejuni* and *C. coli*. The fungistatic agents
227 ~~cycloheximide or~~ amphotericin <u>B or cycloheximide</u> are added in order to prevent growth of yeasts
228 and mould at 37°C (Bolton *et al*., 1988). In some laboratories, incubation takes place at 41.5°C to
229 harmonise with *Salmonella* and *Escherichia coli* O157 isolation protocols ~~(ISO, 2006)~~.

230   iii) Time

231 *Campylobacter jejuni* and *C. coli* usually show growth on solid media within 24–48 hours at <u>37–</u>42°C.
232 As the additional number of positive samples obtained by prolonged incubation is very low, 48 hours
233 of incubation is recommended for routine diagnosis (Bolton *et al*., 1988).

234 ## 1.4. Confirmation

235 A pure culture is required for confirmatory tests, but a preliminary confirmation can be obtained by direct microscopic
236 examination of suspect colony material.

237 ### 1.4.1. Identification on solid medium

238 On Skirrow or other blood-containing agars, characteristic *Campylobacter* colonies are slightly pink,
239 round, convex, smooth and shiny, with a regular edge. On charcoal-based media such as mCCDA,
240 the characteristic colonies are greyish, flat and moistened, with a tendency to spread, and may have
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.

246 ### 1.4.3. Detection of oxidase

247 Take material from a suspect colony and place it on to a filter paper moistened with oxidase reagent.
248 The appearance of a violet or deep blue colour within 10 seconds is a positive reaction. If a
249 commercially available oxidase test kit is used, follow the manufacturer's instructions.

250 ### 1.4.4. Aerobic growth at 25°C

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.

253 ### 1.4.5. Latex agglutination tests

254 *Latex agglutination tests* for confirmation of pure cultures of *C. jejuni* and *C. coli* (often also including
255 *C. lari*) are commercially available.

256 ## 1.5. <u>Biochemical</u> identification of *Campylobacter* to the species level

257 Among the *Campylobacter* spp. growing at 42°C, the most frequently encountered species from samples of animal
258 origin are *C. jejuni* and *C. coli*. However, low frequencies of other species, including *Helicobacter* species, have been
259 described. Generally, *C. jejuni* can be differentiated from other *Campylobacter* species on the basis of the hydrolysis
260 of hippurate as this is the only hippurate-positive species isolated from veterinary or food samples. The presence of
261 hippurate-negative *C. jejuni* strains has been reported (Steinhauserova *et al*., 2001). Table 2 gives some basic
262 classical phenotypic characteristics of the most important thermophilic *Campylobacter* species (ISO, ~~2006~~ <u>2017</u>).

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 Biochemical speciation may be supplemented or replaced with MALDI-TOF mass spectrometry. MALDI-TOF can be
266 used to identify *Campylobacter* isolates rapidly and efficiently at the genus and species level (Bessede *et al.*, 2011).

267 ***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.

268 The confirmatory tests for the presence of thermophilic campylobacters and the interpretation (ISO, ~~2006~~ 2017) are
269 given in Table 3. Confirm results of confirmation tests using positive and negative controls.

270 ***Table 3.*** *Confirmatory tests for thermophilic* Campylobacter

Confirmatory test	Result for thermophilic *Campylobacter*
Morphology	Small curved bacilli
Motility	Characteristic (highly motile and cork-screw like)
Oxidase	+
Aerobic growth at 25°C	–

1.5.1. Detection of hippurate hydrolysis

272 Suspend a loopful of growth from a suspect colony in 400 µl of a 1% sodium hippurate solution (care
273 should be taken not to incorporate agar). Incubate aerobically at 37°C for 2 hours, then slowly add
274 200 µl 3.5% ninhydrin solution to the side of the tube to form an overlay. Re-incubate at 37°C for ~~40~~
275 15–30 minutes, and read the reaction. Positive reaction: dark purple/blue. Negative reaction: clear or
276 grey. If commercially available hippurate hydrolysis test disks are used, follow the manufacturer's
277 instructions. The hippurate hydrolysis test is not very robust and the test is often replaced by
278 molecular tests (see below).

1.5.2. Detection of indoxyl acetate hydrolysis

280 Place a suspect colony on an indoxyl acetate disk and add a drop of sterile distilled water. If indoxyl
281 acetate is hydrolysed a colour change to dark blue occurs within 5–10 minutes. No colour change
282 indicates hydrolysis has not taken place. If commercially available indoxyl acetate hydrolysis test
283 disks are used, follow the manufacturer's instructions.

284 ~~Biochemical speciation may be supplemented or replaced with molecular methods or MALDI-TOF mass~~
285 ~~spectrometry. MALDI-TOF can be used to identify *Campylobacter* isolates rapidly and efficiently at the genus and~~
286 ~~species level (Bessede *et al.*, 2011). A variety of DNA probes and polymerase chain reaction (PCR) based~~
287 ~~identification assays has been described for the identification of *Campylobacter* species (On, 1996; Vandamme,~~
288 ~~2000). On & Jordan (2003) evaluated the specificity of 11 PCR-based assays for *C. jejuni* and *C. coli* identification.~~
289 ~~A fast method to differentiate *C. jejuni* and *C. coli* strains is a duplex real-time PCR, targeting gene *mapA* for *C. jejuni*~~
290 ~~identification and gene *CeuE* for *C. coli* identification (Best *et al.*, 2003). Another real-time PCR method commonly~~
291 ~~used to identify and differentiate between *C. jejuni, coli* and *lari* is described by Mayr *et al.* (2010). A gel-based method~~
292 ~~that is commonly used differentiates between *C. jejuni; C. coli, C. lari* and *C. upsaliensis* (Wang *et al.*, 2002).~~
293 ~~*Campylobacter* isolates can also be molecular identified at species level with 16S rRNA sequencing (Gorkiewicz *et*~~
294 ~~*al.*, 2003).~~

1.6. Molecular detection <u>and identification</u> of *Campylobacter*

<u>Multiple</u> PCR-based methods for the detection of *Campylobacter* in animal faecal samples and enriched meat samples have been ~~extensively~~ described in the literature ~~(Bang *et al.*, 2001 Lund *et al.*, 2003; Olsen *et al.*, 1995)~~. <u>Lund *et al.* describe a real-time PCR method to detect *Campylobacter* spp. in chicken faecal samples using magnetic bead DNA isolation followed by a real-time PCR targeting the 16S rRNA gene (Lund *et al.*, 2003; 2004). For food samples, a combined method is described of Bolton broth enrichment and multiplex real-time PCR targeting gene *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).</u> ~~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).~~ Inclusion of positive and negative reference strains and process controls to detect inhibition of the PCR reaction by the sample matrix are required for all molecular *Campylobacter* detection methods.

<u>A variety of DNA probes and PCR-based identification assays has been described for the identification of *Campylobacter* species (Ferrari *et al.*, 2023; Jribi *et al.*, 2017). 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* 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, C. coli* and C. lari is described by Mayr *et al.* (2010). *Campylobacter* isolates can also be identified at species level with 16S rRNA sequencing (Gorkiewicz *et al.*, 2003).</u>

1.7. Antigen-capture-based tests

Enzyme immunoassays are available for the detection of *Campylobacter* in human and animal stool samples <u>(Ricke *et al.*, 2019)</u>. Some are of the lateral flow format. <u>While antigen tests are convenient to use, in an evaluation study 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,</u> the sensitivity and specificity should be critically evaluated through an in-house validation.

2. Serological tests

There are no serological assays in routine use for the detection of colonisation of *C. jejuni* <u>or</u> ~~*/*~~*C. coli* in livestock.

C. REQUIREMENTS FOR VACCINES AND DIAGNOSTIC BIOLOGICALS

There are no vaccines specifically developed for *C. jejuni* or *C. coli* in animals or birds.

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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
460   diagnostic tests and reagents for campylobacteriosis

461   **NB**: FIRST ADOPTED IN 2004. MOST RECENT UPDATES ADOPTED IN 2017.

2   **MEETING OF THE BIOLOGICAL STANDARDS COMMISSION**

3   **Paris, 4–8 September 2023**

4   _____

5   # C H A P T E R   3 . 1 0 . 8 .

6   # T O X O P L A S M O S I S

7   ## SUMMARY

8   *Toxoplasmosis is a zoonotic infection of animals caused by the protozoan parasite* Toxoplasma gondii*. This*
9   *parasite has the potential to infect all warm-blooded animals. Although infection does not result in clinical*
10   *illness in the majority of animal species, in some it causes acute life-threatening disease. In some animals,*
11   *particularly small ruminants,* Toxoplasma *infection may manifest itself as a disease of pregnancy by*
12   *multiplying in the placenta and fetus. In these animals it can result in abortion or the birth of weak offspring.*
13   *Human infections are generally asymptomatic, but they can cause abortion in pregnant women, ocular*
14   *disease, hydrocephalus or intracranial calcifications in congenitally infected children, ocular toxoplasmosis*
15   *in immunocompetent individuals, and serious symptoms and even death in severely immunocompromised*
16   *patients.*

17   Toxoplasma gondii *is an obligate intracellular parasite that has a sexual cycle in Felidae and a two-stage*
18   *asexual cycle in all warm-blooded animals. Globally the* T. gondii *population structure is diverse and the*
19   *various genotypes are associated with the extent of virulence in particular hosts. In the acute phase of*
20   *infection, tachyzoites multiply in host cells to cause varying degrees of tissue destruction. With the onset of*
21   *an immune response, tachyzoites transform into bradyzoites that multiply slowly in cells to produce tissue*
22   *cysts.*

23   ***Detection of the agent:*** *In aborted fetuses and placenta,* T. gondii *is often difficult to find histologically, but*
24   *is more likely to be seen in tissue sections of brain and placenta. Parasitic stages can be identified by*
25   *immunohistochemistry, while nucleic acid-based assays might be used to confirm presence of parasite DNA*
26   *in tissues and may allow genotyping of the parasite in biological specimens.* In-vitro *isolation of* T. gondii
27   *from host samples is expensive, time consuming and rarely used.*

28   *The sexual part of the life-cycle of* T. gondii *takes place exclusively in epithelial cells of the feline intestine*
29   *and can result in the excretion of large numbers of oocysts in the faeces. Oocysts may remain viable in the*
30   *environment for many months. Oocysts of* T. gondii ~~*morphologically*~~ *resemble those from* Hammondia
31   hammondi*, a related but non-virulent parasite that also uses cats as definitive hosts. Nucleic acid-based*
32   *molecular tests are available to distinguish between these related parasites.*

33   ***Serological tests:*** *Among the easy-to-perform serological tests, the indirect fluorescent antibody test* (IFAT)
34   *and the direct agglutination test* (DAT) *allow the titration of sera and the establishment of appropriate cut-*
35   *offs to ensure diagnostic sensitivity and specificity. The IFAT can be used to differentiate IgM and IgG*
36   *antibodies. The DAT is fast and requires no complex laboratory facilities. Enzyme-linked immunosorbent*

37   *assays (ELISA) require more sophisticated laboratory equipment but can process large numbers of samples*
38   *and are easier to standardise.*

39   **Requirements for vaccines:** *A vaccine composed of live* T. gondii *tachyzoites is available commercially for*
40   *use in sheep in certain countries. The vaccine is supplied as a concentrated suspension of tachyzoites with*
41   *an approved diluent and delivery system. The vaccine must be handled strictly according to the*
42   *manufacturer's instructions as it can be hazardous to the user and has a very short shelf life.*

A. INTRODUCTION

44 *Toxoplasma gondii* is a zoonotic, obligate intracellular protozoan parasite with the capacity to infect all warm-blooded
45 animals, including birds. Although clinical toxoplasmosis seldom occurs in the majority of animal species, acute life-
46 threatening disease has been reported in some animals. In small ruminants, in particular, it manifests itself as a disease
47 of pregnancy by multiplying in the placenta and fetus. Acute, potentially fatal, infections have been recorded from a range
48 of wild or zoo animals (Dubey, 2022). Infected humans often show no symptoms, but congenital toxoplasmosis, postnatally
49 acquired ocular toxoplasmosis in immunocompetent individuals, or toxoplasmosis in severely immunocompromised
50 patients represent serious threats (EFSA, 2018).

51 *Toxoplasma gondii* has a two-stage asexual cycle in warm-blooded animals and a sexual cycle in Felidae. A systematic
52 review and meta-analysis have reported a global seroprevalence of 38% in domestic cats and 64% wild felids (Hatam-
53 Nahavandi *et al.*, 2021). The genetic diversity of *T. gondii* is complicated; three archetypal clonal lineages (I, II, and III)
54 prevail in Europe and North America; in South America, Asia, and Africa, much greater genetic diversity is apparent and,
55 furthermore, fewer clonal and non-clonal lineages have been genotyped (Lorenzi *et al.*, 2016). Transport between
56 continents via animal migration, including birds, and human activity such as trade may have contributed to the genetic
57 population structures of *T. gondii* in different geographical regions (Shwab *et al.*, 2018).

58 In the asexual part of the lifecycle, the two developmental stages are the rapidly multiplying tachyzoite and the slowly
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
61 size and shape, but transforms into the bradyzoite stage and multiplies more slowly within tissue cysts to establish a
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
64 infection, and ingestion of bradyzoites in prey is probably the main route of infection to predators, including the felid
65 definitive host. In animals that succumb to acute infection, tachyzoites may be demonstrated in ascitic fluid or in lung
66 impression smears, as well as in tissue sections of the liver and other affected organs.

67 Abortions in sheep and goats due to *T. gondii* are of particular veterinary importance. Toxoplasmosis in small ruminants
68 must be differentiated from diseases caused by other infectious agents, including infections with *Chlamydophila abortus*
69 (see Chapter 3.8.5 *Enzootic abortion of ewes*), *Coxiella burnetii* (see Chapter 3.1.17 *Q fever*), *Brucella melitensis* (see
70 Chapter 3.1.4 *Brucellosis* [*Brucella abortus*, *B. melitensis* and *B. suis*]), *Campylobacter foetus* (see Chapter 3.4.4 *Bovine
71 genital campylobacteriosis*), *Salmonella* spp. (see Chapter 3.10.7 *Salmonellosis*), and the viruses that cause border
72 disease (see Chapter 3.8.1 *Border disease*), bluetongue (see Chapter 3.1.3), Wesselsbron's disease, and Akabane
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* oocysts. Following primary infection of a cat, oocysts may be shed in the faeces for several days, with large
77 numbers contaminating the environment; up to one billion oocysts from domestic cats, and probably similar numbers from
78 wild felids (Shapiro *et al.*, 2019). The oocysts sporulate in the environment over the next 1–5 days (depending on aeration,
79 humidity, and temperature), at which time they become infective. The structure of *T. gondii* oocysts results in extreme
80 resistance to environmental conditions, with the polymeric structure of the walls ~~giving~~ providing strength against
81 mechanical forces and protection against chemical agents (Shapiro *et al.*, 2019). This results in prolonged survival, up to
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
83 to 35°C. Sporulated oocysts are 11 × 13 µm in diameter and each contains eight sporozoites, four in each of two sporocysts
84 (Dubey, 2022). When a susceptible animal ingests sporulated oocysts, the sporozoites are released to penetrate the
85 intestinal lining, become tachyzoites, and establish an infection.

1. Human health risks

87 *Toxoplasma gondii* is a zoonotic parasite and readily infects people. While human infection, as determined by seropositivity,
88 is moderately common globally (local prevalence ranges from under 10% to over 90% [Pappas *et al.*, 2009]), clinical illness
89 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
95	retinochoroiditis, potentially resulting in loss of vision, pneumonitis, and also toxoplasmic encephalitis. The main burden of
96	human disease, based on disability-adjusted life years (DALYs), ranks as having a high contribution to disease burden
97	globally (Torgerson *et al.,* 2015).

98	As with animal infections, people may be infected by ingestion of bradyzoites in raw or lightly cooked meat containing live
99	*T. gondii* tissue cysts or by ingestion of sporulated oocysts. These may be as contaminants of water or of raw or lightly
100	cooked fresh produce; less commonly, people can be infected by ingestion of tachyzoites in non-heat-treated milk. In
101	addition, transmission via blood transfusion or organ transplantation is also possible. Outbreaks of both waterborne and
102	foodborne toxoplasmosis have been described (EFSA, 2018; Shapiro *et al.,* 2019). The largest outbreak to date occurred
103	in Santa Maria, Brazil in 2018, and is considered to be due to contamination of the water supply with oocysts of a virulent
104	strain of *T. gondii*; at least 930 confirmed cases occurred, among which 8% required hospitalisation, with three fetal deaths,
105	10 abortions, and 29 cases of congenital transmission, with 19 infants with ocular lesions (Dubey, 2021).

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.*

B. DIAGNOSTIC TECHNIQUES

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

Method	Purpose					
	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						
PCR (including nested and real-time PCR)	–	–	–	++	+	–
LAMP	–	–	–	++	+	–
Histopathology	–	–	–	+	–	–
Immmuno-histochemistry	–	–	–	+	–	–
Detection of immune response						
IFAT[a]	+	+	+	++	++	+
ELISA[a]	+	+	+	+++[a]	+++	++
DAT/MAT	+	+	+	+	++	+

111	Key: +++ = recommended for this purpose; ++ recommended but has limitations;
112	+ = suitable in very limited circumstances; – = not appropriate for this purpose.
113	PCR = polymerase chain reaction; LAMP = loop mediated isothermal amplification; IFAT = indirect fluorescent antibody test;
114	ELISA = enzyme-linked immunosorbent assay; DAT = direct agglutination test; MAT = modified agglutination test.
115	[a]In IFAT and ELISA, detection of *Toxoplasma*-specific IgG and IgM antibodies may permit some discrimination between acute and
116	chronic cases of infection. In ELISA, assays assessing the avidity of an IgG response to *T. gondii* may provide information regarding
117	how recently the tested animals have experienced a primary *T. gondii* infection.

1. Detection of the agent

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 mineralised centre, and often associated with a mild focal lymphoid meningitis, represent a fetal immune response 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 late gestation caused by advanced lesions in the placentome preventing sufficient oxygen transfer from mother to fetus. Such foci most commonly occur in the cerebral white matter cores, but sometimes also in the cerebellar white 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.

1.2. Immunohistochemistry

Confirmation of the identity of *T. gondii*-like structures in tissue sections from such cases, as well as from instances of acute toxoplasmosis, may be achieved by immunohistochemistry that labels intact *T. gondii* or antigenic debris using polyclonal or monoclonal *T. gondii* specific antibodies (Dubey, 2022). The antigen-antibody reaction can be visualized by avidin-biotin-complex (ABC) or indirect immune-peroxidase and the peroxidase–antiperoxidase (PAP) technique. The method is both convenient and sensitive and is used with fixed tissues (including archived tissues) that may also exhibit a degree of decomposition, where isolation would not be appropriate or possible. However, cross-reactions with related parasites like *Neospora caninum* are possible.

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.

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
169	bradyzoites in tissue cysts (latent infection) and in different biological samples, including animal and human
170	tissues (e.g. heart and skeletal muscle, placenta, brain) and body fluids (e.g. blood, urine, aqueous humour,
171	cerebrospinal fluid, amniotic fluid, milk). In addition, DNA from oocysts in stool (only felids), food and
172	environmental samples (fresh produce, water, soil) can be detected (Slana *et al.*, 2021). No standard method
173	for extracting *T. gondii* DNA exists, but suitable DNA extraction protocols, based on both in-house protocols and
174	commercial kits have been developed (Dzib Paredes *et al.*, 2016). Sample preparation and DNA extraction
175	procedures are likely to have a considerable impact on the sensitivity of the test. Sensitivity is generally higher
176	in DNA-poor matrices than DNA-rich ones (e.g. tap water vs meat samples). Moreover, inhibitors of DNA
177	amplification differ and are related to sample type. Inhibition of DNA amplification can be avoided by DNA
178	extraction optimised for sample type or by using appropriate additives (e.g. bovine serum albumin) during DNA
179	amplification. A specific concentration of the parasite stage (e.g. oocysts) or its DNA (e.g. by magnetic capture)
180	from the matrix might be required prior to DNA extraction as reported for oocysts from water, fresh produce, or
181	faeces (Slana *et al.*, 2021). A validated protocol for DNA extraction from meat and meat products using a
182	commercial kit is available at the website of the European Union Reference Laboratory for Parasites (EURLP)[48].
183	An example of in-house method for DNA extraction from pig tissue (Jauregui *et al.*, 2001) is reported below.

1.4.1. DNA extraction from animal tissue

Procedure

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 Na_2HPO_4, 1.7 mM NaH_2PO_4).

ii) Digest sample with an equal volume of warm (37°C) pepsin-HCl (1.4 mg of pepsin and 10 mg of 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).

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 oocyst~~s~~ <u>walls</u> 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

i) Wash oocysts four times in 15 ml PBS (300 mM NaCl, 2.7 mM KCl, 10 mM Na_2HPO_4, 1.7 mM NaH_2PO_4) in a 15 ml centrifugation tube, with centrifugation between washes (1100 g for 7 minutes without braking).

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48 https://www.iss.it/documents/5430402/5722370/MI_12_rev._1.pdf/a82a4078-f511-affe-8f90-cabc397bc8ce?t=1620381672663

| 218 | | ii) | Incubate pellet (up to 0.5 ml) in 2 ml 5.75 % sodium hypochlorite (sodium hypochlorite, aqueous |

| 218 | ii) | Incubate pellet (up to 0.5 ml) in 2 ml 5.75 % sodium hypochlorite (sodium hypochlorite, aqueous |

ii) Incubate pellet (up to 0.5 ml) in 2 ml 5.75 % sodium hypochlorite (sodium hypochlorite, aqueous solution, ≥ 4% as active chlorine) for 30 minutes at 37°C.

iii) Add double-distilled H$_2$O up to 15 ml.

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 **g** for 7 minutes without brake).

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).

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.

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.

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.

ix) Mix with 500 µl phenol/chloroform/isoamyl alcohol (25/24/1) by inverting 50 times. Centrifuge for 7 minutes at 13,000 **g**.

x) Transfer supernatant to fresh tube and mix with 500 µl phenol/chloroform/isoamyl alcohol (25/24/1) by inverting 50 times. Centrifuge for 7 minutes at 13,000 **g**.

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).

xii) Centrifuge for 15 minutes at 13,000 **g**. Decant the supernatant.

xiii) Wash the pellet using 70% (v/v) ethanol and centrifuge for 15 minutes at 13,000 **g**.

xiv) Discard the ethanol solution and air dry the pellet.

xv) Solve DNA in double-distilled water for at least 12 hours at 4°C.

xvi) Use 2.5–10 µl aliquots for PCR (see Section B.1.2 above).

The same PCR-based and LAMP assays detailed in Section B.1.2 have been also used for oocyst detection, with B1 gene and the 529RE being targets of choice (Slana *et al.*, 2021).

Another important issue is the possibility of combining detection with information on oocyst viability. Bioassays, currently the only definitive way to detect viable oocysts, are expensive and relatively few laboratories have the necessary facilities. Reverse transcription (RT) real-time PCR (real-time RT-PCR) or propidium monoazide-based real-time PCR have shown some promise for assessing oocyst viability in complex sample matrices (Kim *et al.*, 2021; Rousseau *et al.*, 2018).

1.4.3. Nucleic acid detection methods

Although single copy genes (e.g. SAG1, SAG2, SAG3, GRA6, and GRA7) have been used, multi-copy genes or genetic elements (e.g. 18S rDNA, B1, ITS1, 529RE) are preferred as they provide a higher sensitivity (Dzib Paredes *et al.*, 2016; Slana *et al.*, 2021). For instance, 35 copies of the B1 gene and 200–300 copies of the 529 bp repetitive element (529RE) are present in the *T. gondii* genome, and 10 to 100-fold higher sensitivity is generally observed in amplification targeting the 529RE compared with B1, although this also reflects the type of assay and sample being analysed (Belaz *et al.*, 2015). In addition, some strains may have partially lost, or have a mutated, 529RE, and this could compromise diagnostic sensitivity (Wahab *et al.*, 2010).

Conventional PCR targeting B1 was the first to be used in clinical diagnostics in people (Burg *et al.*, 1989). To obtain details on *T. gondii* genotype (e.g. for outbreak investigation, infection source tracing) the methods of choice are multi-locus PCR combined with restriction fragment length polymorphism (PCR-RFLP) or sequencing and multi-locus microsatellite typing (Ajzenberg *et al.*, 2010; Su *et al.*, 2006).

To increase sensitivity, several nested PCR assays have been implemented (Dzib Paredes *et al.*, 2016). The reaction consists of two successive rounds of amplification. The product of the first amplification serves as template for the second amplification, using one or two internal primers. The risk of cross- and carry-over contamination and false positives is increased with nested PCR, and precautions should be taken to mitigate the risk (Dzib Paredes *et al.*, 2016).

269	There are several real-time PCR protocols and real-time PCR in combination with a hydrolysis probe
270	is the most frequently applied (Slana *et al.*, 2021). This has largely improved both sensitivity and
271	specificity of detection of *T. gondii* DNA, with the advantage of avoiding post-amplification
272	manipulations and thus limiting the risk of carry-over contamination. Although sensitivity can be
273	satisfactory with both conventional and real-time PCR using pure genomic *T. gondii* DNA, assay
274	specificity might be affected when testing field samples. Conventional PCR can result in non-specific
275	amplification, whereas this is not detected by real-time PCR due to the probe detection, despite the
276	amplified target being the same. Furthermore, real-time PCR can be multiplexed and simultaneous
277	amplification of an internal amplification control (a heterologous DNA fragment) can be used to
278	monitor for the presence of inhibitors. In addition, amplifying two *T. gondii*-specific targets at once
279	may increase sensitivity. Another advantage of real-time PCR is the possibility of quantification of *T.*
280	*gondii* DNA.

281	A selective enrichment of target DNA combined with real-time PCR (i.e., magnetic capture PCR) has
282	been reported to increase *T. gondii* detectability in animal samples (Gisbert Algaba *et al.*, 2017). The
283	principle relies on separating and concentrating *T. gondii* DNA from sample DNA by specific DNA
284	probes, complementary to the targeted parasite 529RE genomic region, which are conjugated to
285	magnetic beads and followed by real-time PCR (Gisbert Algaba *et al.*, 2017). However, this method
286	is expensive, time consuming, and requires further expertise, so might be not suitable for routine
287	analysis or large surveys.

288	As an alternative to PCR, LAMP has been considered for *T. gondii* DNA detection, and diagnostic
289	purposes, in environmental, veterinary, and human samples. LAMP takes advantage of a DNA
290	polymerase (originally Bst) having both high strand displacement and replication activities. Nucleic
291	acid amplification is performed under isothermal conditions (60–65°C), without the need of a DNA
292	denaturation step. Both, B1 and 529RE have been widely used as targets in different LAMP assays,
293	and LAMP is reported as comparable to real-time PCR for the detection of *T. gondii* in blood and
294	animal tissues (Robert *et al.*, 2021). Although LAMP provides an opportunity for development of point-
295	of-care testing or implementation of molecular tests in settings with limited facilities, there are several
296	drawbacks including design of appropriate primers and the high risk of carry-over contamination.
297	Commercial assays for both real-time PCR and LAMP are available for clinical diagnosis of
298	toxoplasmosis.

299	Overall, the reported sensitivity of published molecular methods can be as low as one (or even less)
300	genome equivalent per reaction. However, this largely depends on sample type, DNA extraction,
301	copies of the targeted gene or sequence, amplification and detection reagents, procedures and
302	platforms. The lack of accepted standard methods prevents robust comparison of sensitivity and
303	specificity of the currently applied molecular tests.

2. Serological tests

305 There are several serological tests available for the detection of *T. gondii* antibodies (Table 1). All serological tests have
306 limitations in diagnostic sensitivity and specificity and need proper validation to ensure confidence in results (refer to
307 Chapter 1.1.6. *Principles and methods of validation of diagnostic assays for infectious disease* and Chapter 2.2.1.
308 *Development and optimisation of antibody detection assays*).

309 The Sabin-Feldman dye test (DT) is a reference serological test for *T. gondii* antibody in humans (Dubey, 2022). Although
310 the DT appears both specific and sensitive in humans, it is not extensively validated in other species. In addition, it is
311 potentially hazardous as live parasite is used, is expensive, and requires a high degree of technical expertise.

2.1. Preparation of antisera and antigens

| 313 | Antisera to *T. gondii* and conjugated antisera for use in IFAT and ELISA, to allow screening of a variety of animal |
| 314 | species, may be obtained commercially. International standards for animal sera are not available. |

| 315 | Below are protocols for the preparation of tachyzoite antigen for use in the IFAT and ELISA. Tachyzoites may be |
| 316 | grown in tissue culture and retained as whole parasites for the IFAT, or prepared as soluble antigen for the ELISA. |

2.2. Preparation of frozen stabilates of *T. gondii* tachyzoites

2.2.1. Test procedure

i) Produce tachyzoites in tissue cell culture as described. Suitable *T. gondii* 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]).

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 *T. gondii.* Final concentration of the tachyzoite suspension should be approximately 1.5×10^8 tachyzoites/ml.

iii) Combine dimethyl sulphoxide (DMSO), normal horse serum (free from antibody to *T. gondii*) 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.

iv) Allow the preparation to stand on the bench for 1 hour (4-10°C; optimally on ice).

v) Dispense into 1-ml aliquots using screw-topped tubes appropriate for liquid nitrogen storage.

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.

vii) The next day transfer to liquid nitrogen, keeping well insulated while transferring.

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).

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 *T. gondii* and add suspension to cell culture.

2.3. Production of *Toxoplasma* tachyzoites in cell culture

2.3.1. Test procedure

i) *Toxoplasma gondii* 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).

ii) Cells and parasite are grown in IMDM supplemented with 50 IU/ml penicillin, 50 µg/ml streptomycin, and 2% fetal bovine serum; there are other cell culture media suitable as well.

iii) Tachyzoites are harvested from tissue culture flasks by scraping the cell monolayer using a sterile cell scraper.

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.

2.4. Preparation of whole tachyzoites for IFAT and agglutination

2.4.1. Test procedure

i) Produce 4×10^7/ml suspension of *T. gondii* tachyzoites in PBS.

ii) Add formaldehyde (40%) to give a final concentration of 0.2% (v/v).

iii) Incubate at 4°C overnight and divide into aliquots in suitable tubes and store frozen until required (–20°C).

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49 American Type Culture Collection, P.O. Box 1549, Manassas, Virginia 20108, United States of America.

2.5. Production of soluble antigen for ELISA

2.5.1. Test procedure

i)   Produce a suspension of *T. gondii* tachyzoites in PBS.

ii)   Centrifuge at 2000 **g** for 15 minutes, retain the pellet and resuspend it in nine times its volume of distilled water.

iii)   Lyse the tachyzoites by freezing and thawing three times.

iv)   Sonicate the antigen preparation for 20 seconds at 4°C at an amplitude of 20 microns.

v)   Remove any cellular debris by centrifugation at 10,000 **g** for 30 minutes at 4°C.

vi)   Retain the supernatant and store at –20°C until required; protein estimation should be between 2 and 4 µg/ml.

2.6. Indirect fluorescent antibody test

The indirect fluorescent antibody test (IFAT) (Dubey, 2022) is a simple and widely used method. Whole, killed *Toxoplasma* tachyzoites are incubated with diluted test serum, the appropriate fluorescent labelled secondary antibody is added, and the result is then viewed with a fluorescence microscope. Fluorescent-labelled species-specific secondary antibodies are available commercially, the method is relatively inexpensive, and kits are commercially available. However, the results are read by eye, so subjective variation may occur. It may be difficult to find some species-specific conjugates and there is a risk of possible cross-reactivity with rheumatoid factor and anti-nuclear antibodies. The following is a protocol for carrying out an IFAT for anti-*Toxoplasma* IgG antibodies in sheep serum. It only requires minor modifications for testing different species or for measuring IgM antibody.

2.6.1. Test procedure

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.

ii)   Apply 5 µl of a whole tachyzoite preparation (Section B.2.4.1 above) on to each well and allow to air dry.

iii)   Fix in methanol for 10 minutes.

iv)   Wash twice (10 minutes for each wash) in 0.3 M PBS, pH 7.4.

v)   Prepare serial dilutions of the test sera in PBS (e.g. 1/16, 1/32, etc. up to 1/1024).

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.

vii)   Wash twice (10 minutes each time) in PBS.

viii)   Add 5 µl 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 µm sterile filter), to each well and incubate for 30 minutes at room temperature.

ix)   Wash three times for 10 minutes each time in PBS.

x)   Mount the slides under cover-slips with buffered glycerol (nine parts PBS, one part glycerol).

xi)   Examine using a fluorescence microscope, fitted with ×20 and ×40 objective lenses.

With a negative test serum result, the whole parasites will appear red due to the autofluorescence of the Evans blue dye. They may also present with a green fluorescent cap at the parasite pole (nonspecific polar fluorescence). With a positive test serum, the parasites will fluoresce red and at least 80% of them within a given well will be surrounded by an unbroken band of green fluorescence. In an adult sheep/goat a positive titre could be defined as ≥1/64 and a negative titre as ≤1/32. For lamb/kid and fetal sera, respective titres could be defined as ≥1/32 and ≤1/16. These cut off values should be validated locally as results may vary between laboratories, depending on, e.g., the fluorescence microscope and the operator. Optimally, each slide should include positive control and negative controls.

2.7. Modified agglutination test

The modified agglutination test (MAT) (Dubey, 2022) is both sensitive and specific. Formalinised *Toxoplasma* tachyzoites are added to U-shaped well microtiter plates and dilutions of test sera are then applied. Positive samples will produce agglutination that can be graded, whereas negative samples will produce a 'button' of precipitated 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 false positives due to non-specific IgM. The MAT has been used extensively for detection of *T. gondii* antibodies in 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.

2.7.1. Serum-diluting buffer

i)   Dissolve 42.5 g NaCl, 1.54 g NaH$_2$PO$_4$, and 5.4 g Na$_2$HPO$_4$ 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.

2.7.3. Serum dilutions

i)   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

i)   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.

2.8. Enzyme-linked immunosorbent assay

One of the first *T. gondii* enzyme-linked immunosorbent assay (ELISA) (Voller *et al*., 1976) used a soluble antigen preparation made from *T. gondii* RH strain tachyzoites (as described below) and layered into wells in an ELISA microtiter plate. Test sera are added, followed by a species-specific secondary antibody conjugated with a reactive 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.

458 To improve the specificity of the conventional ELISA, native purified *T. gondii*-specific antigens have been used
459 (Basso *et al.*, 2013). In addition, severam recombinant antigens have been established, and these seem suitable for
460 replacing native antigens for serological diagnostic tests. For many of these recombinant antigen ELISAs, thorough
461 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.

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
468 (Europe and New Zealand) to reduce the effects of *T. gondii* infection (e.g. early embryonic death, abortion). It consists of
469 $\geq 10^5$ 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
471 weeks prior to mating and only for use in healthy, non-pregnant female sheep. Despite the acknowledged importance of
472 human toxoplasmosis, human vaccines are currently unavailable, and the vaccine for sheep has disadvantages, such as
473 a short shelf-life (10 days), strict storage conditions, and, as a live vaccine, potential risk to operators. Information on the
474 production details of this vaccine and QC requirements are not available.

475 Although the importance of a killed or non-live vaccine is acknowledged – for vulnerable humans (e.g. women before they
476 are pregnant), for reducing abortions in sheep, for reducing tissue cysts in meat animals (pigs, cattle, chickens, etc.), and
477 for limiting oocyst shedding from kittens – to date this remains elusive (Innes *et al.,* 2019). However, with recent scientific
478 advances, including availability of genetic, transcriptomic, and metabolomic data, the potential for developing knockout
479 variants, and other new technologies suggest new possibilities for development of such a vaccine (Mevelec *et al.,* 2020;
480 Zhang *et al.*, 2022).

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557   **NB:** At the time of publication (2024) there were no WOAH Reference Laboratories
558   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

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Annex 20. List of Main Focus Areas and Specialties for WOAH Collaborating Centres

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The role of WOAH Collaborating Centres is anchored to the WOAH's founding mandate and to the Seventh Strategic Plan (2021–2025)[50].

1.   Animal health management

WOAH has the responsibility to collect, analyse and disseminate relevant scientific information, especially on disease control methods, and to provide expertise in the control of animal diseases including zoonotic diseases, as well as health threats at the animal–human–ecosystems interface, ~~while taking into account~~ under the "One Health" approach ~~concept whenever possible~~. This topic covers issues primarily, but not exclusively, related to Sections 2 and 4 of the _Terrestrial_ and _Aquatic Codes_ and to Part~~s~~ 3 of the _Terrestrial Manual_ and Part 2 of ~~both~~ the _Terrestrial_ ~~and~~ _Aquatic Manuals,_ respectively. The topic is expected to assist WOAH and its Members to fulfil the core missions of the 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 disease~~s~~ control
- Biothreat reduction
- Impact of climate change on animal health

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~~ 4.9 of the _Aquatic Code_.

- Animal welfare
- Animal production food safety
- Sustainable animal production (including integrated health management)
- Safety of animal feed
- Climate change and impacts

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
- Laboratory information systems technology
- Validation ~~procedures for diagnostic tests~~ of laboratory methods
- Development and application of innovative technologies

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50   https://www.woah.org/en/document/seventh-strategic-plan/

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 para-professionals 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 education (under~~-~~ and post-graduate) ~~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

5. Veterinary products

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
- Managing antimicrobial ~~agents~~ resistance
- Alternatives to antimicrobials
- New technologies

6. Wildlife health and biodiversity ~~Environment and climate change~~

~~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 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 expected to address animal health issues, including aquatic animals, connected to wildlife, biodiversity, climate change, and emerging risks.~~ Wildlife plays a vital role in maintaining healthy and functioning ecosystems, thus contributing to the preservation of biodiversity. Wildlife is an asset, supporting livelihoods through the provision of income, whether it be through tourism or as a source of food. Importantly, wildlife has a positive effect on human well-being, contributing to education, physical and mental health, social values, culture and spirituality. This topic is expected to address animal health issues, including terrestrial and aquatic animals, connected to wildlife, biodiversity, and emerging risks.

- Threats to ~~livestock or~~ wildlife health and welfare and biodiversity
- Impact of climate change ~~and biodiversity~~ on wildlife
- ~~Disease related (including vector-borne)~~
- Wildlife disease epidemiology, modelling, and surveillance
- 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