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# Harmonisation of the diagnostic performances of ELISA tests for *C. burnetii* antibodies in ruminants: optimal positivity thresholds and performance reassessment

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Context

- Several ELISAs are available for the serological diagnosis of *C. burnetii* infection in ruminants.
- None can be considered as a gold standard or as a reference test.
- **Differences between their diagnostic performances** (Lurier *et al.* 2021)  $\Rightarrow$  decrease the agreement between the test results from different veterinary laboratories, and  $\Rightarrow$  limits the comparability of surveillance data.
- Harmonisation of the tests is possible by changing their positivity thresholds.
  - Proposed ROC curves estimation methods (ordered multinomial model Wang et al. 2007 and mixture model – Choi et al. 2006) are not applicable due to the conditional dependency of the tests and to the non-Gaussian distribution of their quantitative measurement.

# **Objectives**

- Identify the positivity thresholds that maximise the agreement between the tests.
- Estimate the impact of changing the thresholds.

#### Method

- Data: results of three ELISAs from 1258 cattle, 1474 goats and 1432 sheep.
- Thresholds maximising Hubert's kappa were identified using a differential evolution algorithm.
- The uncertainty around these optimal thresholds was estimated by bootstrapping.
- Sensitivities and specificities of the tests at the optimal thresholds were estimated by Bayesian inference using the hierarchical zero-inflated beta-binomial latent class **model** previously described by Lurier *et al*. (2021).

# Results

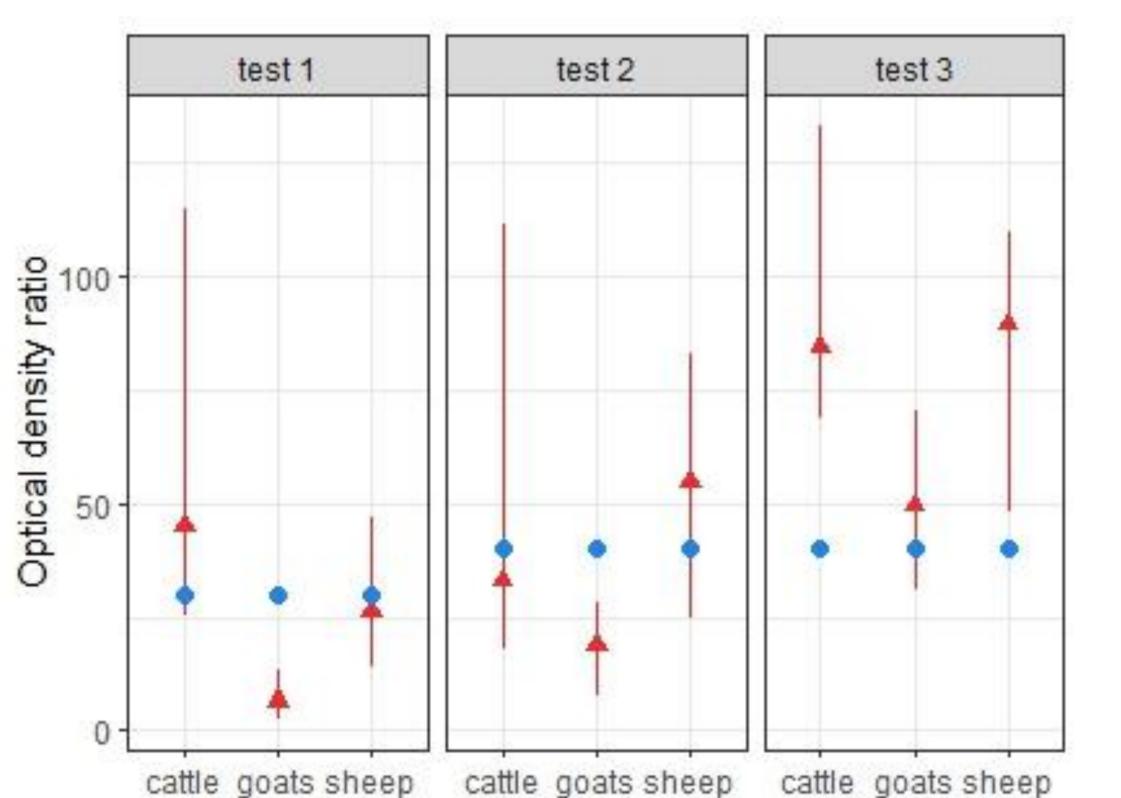
**Optimal positivity thresholds** 

# Impact of the threshold change

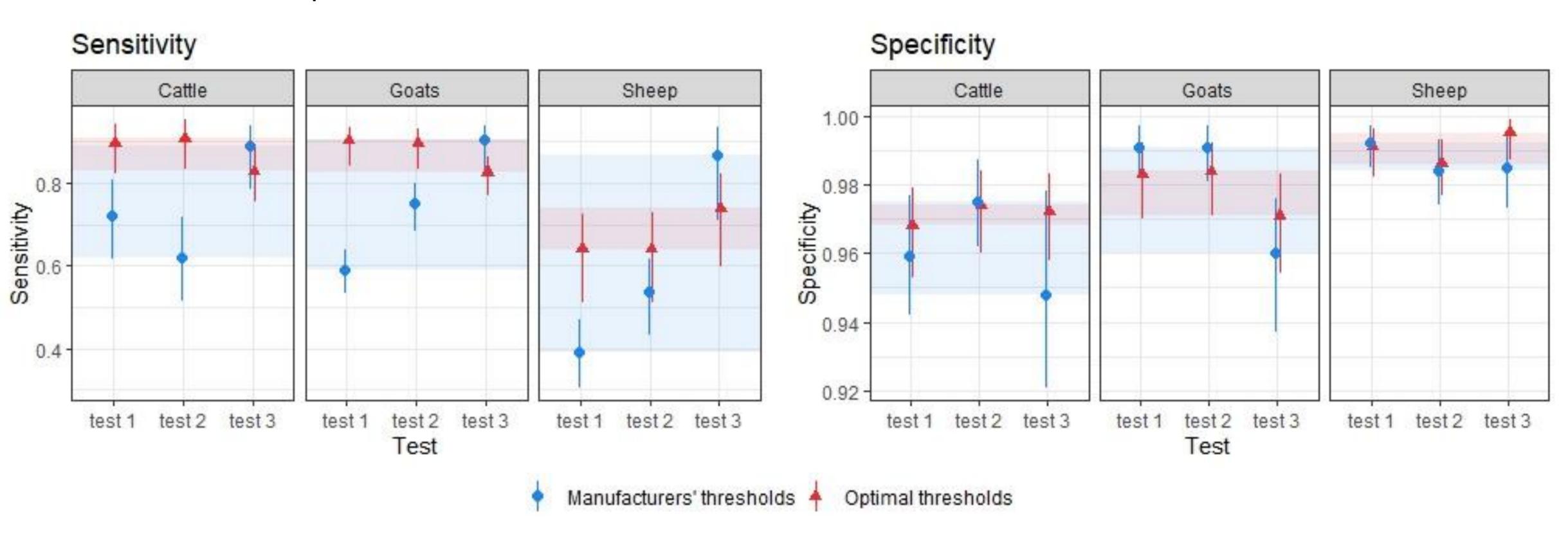
- **Diagnostic performances**

**Optimal thresholds** are different (higher or lower) from the manufacturers' thresholds.

The uncertainty around the optimal thresholds is high, but should be interpreted with caution as the coverage of these bootstrap intervals is not guaranteed in this case.



At the optimal thresholds, the sensitivities of tests 1 and 2 increased without major changes in their specificities. The **sensitivity of test 3 decreased** while its **specificity increased**. The sensitivities and specificities are **closer** from one test to another.



Sensitivity and specificity of the tests at manufacturers' and optimal thresholds and their 95% credibility intervals

**Observed disagreement** 

The proportion of discordant cases is **divided by approximately 2** compared to the manufacturers' thresholds (e.g. from 20 to 12% for goats).

#### Species

Manufacturers' thresholds 🔺 Optimal thresholds

Manufacturers' thresholds and thresholds that maximise Hubert's kappa and their 95% bootstrap intervals

#### Apparent prevalence

The apparent prevalences are more similar from one test to another at the optimal thresholds (e.g. in sheep they range from 7.7 to 17% at the manufacturers' thresholds and from 8.7 to 10.3% at the optimal thresholds).

Findings of the study Improved harmonisation of the test results and of their diagnostic performances  $\rightarrow$  Improved comparability of surveillance data

### **Discussion and perspective**

The increase in sensitivity of tests 1 and 2 estimated in this study is not consistent with the increase in their thresholds compared to the manufacturers' ones.

The development of methods to evaluate ROC curves in the absence of a gold standard, especially considering the conditional dependence between tests, remains an important area for future research.

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Wang, Chong, Bruce W. Turnbull, Yrjö T. Gröhn, and Søren S. Nielsen. 2007. 'Nonparametric Estimation of ROC Curves Based on Bayesian Models When the True Disease State Is Unknown'. Journal of Agricultural, Biological, and Environmental Statistics 12 (1): 128–46. https://doi.org/10.1198/108571107X178095.