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Interlaboratory diagnostic test evaluation using Bayesian latent class models

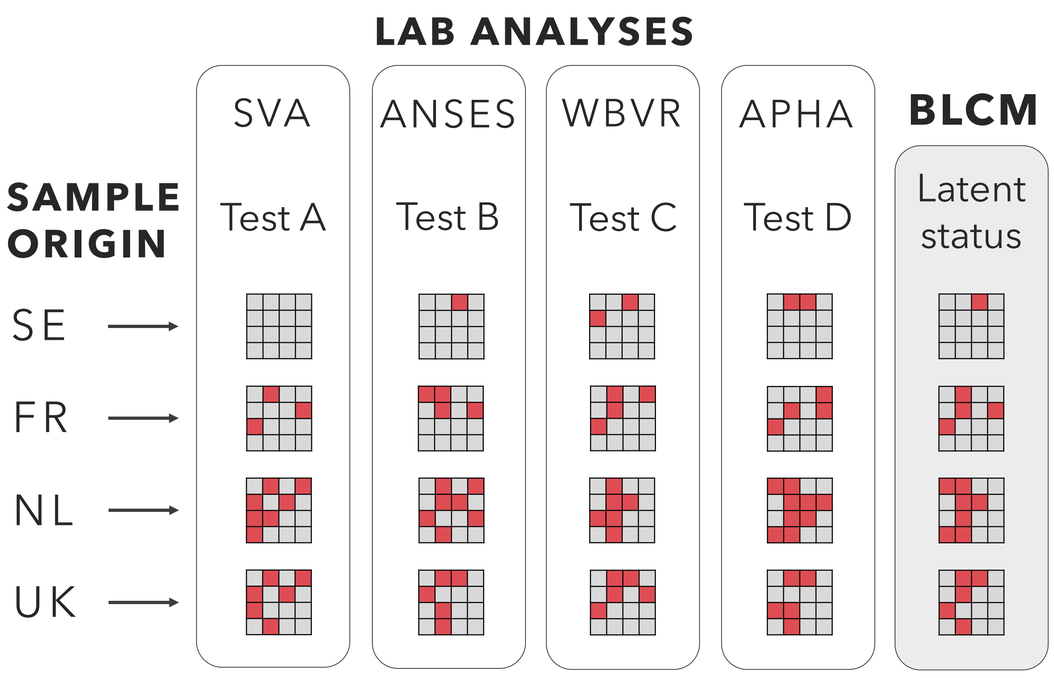
Knowing diagnostic test accuracy is essential when using test results to infer the presence, prevalence, or absence of a disease in a population

AIM

Proof of concept for cost-effective diagnostic test evaluation

CHALLENGES → SOLUTIONS

- Lack of gold standard test → Bayesian latent class models
- Lack of positive samples → Samples from multiple countries
- Budget constraints → Inter-laboratory evaluation



The Bayesian latent class model (BLCM) combines the test results in each population and infer the most likely *true* value for each sample, i.e. the *latent status*.

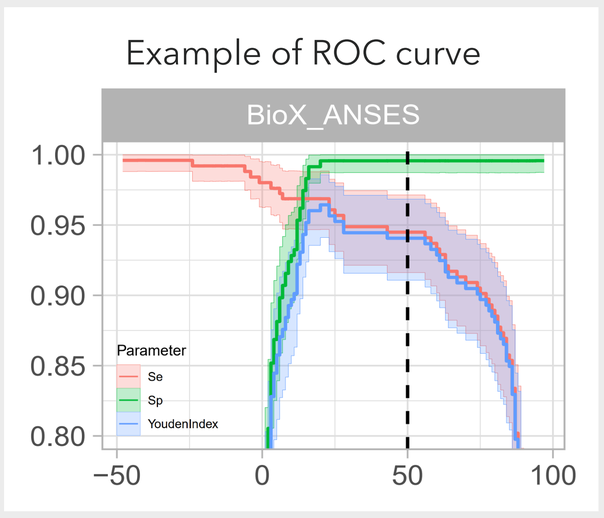
CASE STUDIES

Serological detection of **BVD** & **IBR**

- **2** diseases
- **4** countries
- **4** laboratories
- **6** epidemiologists
- **6** laboratory experts
- **7** serological tests for IBR
- **8** serological tests for BVD

STUDY DESIGN

- Sample sharing between labs
- Each lab applied their own routine diagnostic test(s)
- Joint data analysis using BLCM



Example of population-specific estimates (Svanovir_SVA)

Population	Median specificity	Credible interval
SE	0.995	[0.977 – 1.000]
NL	0.345	[0.198 – 0.509]
UK	0.916	[0.780 – 0.997]
FR	0.988	[0.948 – 1.000]
Combined	0.903	[0.863 – 0.938]

OUTCOMES

- Diagnostic sensitivity and specificity of ELISA tests for detection of Bovine Viral Diarrhoea (BVD) and Infectious Bovine Rhinotracheitis (IBR)
- Population-specific accuracy and ROC curves for cut-off optimization
- R code for inter-laboratory diagnostic test evaluation using BLCM:
  - → algorithm for **optimal sample allocation** among populations
  - → methods for **assessing the empirical fit** of Hui-Walter models
  - → **post hoc accuracy estimates** for tests that were excluded from the model

CONCLUSIONS

- Cost-effective approach to diagnostic test evaluation
  - → especially valuable for routine diagnostics
- Useful insights to the labs and tools for test optimization
- Applying several tests entails:
  - 👎 correlations that increase model complexity
  - 👍 possibility to remove tests and get post hoc estimates
- Requires deep understanding of tests, populations and underlying latent status