

INTERNATIONAL
EpiLab



Probability of freedom from paratuberculosis in Danish dairy herds

Evan Sergeant



Objective

- To develop a method to estimate probability of freedom from paratuberculosis in Danish dairy herds based on individual milk-ELISA testing;



Terminology

- Sensitivity – probability that an infected individual will test positive
- Specificity - probability that an uninfected individual will test negative
- Design prevalence – pre-determined prevalence level that you wish to be confident of detecting with your testing strategy
- Probability of freedom – probability that the true prevalence is less than the design prevalence
- **Note:** some high PFree herds may still be infected but at less than the design prevalence



Methods

- Simulation model implemented in R and on web
- Age-specific sensitivity and specificity estimates used
- A distribution of observed values used for design prevalence (5% - 85%, mean = 34%)
- Analysis based on actual or simulated test results for 19 herds
- Calculate average Se and Sp for each herd based on age structure
- Calculate probability of observed result for uninfected herds and for herds infected at design prevalence
- Calculate probability of freedom (P_{Free}) for observed result
- Simulate for 1,000 iterations



For each Animal

Age of animal

Age-specific
sensitivity

Age-specific
specificity

Design
Prevalence

$P(\text{Test positive})$

For each Herd

Mean values for
 Se , Sp and $P(T+)$

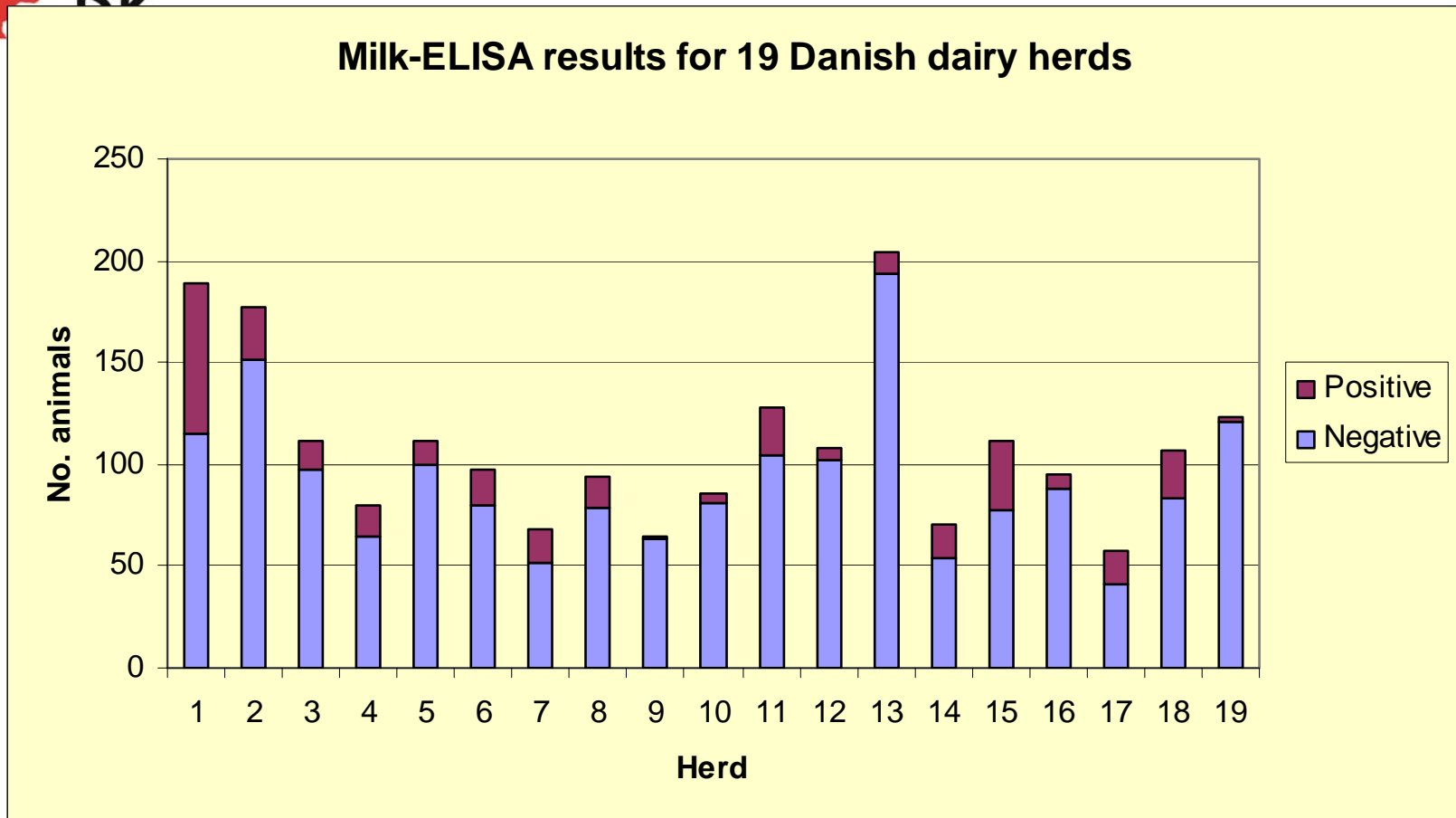
True Prevalence

Probabilities of
observed number of
reactors for infected
and uninfected herds

$P(\text{Free})$



Results

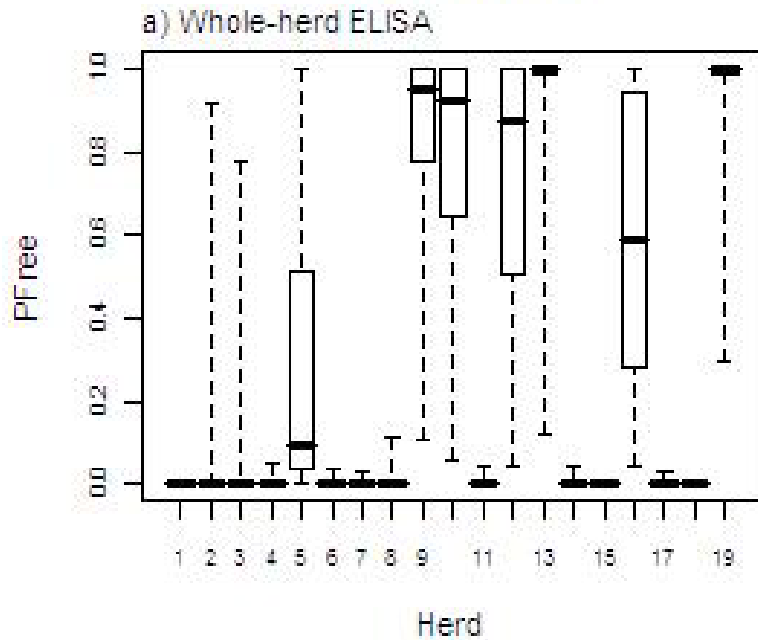


- 2 herds < 5%, 4 herds 5 – 7%
- 7 herds 10 – 20%
- 6 herds > 20%

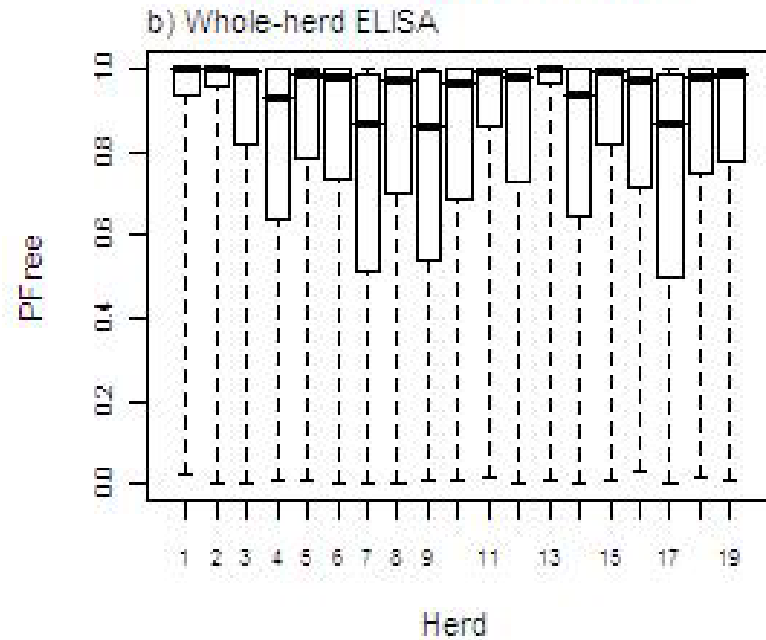


Results

Actual test results

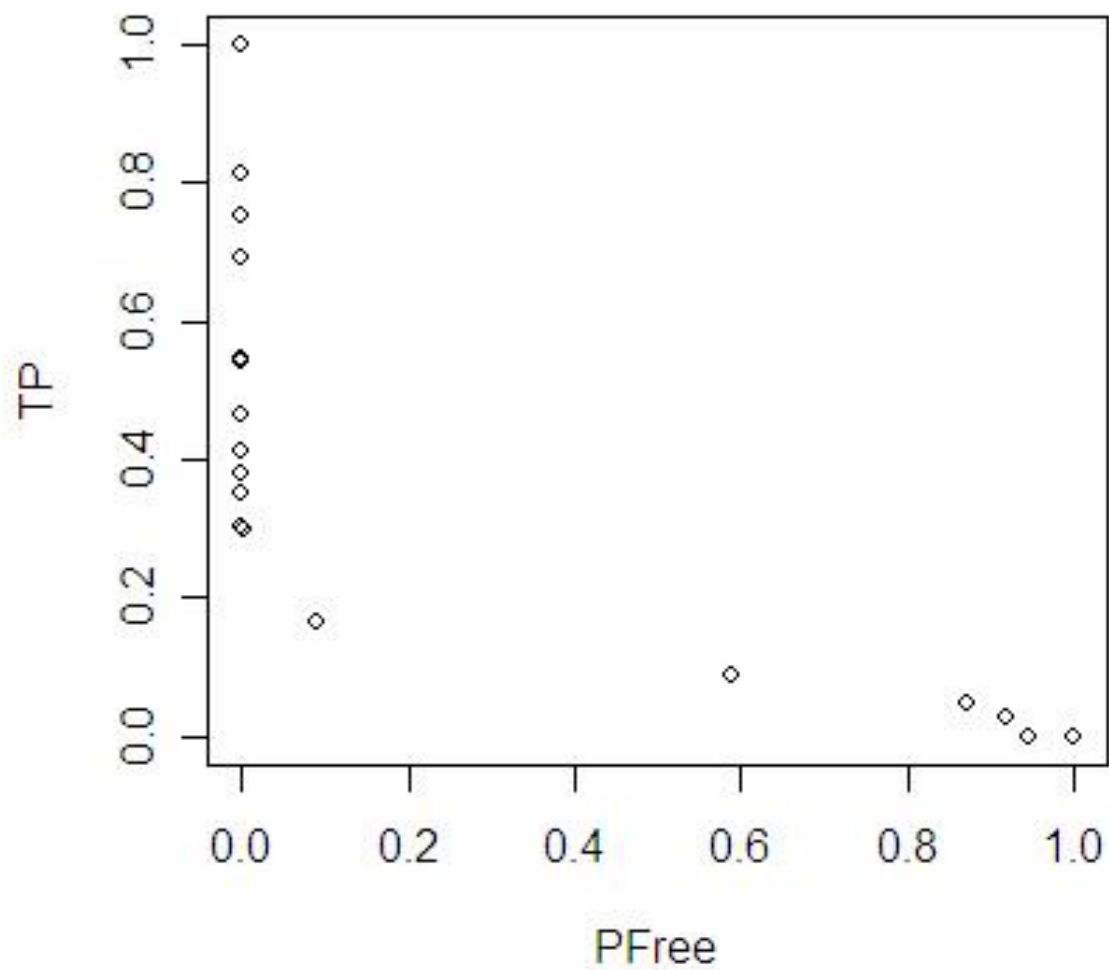


Simulated Uninfected herds





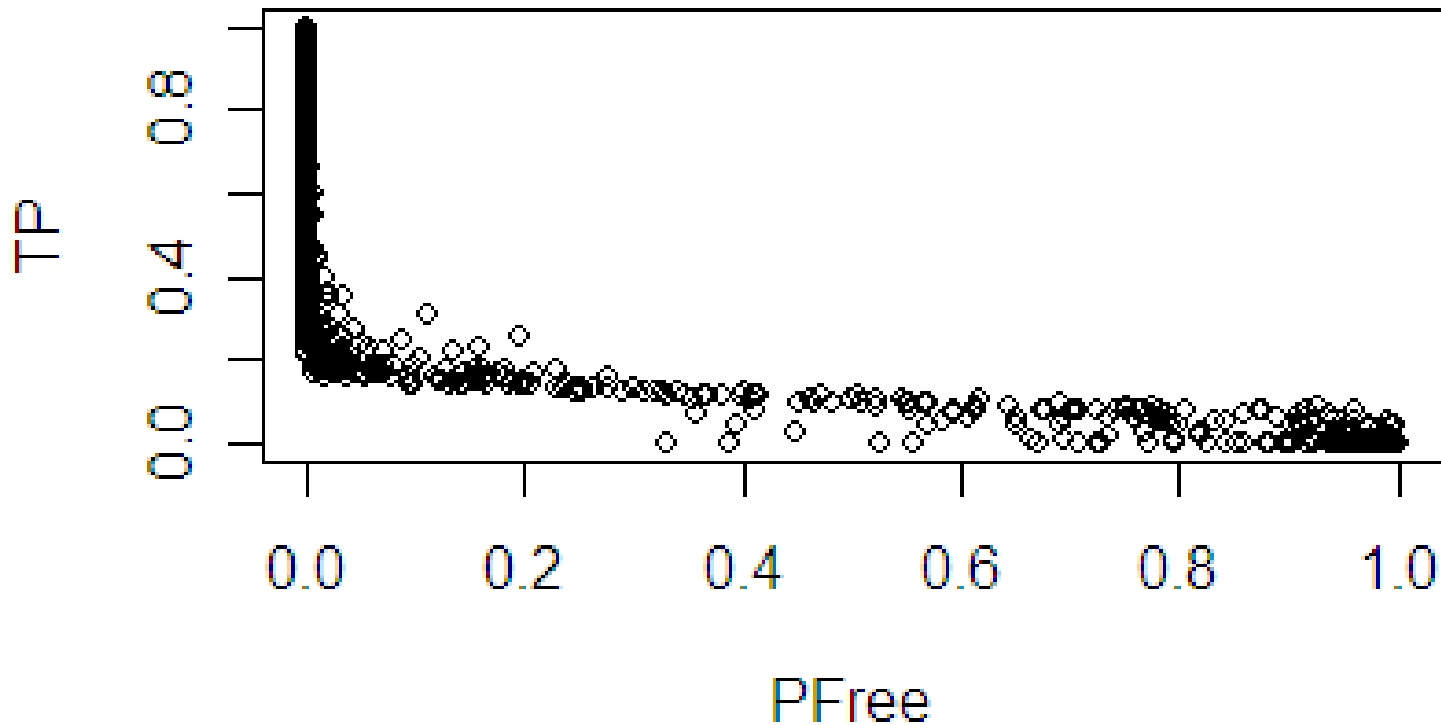
TP vs PFree





844 herds (Operation ptb)

TP vs PFree





Discussion

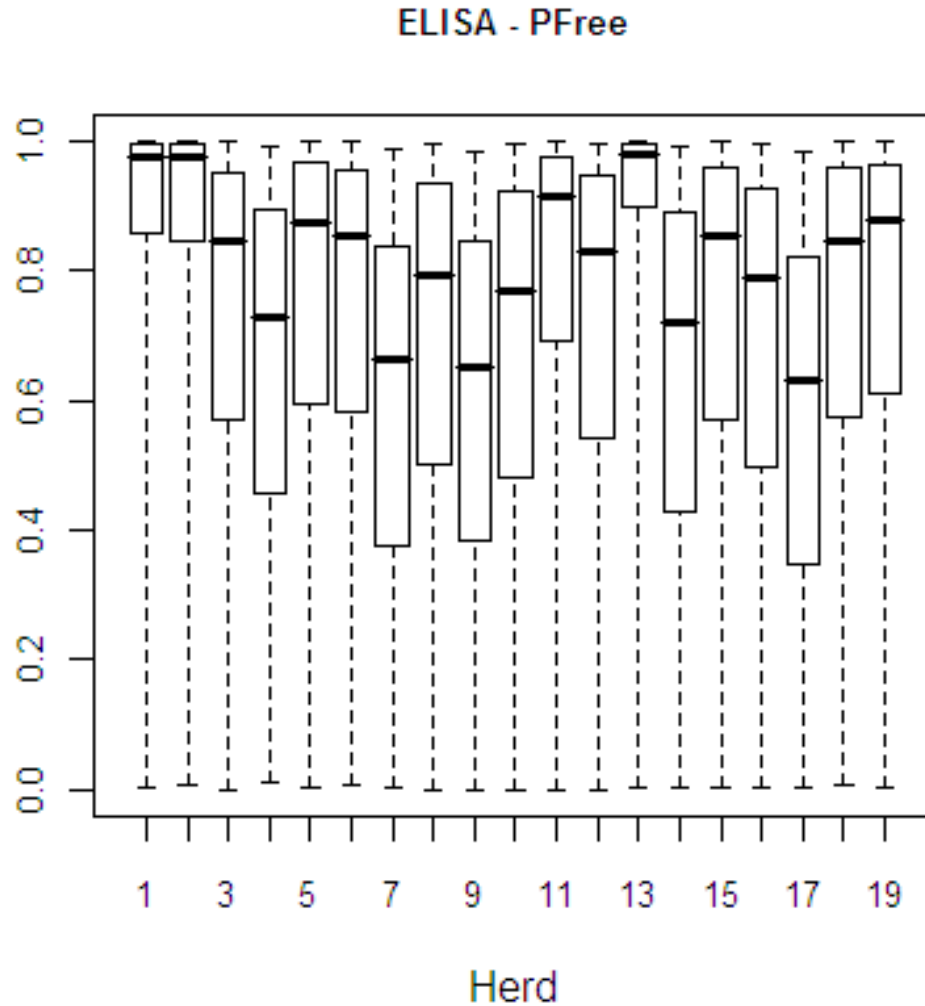
- Only 2/19 herds had high PFree values
- 12/19 PFree = 0
- 5/19 had intermediate values
- 2 infected herds had high PFree
 - 1 had prior testing, possibly biased result
 - 1 low-prevalence only confirmed 1 year later
- Map never cultured from 1 herd with low PFree – why?



Design prevalence issues

- Distribution rather than point value reflects industry situation
- Generally high (mean 34%) compared to more acute diseases
- Some herds likely to be infected at low prevalence despite high PFree
- May need to review and decrease if prevalence drops as control improves
- Problems with identifying high PFree herds at lower DP – ? update estimates over time

PFree for DP=20% in simulated uninfected herds





Issues to consider

- Good for identifying infected herds, not so good for identifying uninfected herds (better tests required)
- Poor power in smaller herds
- Updating PFree estimates with repeated testing:
 - Discounting value of historical data
 - Lack of independence between repeated tests in individual herds
- Calculation based on probability of actual result, not on hypothetical cut-off (not HSe/HSp)



Acknowledgment

- Project funded by the International EpiLab, Denmark