

Within-herd transmission of *Mycoplasma bovis* infection in 20 Dutch dairy herds

Marit Biesheuvel¹, Caitlin Ward², Patty Penterman³, Erik van Engelen³, Gerdien van Schaik^{3,4}, Rob Deardon², Herman Barkema¹ ¹Faculty of Veterinary Medicine, University of Calgary, Calgary, Canada ²Department of Mathematics and Statistics, Faculty of Science, University of Calgary, Calgary, Canada

³Royal GD, Deventer, The Netherlands

⁴Department of Farm Animal Health, Faculty of Veterinary Medicine, Utrecht University, Utrecht, The Netherlands

Background

- *Mycoplasma bovis:* naturally resistant to antibiotics targeting cell wall synthesis
- **PREVALENT** worldwide in dairy and beef industry

Objectives

- Estimate the within-herd transmission rate between calves, youngstock and cows
- 2. Identify the **most likely** withinherd transmission **pathways**

Materials and Methods

Longitudinal data were collected on **20 clinically infected dairy farms** in the Netherlands between 2016 – 2017.

• **SYMPTOMS**: mastitis or arthritis in cows



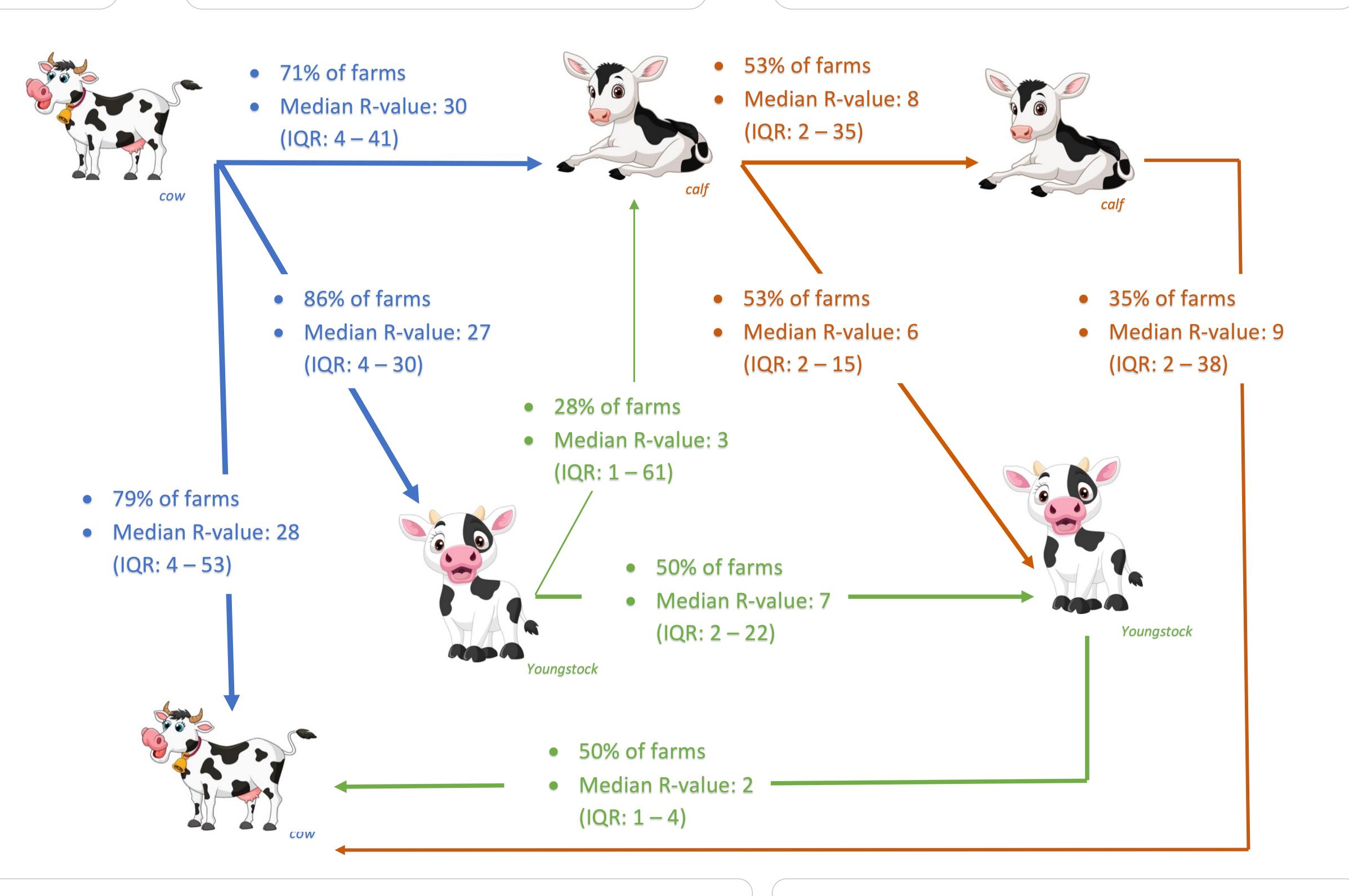
- **CAUSES** <u>mastitis</u> or <u>arthritis</u>, but also <u>pneumonia</u>, keratoconjunctivitis, otitis media, abortion, subcutaneous abscesses
- RISK FACTORS: purchase of animals, increased stress levels, immunosuppression.
- IMPACT: <u>High economic impact</u> hard to treat and often results in culling
- **TRANSMISSION:** During milking and noseto-nose contact, or through environment

 Identify farm management factors explaining transmission rates and pathways.



- **SAMPLING:** Each farm was sampled 5 times in 3 months
- **RISK ASSESSMENT:** Farm management factors were assessed for each farm

ANALYSIS: An age-stratified SIR model on 3 aggregated individual test results to calculate disease transmission rates and pathways, and a Fisher's exact test to explore potential explanatory farm factors



Results

• <u>R-value (transmission rate</u> (β) / removal rate (γ)):

'The average number of secondary infections in age group *i* caused by 1 infectious animal in age group *j* over the entire course of its infectious period in an otherwise fully susceptible herd'.

Conclusions

- Transmission was highly variable among herds and cattle age groups, with highest transmission rates from cows to cows, youngstock and calves.
- Most important transmission pathways: cow-to-cow, cow-to-calf, cow-to-youngstock. But also, calf-to-calf, calf-to-youngstock, youngstock-to-youngstock and youngstock-to-cows.
- Associated farm factors: internal biosecurity (separate caretakers for different age groups, number of people involved), external biosecurity (contractors, external employees) or indirect transmission routes (number of feed and water stations).

Take away message

- First study to estimate transmission rates for *M. bovis* and results will help better understand the on-farm intervention strategies regarding *M. bovis* outbreaks.
- Future steps: design a Bayesian individuallevel model to account for low test sensitivity and use the 3 diagnostic results separately without manufacturer cut-off

