

Simulation of Salmonella Dublin infection dynamics in dairy herds



Liza Rosenbaum Nielsen¹, Anne Braad Kudahl², Søren Østergaard²

- ¹ Department of Large Animal Sciences, University of Copenhagen, Denmark, Irn@life.ku.dk
- ² Department of Animal Science, Foulum, Aarhus University, Denmark

BACKGROUND

National prevalence of test-positive dairy herds reduced from 26% to 8.3% from 2002 to 2012

Growing demand for decision support for control strategies in the Danish cattle industry to target control options in the remaining infected dairy herds

OBJECTIVE

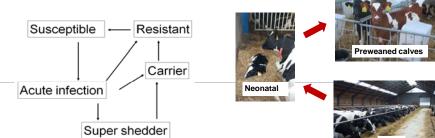
To construct a simulation model to mimic real life infection dynamics of Salmonella Dublin in Danish dairy herds with different herd sizes, hygiene and management levels



METHODS

Age-structured stochastic, mechanistic and dynamic simulation model (Dublin-Simherd)

5 infection states, 6 age groups





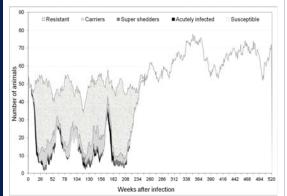
Breeding heifers



Key facts about the modelling and the scenarios

Weekly time steps,10-years simulations, 1000 iterations + sensitivity analyses
Infection state probabilities, morbidity, mortality mainly from field data and literature
Herd susceptibility determined by management. Number of contacts per week determined by hygiene level
Introduction of infection to a naïve dairy herd by purchase of one acutely, asymptomatically infected heifer
48 scenarios combining 4 hygiene levels, 4 herd susceptibility levels 3 herd sizes (70 cows, 200 cows and 400 cows)

RESULTS



Example of infection dynamics among growing heifers in one iteration out of 1000

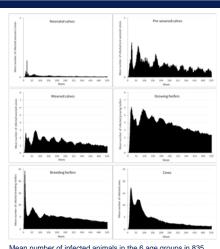
Reduced hygiene and increased herd susceptibility lead to

- increased probability of spread
- increased duration of infection
- larger epidemic size
- decreased probability of extinction

Common for S. Dublin to become endemic – in particular in large herds (>200 cows)

Persistent carriers

but not super shedders –
 were required in the model to mimic real life infections



Mean number of infected animals in the 6 age groups in 835 iterations with spread of Salmonella Dublin in a simulated 200 cow dairy herd. Note the multi-peak patterns, representing repeated waves of infections a approximately yearly intervals