Tutorial 1: Designing a mathematical model for bovine TB



Background information

Bovine TB – a chronic infectious disease

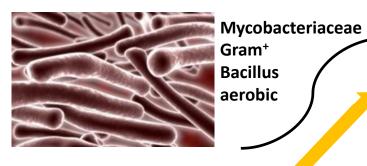
4th most significant livestock disease in developing countries (Perry et al. 2002)

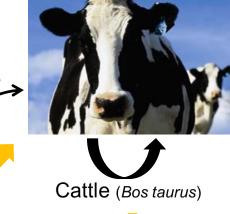


- One of the most persistent animal health problems
 - Endemic in many countries
 - Huge financial losses (total costs over £275 million per annum)
- An increasingly important public human health concern
 - » zoonotic transmission
 - » 10-15% of human TB cases caused by bTB in developing world

Bovine TB: Transmission routes

Mycobacterium bovis





zoonotic

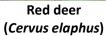


aerosol

droplets

milk



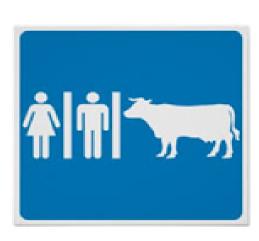




Badgers (Meles meles)



Possums (Trichosurus vulpecula)





bTB Diagnostics

Main diagnostic test on life animals: Skin test

- Single Intradermal Comparative Cervical Test (SICCT)
- > 99% Specificity (i.e. very few false positives)
- ~ 55-70% Sensitivity (i.e. many false negatives: many infected animals are 'non-reactors')
 - Infected animals only react to the test after 30-50 days post infection
 - Test outcome also compromised by co-infection (e.g. fluke, Para-TB)

Abattoir inspection:

- Tubercles (lesions) in lungs and lymph nodes
- ~30-50% Sensitivity

Many undetected cases!







Current control strategies (UK)

Aim: make UK officially bTB free by 2038

- Routine herd testing & culling of infected cattle; movement restrictions until herd is declared bTB free
 - Strategy not effective in the presence of wildlife reservoir
 - Controls focusing on a single transmission route are unlikely to be effective (Brooks-Pollock et al. 2014)

Badger culling

- Inconclusive results from Randomized Budger Culling Trial
- Not effective in the long-term (Jenkins 2010)

Vaccines:

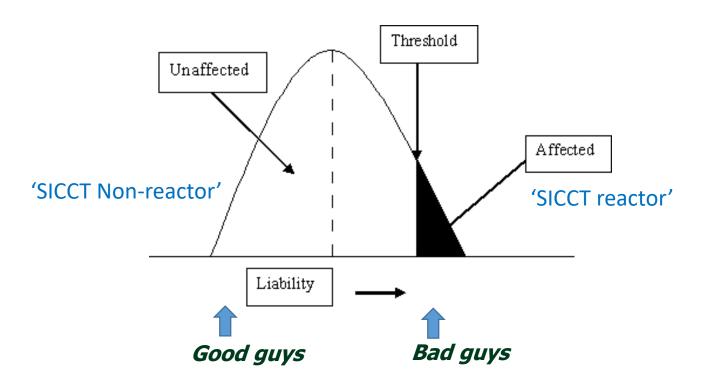
- attenuated vaccines; not fully protective
- vaccinated animals indistinguishable from infected animals using standard tuberculin tests
- No vaccine applied currently



Several millions \$/yr spent on ineffective control measures Complementary strategies needed!

Genetic selection for bTB resistance

- Strong evidence for genetic variation in bTB resistance of cattle
 - Heritability: 0.08-0.23; resistance controlled by many genes
 - Prediction accuracy: 72%
- Genomic selection of animals with low genetic risk for bTB infection implemented in the UK (2016)



Question:

To what extent can genetic selection help to eliminate bovine TB in cattle within the next 20 years?

Task:

Design a mathematical model to answer this question & to serve as decision making tool for developing efficient bTB control strategies