

A mathematical model for interpretation of antibody titre distribution

A.F. Viet ^{a,b}, G. Medley ^b

^a Unit of Animal Health Management, Veterinary school & INRA, BP 40706, 44307 Nantes Cedex 03, France (corresponding author)
^b Ecology and Epidemiology group, Dept. Biological Sciences, The University of Warwick, Coventry, CV4 7AL, United Kingdom

Context and objective

Measurements of individual antibody titres define of a distribution of antibody titres at the herd level (e.g. Fig.1 with 3 antibody titre levels).

This distribution can potentially contain information on time since virus introduction

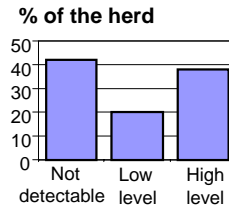


Fig 1. Example of antibody titre distribution

Objective :

To define a modelling framework to interpret distribution of antibody titres at herd level in terms of possible period of time since pathogen introduction.

Mathematical model

- Stochastic SIR model with K antibody titre levels ($K=3$ in Fig.2).
- Dynamic of the antibody titre:
 - Increase after an infection (I) or a re-exposure (R),
 - Decrease if maternal antibodies (S) and after the infection (R).
- Horizontal transmission (rate $\lambda(S_\alpha \rightarrow I_\alpha)$) : frequency-dependent form, taking into account a variability of the sensibility (q_α) and of the excretion (p_α) related to the antibody titre level α .

$$\lambda(S_\alpha \rightarrow I_\alpha) = q_\alpha c \left(p_1 \frac{I_1}{N} + p_2 \frac{I_2}{N} + p_3 \frac{I_3}{N} \right) \begin{cases} 0 \leq p_3 \leq p_2 \leq p_1 \leq 1 \\ 0 \leq q_3 \leq q_2 \leq q_1 \leq 1 \end{cases}$$

where I_α is the number of infected animal with the antibody titre level α and N the total number of animals.

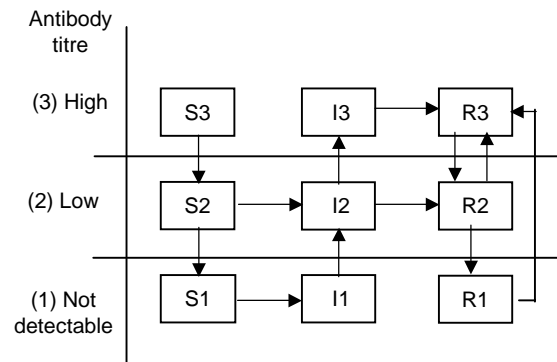


Fig 2. Diagrammatic representation of the model: $S_\alpha \leftrightarrow$ susceptible with the antibody titre level α , $I_\alpha \leftrightarrow$ infected with the antibody titre level α , $R_\alpha \leftrightarrow$ recovered with the antibody titre level α ; $\alpha \in \{1 \text{ (not detectable)}, 2 \text{ (low)}, 3 \text{ (high)}\}$.

Model use

- Definition of the model for the considered pathogen and the population
- Simulation of the model
- For a given distribution (e.g. Fig.1), definition over time of the probability to have the observed distribution (e.g. Fig. 3 with 5000 simulations)

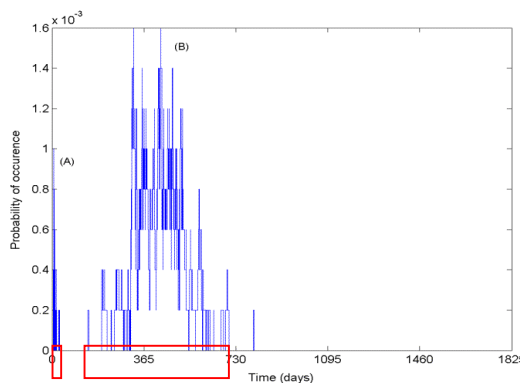
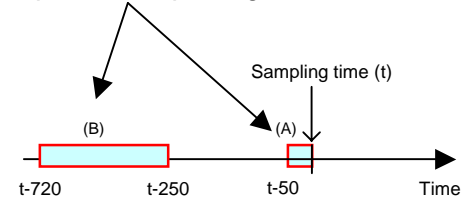


Fig 3. Proportion of simulations, over time, where the considered distribution was observed

- Identification of most likely periods of pathogen introduction:



Discussion and conclusion

- Uses the dynamics of antibody titre distribution.
- Sensitivity and specificity of the test need to be considered.
- Other possible use: Determination of the probability that the pathogen was present at the time of sampling.