

Bayesian data assimilation for vector borne disease response: *Theileria orientalis* (Ikeda) in NZ cattle.

Chris Jewell

CHICAS
Lancaster University
c.jewell@lancaster.ac.nz

16th April, 2015

Lancaster
Medical School



MASSEY UNIVERSITY
TE KUNENGA KI PŪREHUROA
UNIVERSITY OF NEW ZEALAND

- 1 Motivation
- 2 Modelling
- 3 Inference and forecasting
- 4 *T. orientalis* Ikeda prediction
- 5 Roadmap

Outline

- 1 Motivation
- 2 Modelling
- 3 Inference and forecasting
- 4 *T. orientalis* Ikeda prediction
- 5 Roadmap

Motivation

Welfare and Economics

- **Foot and Mouth Disease**
 - 2001: £8 billion, 6.5 million slaughtered
 - 2007: £100 million, 2610 slaughtered
- **Avian Influenza**
 - worth >3.5 billion
 - 40% UK primary meat market (2004)
- **bTB?**
- **Enterobacteriaceae?**

- Casts **doubt** on food safety



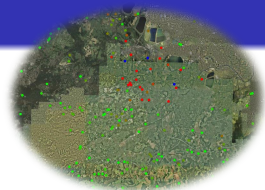
Epidemic Control

Framework Response Plan for Exotic Animal Diseases:

- 1 Minimise the number of animals which need to be culled either to control the disease or on welfare grounds, and which keep animal welfare problems to a minimum.
- 2 Protect public health.
- 3 Cause the least possible disruption to the food, farming and tourism industries, to visitors to the countryside, and to rural communities in the wider economy.
- 4 Minimise damage to the environment.
- 5 Minimise the burden on taxpayers and the public.

Defra 2007

Epidemic control

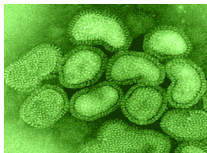


Forecasts of disease spread inform control:

- Provide **real-time** predictions of **risk**
 - Understanding the **determinants** of transmission
 - How many **occult** (undetected) infections are there?
 - Who is likely to be **infected next**?
- Explore a full range of possible outcomes **given the current state of the epidemic**
 - Movement restriction
 - Vaccines
 - Culls

Emerging infectious diseases

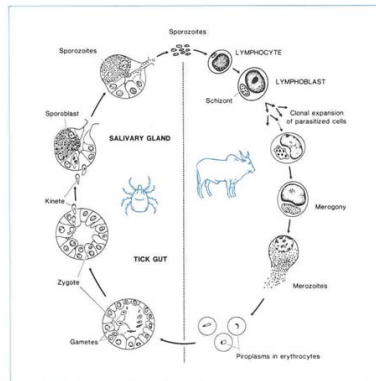
- Broad definition - Brown, C (2004) *Rev. sci. tech. Off. int. Epiz* **23**:435
 - a known agent appearing in a new geographic area
 - a known agent or its close relative occurring in a hitherto unsusceptible species
 - a previously unknown agent detected for the first time.
- Our only experience of how they behave is the **current epidemic**
- Model parameters may be **uncertain**



Theileria orientalis Ikeda

NZ Ministry for Primary Industries contract, Massey EpiCentre

- Protozoal vector borne disease
 - Host: **Cattle**
 - Vector: **Tick** *H. longicornis*
- Endemic, but...
- August 2012 new virulent subspecies **Ikeda**
- Case morbidity < 35%
- Mortality \approx 1%
- Cost NZ\$25k per farm

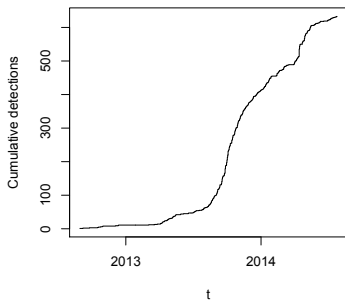


Theileria in NZ cattle

Questions

- What are the main determinants of transmission?
 - **Environmental** (spatial) spread
 - Spread via **NAIT** network
 - Importance of **tick presence**
- Where might any **undetected infections** be?
- **How fast** will the epidemic spread?

Theileria in NZ cattle



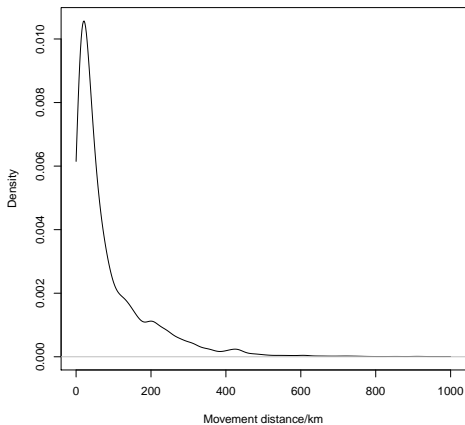
Epidemic at 1st Aug 2014

- 655 Case detections
- 136837 Farm locations
- Dairy/non-dairy
- NAIT cattle movement network frequencies

Databases: AgriBase, FarmsOnLine, NAIT

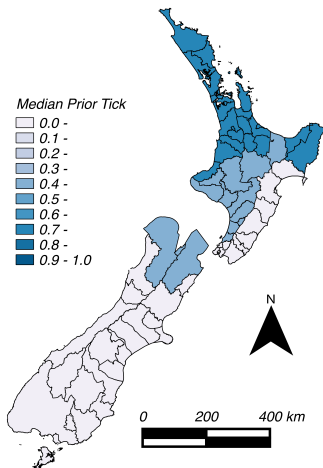
National Animal Identification and Tracing (NAIT)

- Mandatory tracing of cattle and deer
- Began 2012
- 517328 recorded movements
- 0.0003% connectivity
- Represented as sparse matrix



Theileria in NZ cattle

Tick habitat



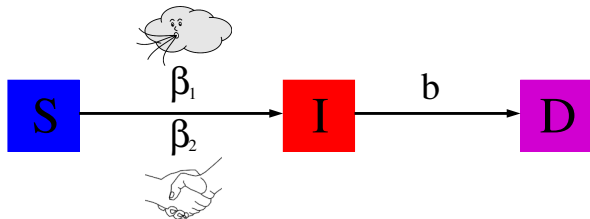
- Samples tested from BVD surveillance
- All *T. orientalis* spp.
- Aggregated at TLA level
- Prior risk: Alan Heath

Outline

- 1 Motivation
- 2 Modelling**
- 3 Inference and forecasting
- 4 *T. orientalis* Ikeda prediction
- 5 Roadmap

SID models & population heterogeneity

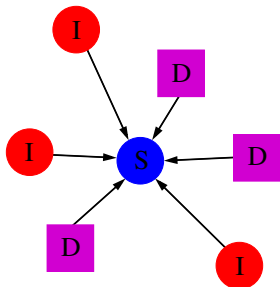
- 1 Individual herds **infect** each other
 - Spatial rate β_1
 - NAIT network β_2
- 2 Once infected, herds are infectious forever
 - **Detected** at rate b



$$\beta_1 = ?, \beta_2 = ?, b = ?$$

Infection process

- At any time t , susceptible j has **infectious pressure** exerted on it by
 - all infected or notified farms $i \in \mathcal{I}(t), \mathcal{D}(t)$
 - “Background” (not explicitly modelled)



In a small interval $\Delta t = (t, t + \delta]$:

$$P(j \text{ infected}) \approx \lambda_j(t) \cdot \Delta t$$

$$\lambda_j(t) = \epsilon + \sum_{i \in \mathcal{I}(t)} \lambda_{ij}(t) + \sum_{i \in \mathcal{N}(t)} \lambda_{ij}^*$$

Transmission model

SID model

$$\begin{aligned}\lambda_{ij}(t) &= s(t; \alpha, \nu) \zeta^{\kappa_j} p_{k(j)} [\beta_1 K(i, j; \delta) + \beta_2 c_{ij}], \quad i \in \mathcal{I}, j \in \mathcal{S} \\ \lambda_{ij}^*(t) &= \beta_{ij}(t), \quad i \in \mathcal{N}, j \in \mathcal{S}\end{aligned}$$

$$K(i, j; \delta) = \frac{\delta}{(\delta^2 + \|x_i - x_j\|^2)^{1.2}}$$

$p_{k(j)}$ = tick occurrence in TLA k

ζ = effect of dairy cf. beef

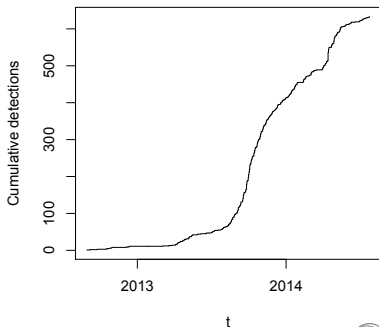
$$D - I \sim \text{Gamma}(4, b)$$

Unknowns

$$\theta = \{\beta_1, \beta_2, \delta, \mathbf{p}, \zeta, \alpha, \nu, b\}$$

Seasonality

- Biannual peak incidence – autumn/spring
 - Due to **vector ecology**
-
- Vector seasonality
 - Smooth? (common sense!)
 - On/off threshold? (literature*)
 - Candidates:
 - Periodic piecewise cubic spline
 - Periodic square wave



*e.g. Stafford KC (1994), Ogden *et al.* (2004)

Tick model – dynamics

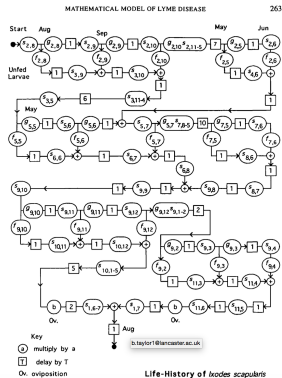
IMA Journal of Mathematics Applied in Medicine and Biology (1999) **16**, 261–296

A mathematical model of the ecology of Lyme disease

TRAVIS C. PORCO

*Community Health Epidemiology and Disease Control, 25 Van Ness Avenue, Suite 710,
San Francisco Department of Public Health, San Francisco,
California 94102, USA*

[Received 14 February 1994 and in revised form 17 June 1998]



Tick model – proxy sampling

- BVD samples collected during 2013
- Tested for *Theileria orientalis* spp.
- Implies **Binomial** sampling model, TLAs $k = 1, \dots, m$

$$x_k \sim \text{Binomial}(n_k, p_k)$$

n_k herds sampled, x_k *Theileria* +ve, $p_k \propto$ tick occurrence

- Independent sampling \rightarrow joint likelihood
- Robust to test **Sensitivity** and **Specificity**.

Outline

- 1 Motivation
- 2 Modelling
- 3 Inference and forecasting**
- 4 *T. orientalis* Ikeda prediction
- 5 Roadmap

Approach to inference

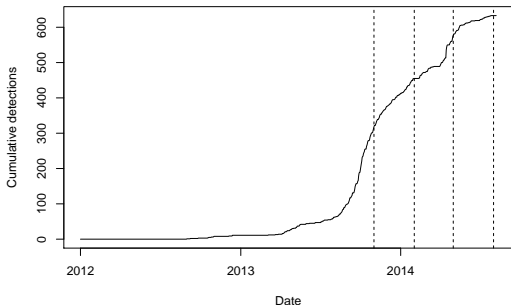
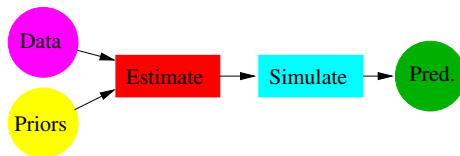
Bayesian approach for risk forecasting

$$P(\theta|X) = \frac{L(X|\theta)f(\theta)}{\int_{\Theta} L(X|\theta)f(\theta)d\theta}$$

- Coherent inclusion of **Prior** information
- **Likelihood** assimilates *all* data
- **Posterior** encodes **uncertainty** → **Predictive** distribution
- Complicated integral → MCMC
 - Unobserved infection times
 - Occult infections

Approach to real time forecasting

At any point **during** an epidemic:



Approach to inference

- **Bayesian** approach for risk forecasting

$$\begin{aligned}
 L(\boldsymbol{\theta} | \boldsymbol{I}, \boldsymbol{D}) &= \sum_{j: I_{\kappa} < I_j < T_{obs}} \left(\lambda_j(I_j^-) \right) - \exp \left[\int_{I_{\kappa}}^{T_{obs}} \left(\sum_{j \in \mathcal{P}, j \neq \kappa} \lambda_j(t) \right) dt \right] \\
 &\times \prod_{k=1}^m p_k^{x_k} (1 - p_k)^{n_k - x_k} \\
 &\times \sum_{j: D_j \leq T_{obs}} (f_D(D_j - I_j)) + \sum_{j: D_j > T_{obs}} (1 - F_D(T_{obs} - I_j))
 \end{aligned}$$

- Jump algorithm integrates over dimension of \boldsymbol{I}
- GPU as a **likelihood coprocessor**
 - NVIDIA CUDA software libraries
 - Parallelise likelihood **within** MCMC.

MCMC algorithm

Repeat the following steps

- 1 Model parameters (adaptive logRWMH)
 - 1 Update $\{\beta_1, \beta_2, \zeta, \epsilon, \delta\}, \mathbf{p}, \alpha, \nu$
 - 2 Update b
 - 3 Update b with $U_i = bD_i \sim \text{Gamma}(4, 1)$ (non-centred)
- 2 Infection times - repeat z times:
Equal probability
 - 1 Move $I_s \rightarrow I_x^*$
 - 2 Add $\{I + s\} \rightarrow I^*$
 - 3 Delete $\{I - s\} \rightarrow I^*$

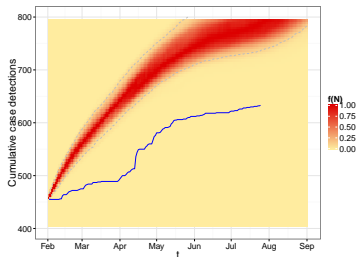
Outline

- 1 Motivation
- 2 Modelling
- 3 Inference and forecasting
- 4 T. orientalis Ikeda prediction**
- 5 Roadmap

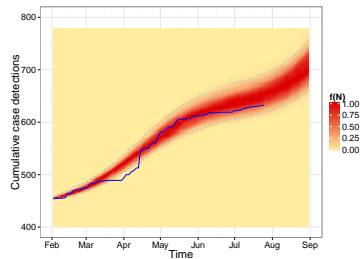
Theileria in NZ cattle

Prediction: in sample cumulative detections

Cubic Spline



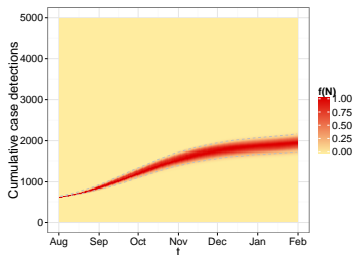
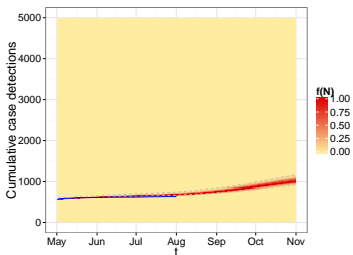
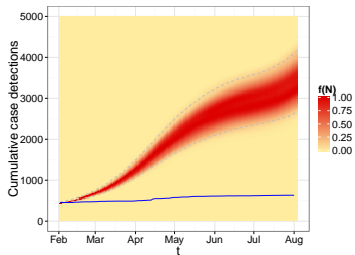
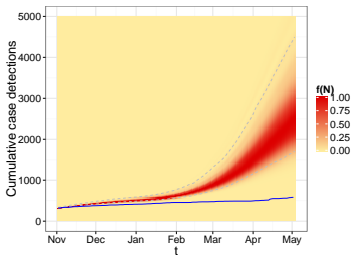
Square Wave



Analysis at 1st Aug 2014, simulate from 1st Feb 2014

Theileria in NZ cattle

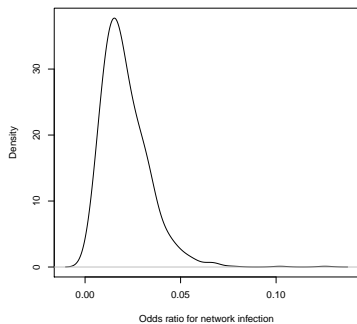
Prediction: out of sample cumulative detections – justification for square wave



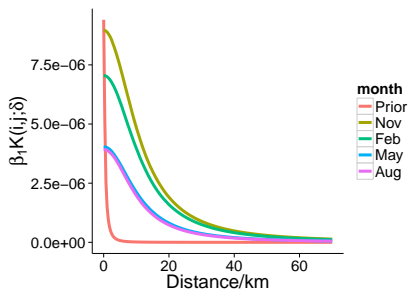
Theileria in NZ cattle

Parameter estimation

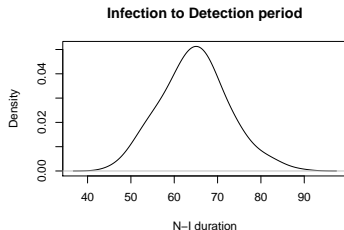
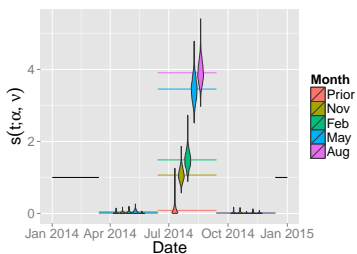
NAIT vs Environmental spread



Distance Kernel



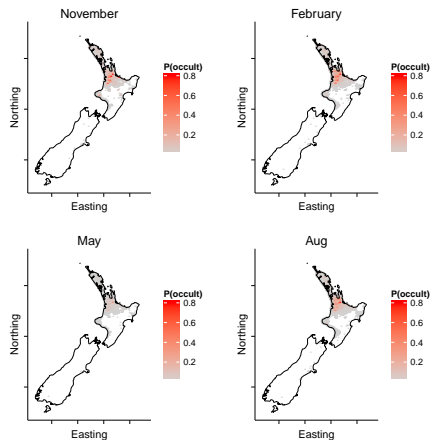
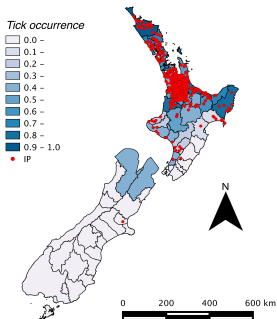
Seasonality and infectious period



Caveat: non-identifiability between phase and infec-detect time!

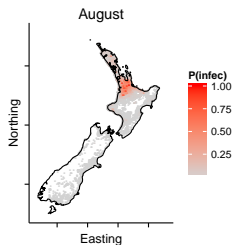
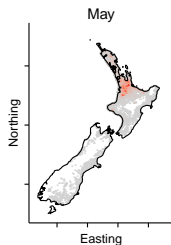
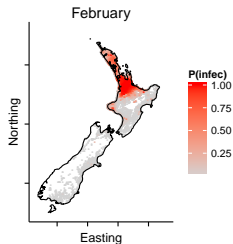
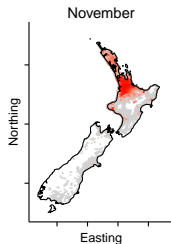
Theileria in NZ cattle

Tick occurrence probability and occults



Theileria in NZ cattle

Prediction: 6 month infection risk



Outline

- 1 Motivation
- 2 Modelling
- 3 Inference and forecasting
- 4 *T. orientalis* Ikeda prediction
- 5 Roadmap**

Theileria in NZ cattle

Conclusions

- **Environmental** spread main driver
 - **NAIT** spread limited (currently)
 - Caveat – AgriBase, FOL, NAIT joining issues
- Epidemic **appears** to be slowing
 - Complacency vs. effective pop size.
- Modelling issues
 - **Ticks**
 - Seasonality – tie in with climate data (resolve $Cor(b, \nu)$)
 - e.g. MAXLIK and MAXENT, etc.
 - **Endemic stability**
 - Require **SIDS** model
 - Within-farm epidemic
- Lack of **coherent** livestock databases (CEBRA/MPI)

Thanks to...

- IFS
 - Richard Brown

- MPI
 - Daan Vink
 - Kevin Lawrence
 - Andy McFadden
 - Mary van Anandel

- AsureQuality
 - Robert Sanson

