A movement rewiring algorithm applied to epidemic spread among bovines



SCIENCE & IMPACT

Vétérinaire, Agroalimentaire et de l'Alimentation

Nantes Atlantique

École Nationale

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ModStatSAP meeting – 12/03/2019



Context and objectives

Hypothesis: cattle movement through commercial exchanges is a major factor of disease transmissions between farms

Question: How can we change the movements to reduce infection risks without reducing cattle exchange ?

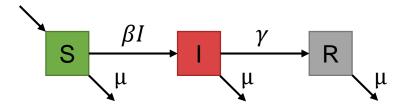
Movement rewiring algorithm

Tested on data-based movements +

mechanistic epidemic model

The computational model

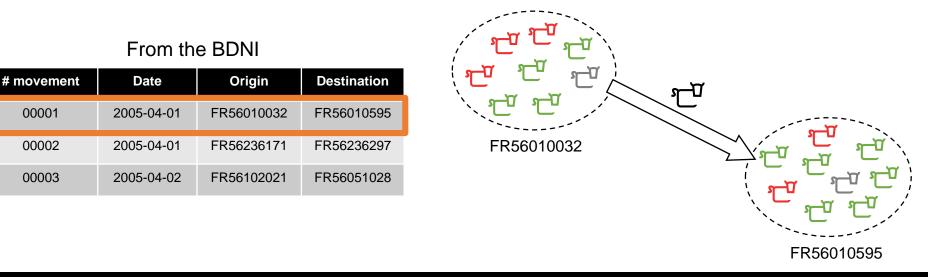
Intra-herd dynamics without variation in herd size



SIR model with 3 parameters:

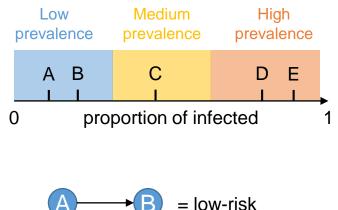
- The infection rate β
- The mean infection duration $1/\gamma$
- The death rate μ

Inter-herd dynamics based on real data



The rewiring algorithm

Each herd is assigned a **prevalence class**:



Movements are **high risk** if the prevalence class of the source is **higher** than the prevalence of the destination A = low-risk



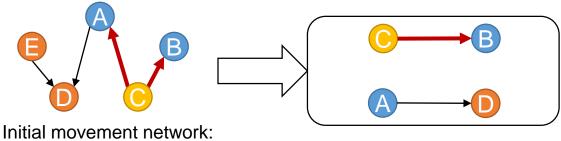
The objective: change the source of high-risk movements to create low-risk movements

• The algorithm works on the movements occurring on a given day only

under 4 constrains:

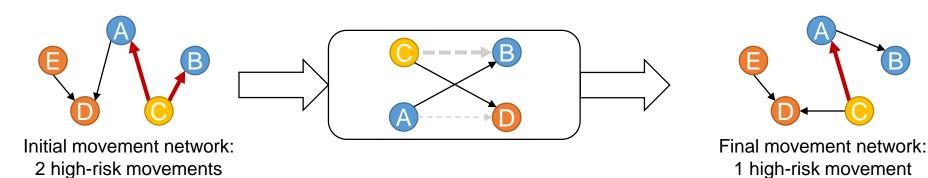
- The (in- and out-) degrees of each herd must not change
- No loops
- The prevalence class of the new source is always lower for high-risk movements

• Step 1: swap the origins of two movements

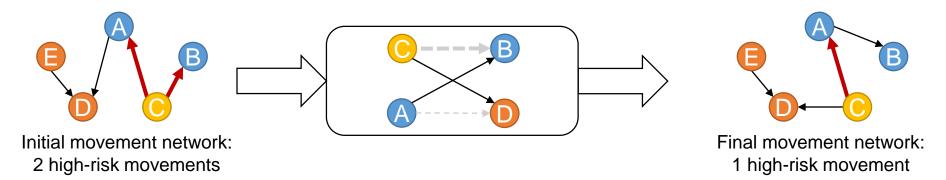


Initial movement network: 2 high-risk movements

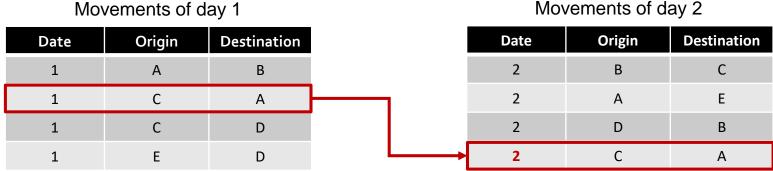
• Step 1: swap the origins of two movements



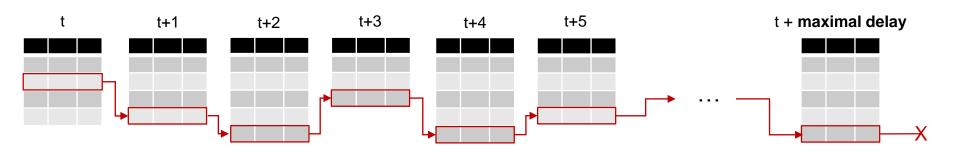
Step 1: swap the origins of two movements ٠

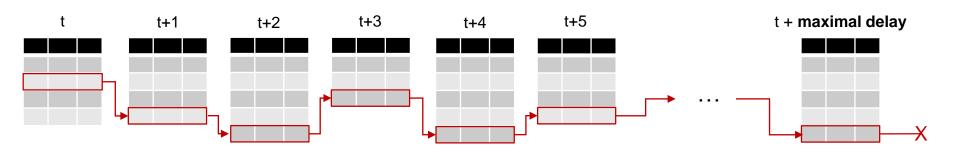


Step 2: push back the remaining high-risk movements to the next day ٠

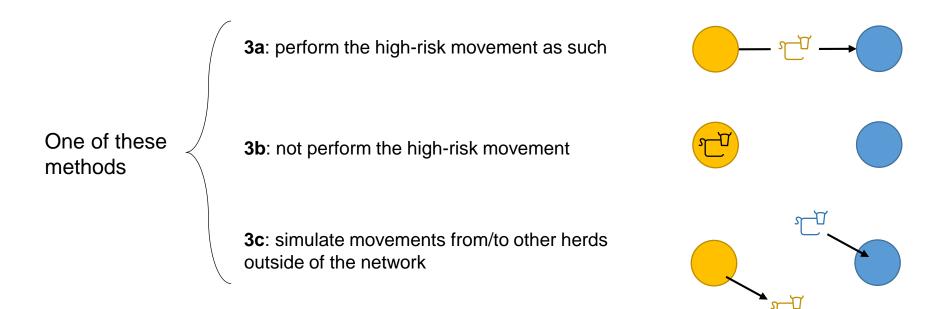


Movements of day 2





• Step 3: managing the high-risk movements reaching the maximal delay



Testing the algorithm



What ?



Prim'Holstein

Where?

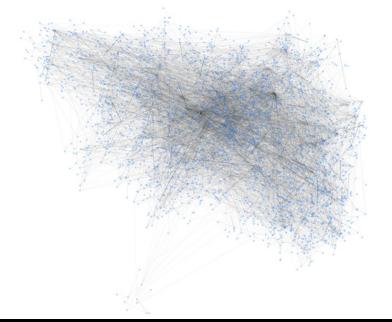


Morbihan

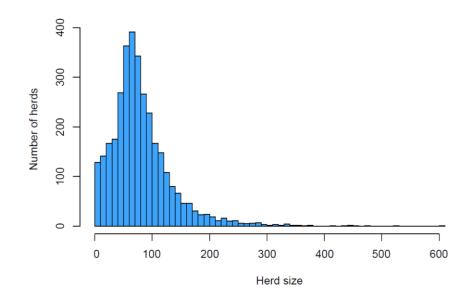
When ?



15429 movements between 3300 herds



Herd sizes from 1 to 605 animals



Parameters of the simulation

3 sets of parameters considered:

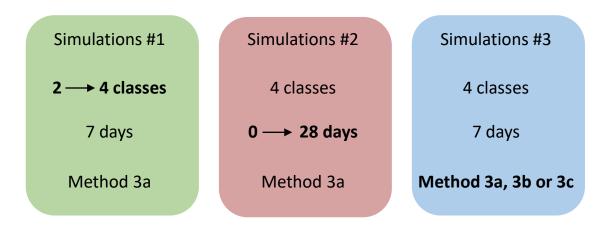
- Number of classes of prevalence
- Maximal delay in step 2
- Method used in step 3

Other parameters:

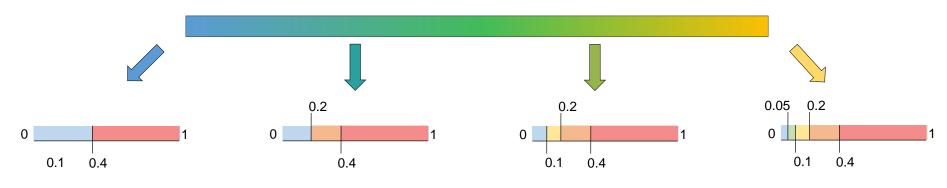
• Epidemiological parameters: $\beta = 0.005$, $1/\gamma = 50$, $\mu = 0.02$

R varies between 0.05 (*N* = 1) and 30.25 (*N* = 605)

- 660 herds (20%) infected initially (chosen at random)
- 25% of status / in the herds infected initially
- 500 repetitions per scenario

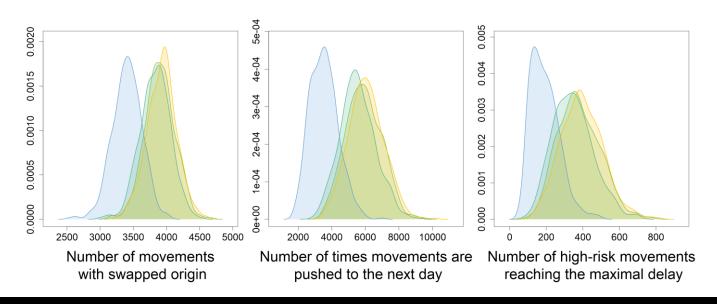


2 to 5 classes of prevalence

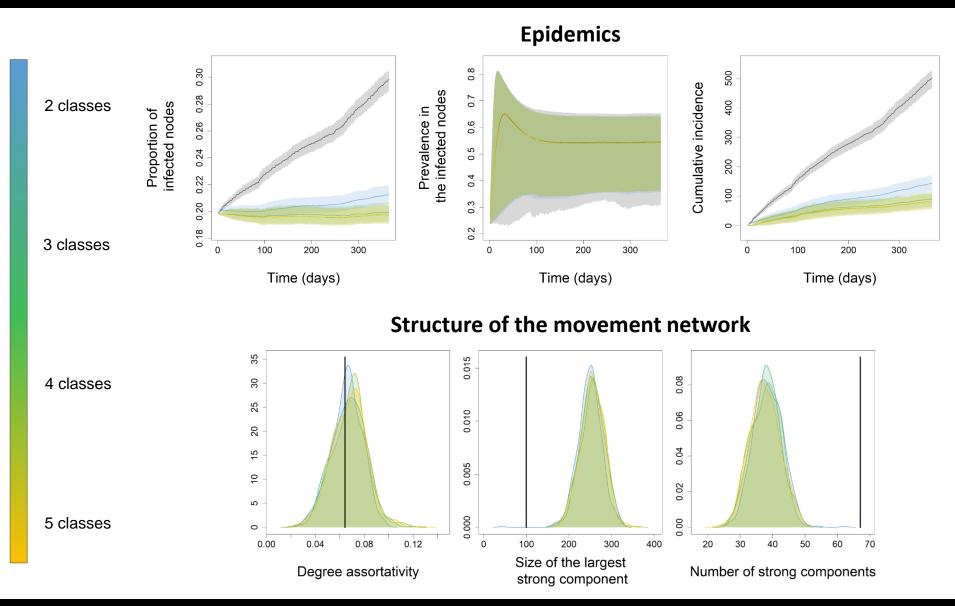


(+ Maximal delay = 7 days + Method = 3a: high-risk movements performed)

Algorithm functionning

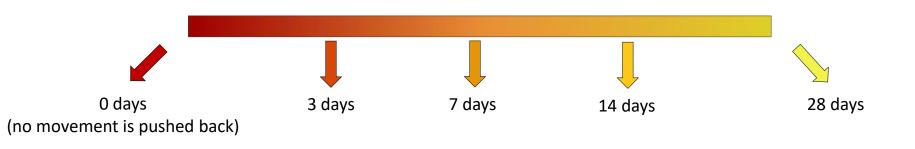


2 to 5 classes of prevalence



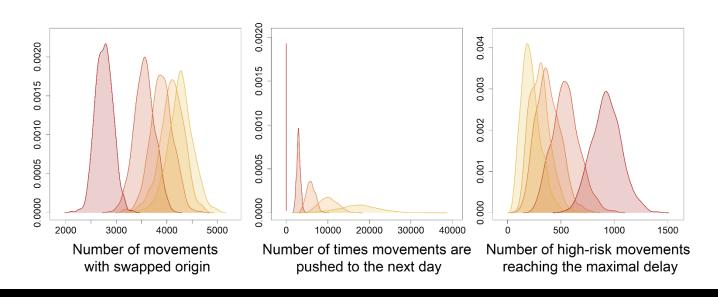
10/17

0 to 28 days of maximal delay

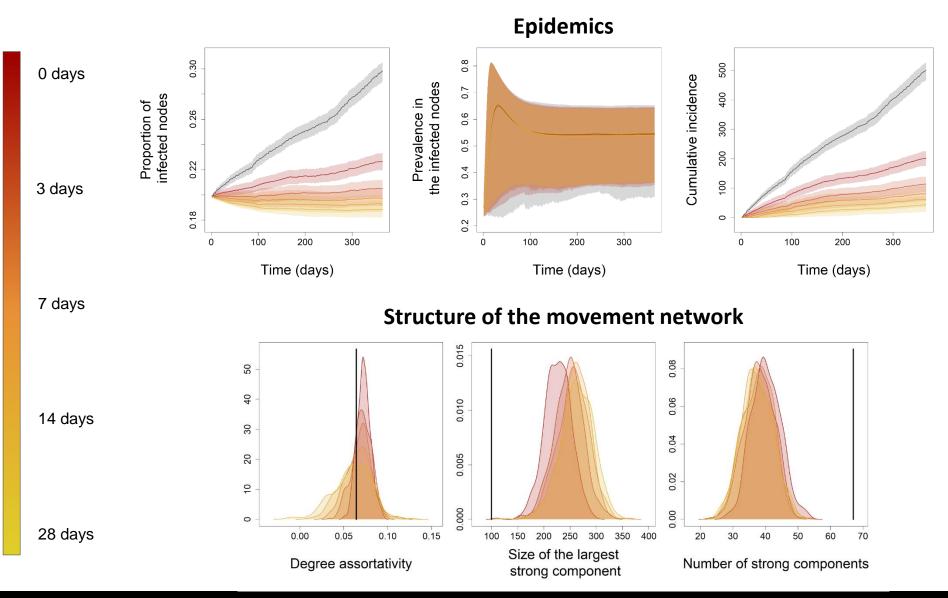


(+ 4 classes of prevalence + Method = 3a: high-risk movements performed)

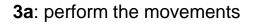
Algorithm functionning



0 to 28 days of maximal delay



3 methods for the last high-risk movements

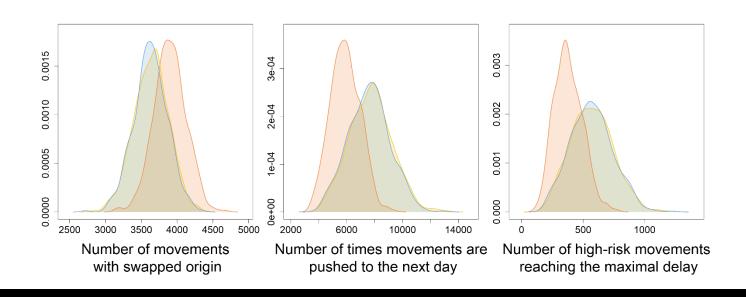


3b: remove the movements

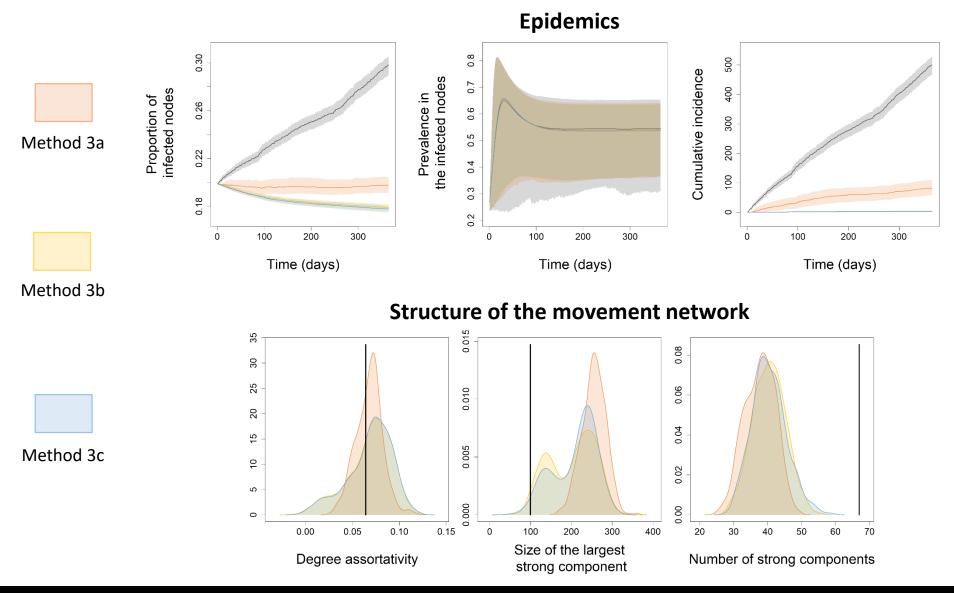
3c: move to the outside

(+ 4 classes of prevalence + Maximal delay = 7 days)

Algorithm functionning



3 methods for the last high-risk movements



Perspectives

Sensitivity analyses

- to the three epidemiological parameters (β , γ and μ)
- to the rewiring parameters

Mimic real bovine diseases

• which parameters result in a « realistic » infection without rewiring?

Different ways to handle the prevalence classes

- how often are they updated? (*daily in the current model*)
- which error on the estimation of prevalences (*no error in the current model*)

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THANK YOU



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