

Why using epidemiological models to evaluate control strategies for livestock infectious diseases?

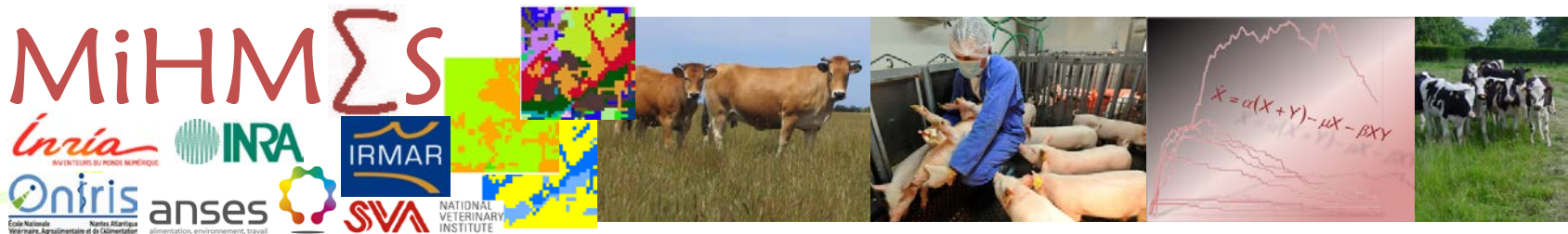
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Infectious diseases in livestock

- Animal health & welfare
- Zoonoses
 - ➔ Food safety
 - ➔ Veterinary public health



- ➔ Economic losses in animal productions
 - ➔ Worse animal performances, production losses, culling
- ➔ **Effective preventive and control measures**

Infectious diseases in livestock

- What need to be understood?
 - Which pathogen, which source / origin?
 - Which transmission route(s)?
 - Direct, indirect (vector, environment, wind, ...), vertical (*in utero*), etc.
 - Which host and environmental factors favouring?
 - Infection transmission, clinical sign appearance (disease)
 - Which efficient control measures?
 - Vaccination, test & cull, hygiene, etc.



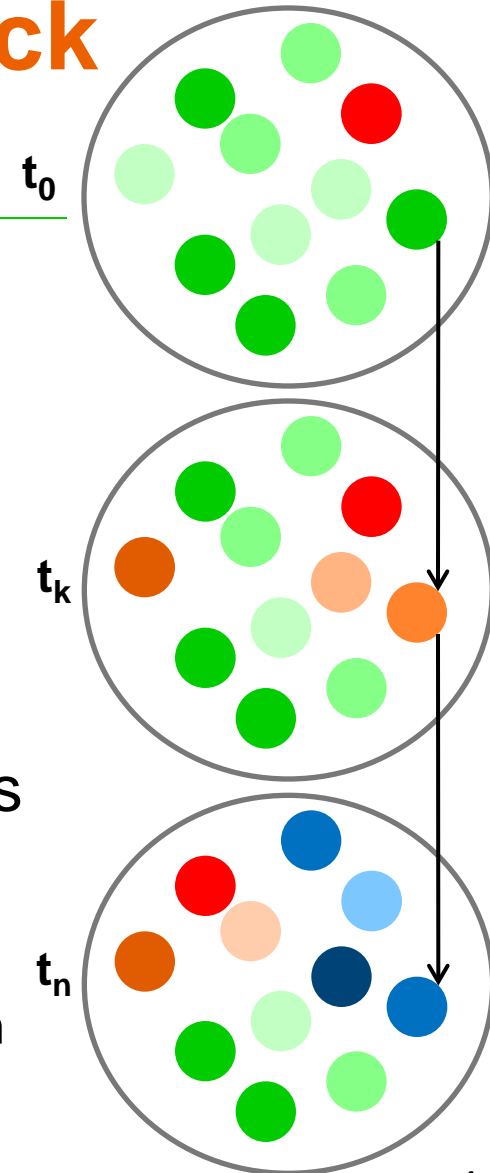
→ **Dynamic spread of infectious pathogens in a host population under various scenarios**

Infectious diseases in livestock

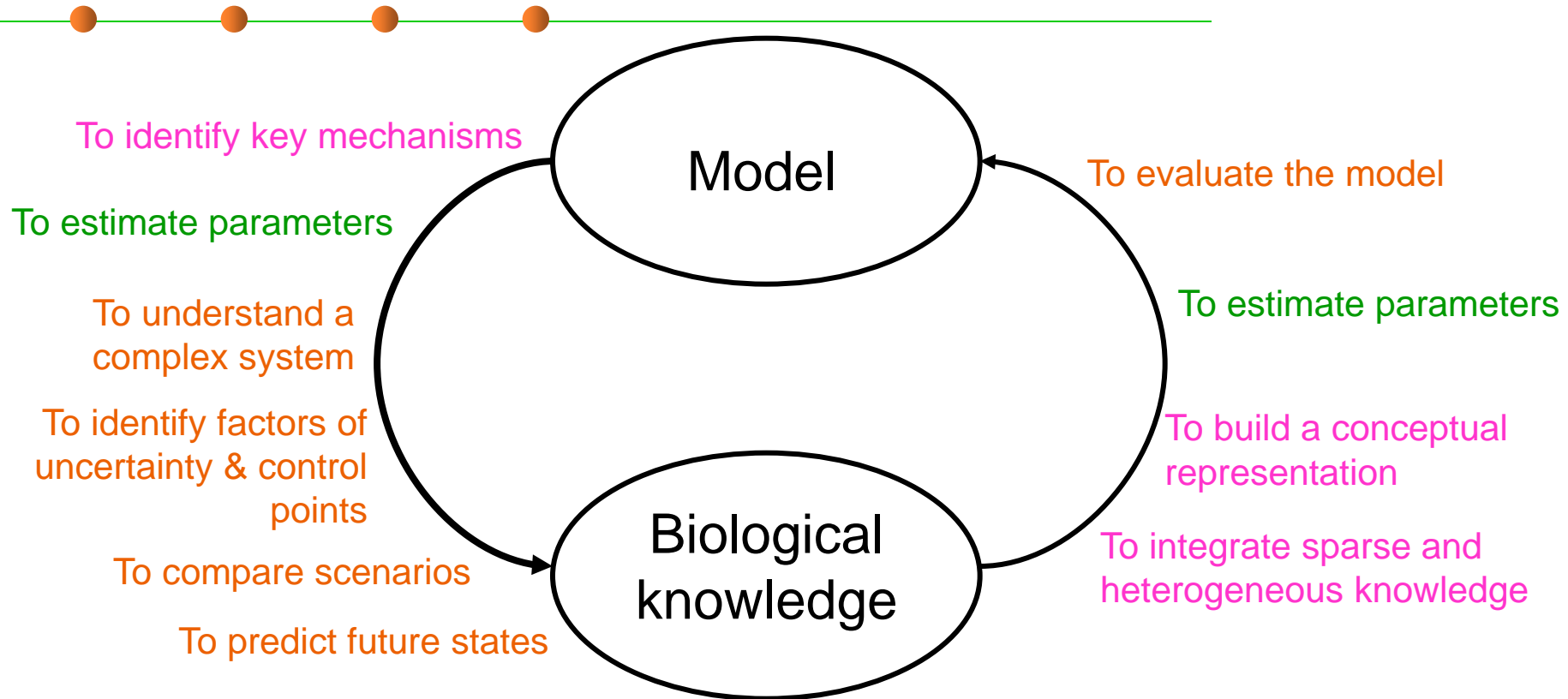
■ A complex biological system

- Interactions between hosts, pathogens, farming environment, farmers' decisions
 - Progression of the individual health status
 - Variability among susceptible / infected hosts
 - Variety of time and space scales
- Structured and managed host populations
 - In groups of animals within a herd
 - In herds localised in a region
 - In a supply chain of primary animal production

➔ **Integrative modelling approach**



An integrative modelling approach




**Which influent parameters?
Which influent assumptions?
How to control the system?**

**Which health states?
Which transitions?
Which assumptions?**

An integrative modelling approach

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-
-
-
- Integrated & simplified representation of a process
- To understand a biological system functioning & to predict its change over time / space
 - To predict numbers per health status in space and time
 - To estimate unobservable parameters
 - To evaluate biological assumptions
 - To test control scenarios

An integrative modelling approach

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- A compromise between **realism & parsimony**
 - Several **scales of integration** can be needed
 - Accounting for **farmers' decisions** improve the predictive capability of epidemiological models and of associated **decision tools**

An integrative modelling approach

■ Integration scales

■ Within-host scale

To represent interactions between the pathogen and the host immune system and the host shedding level

■ Between-host / within-herd scale

To account for host heterogeneity in susceptibility and shedding
To account for herd structure, management and farmer's decisions

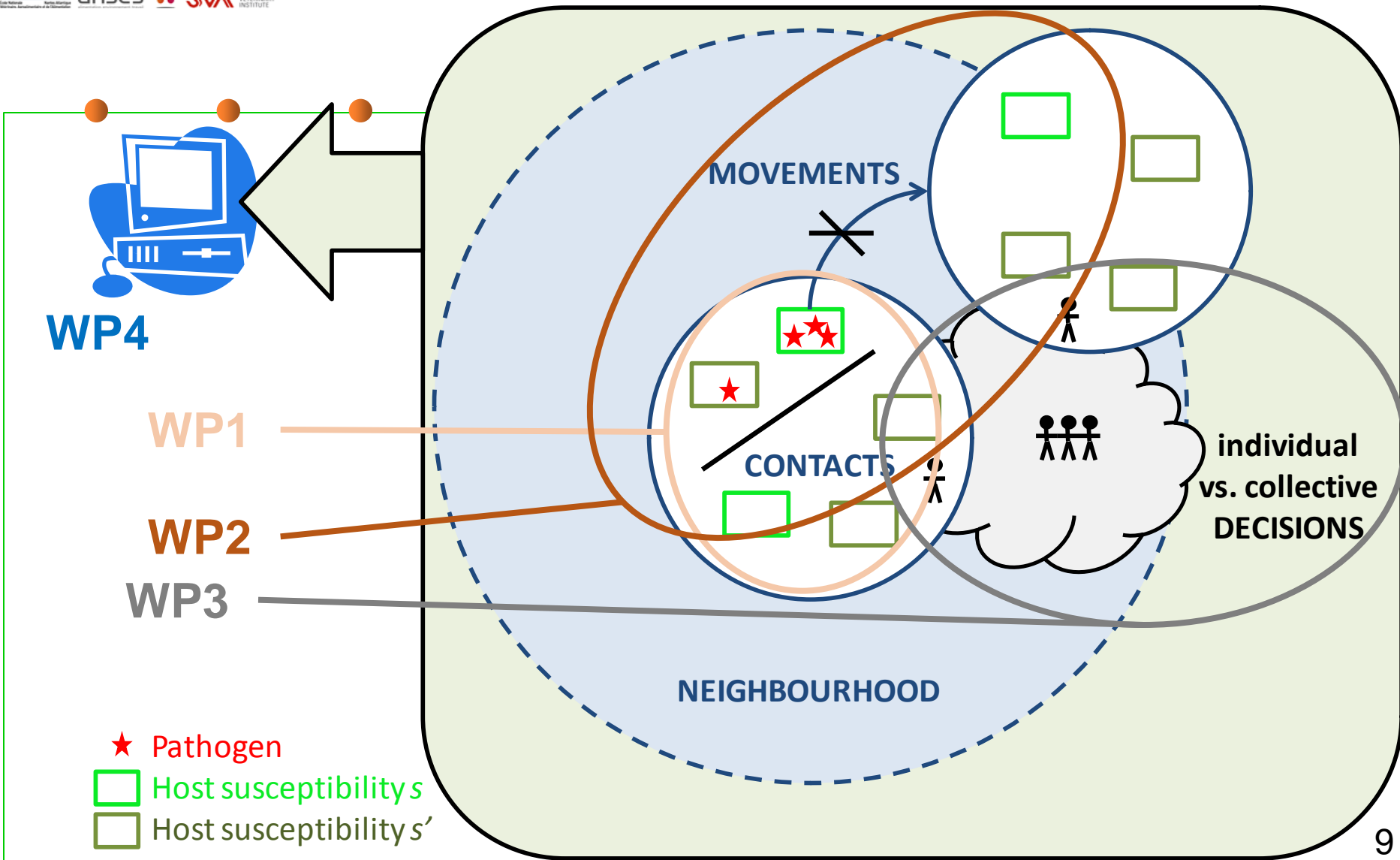
■ Between-herd / animal (meta)population scale

To represent animal movements, and neighbouring and indirect contacts between herds

To account for individual & collective farmers' decisions



multi-scale Modelling - from the animal Intra-Host to the Metapopulation - of pathogen spread to Evaluate control Strategies

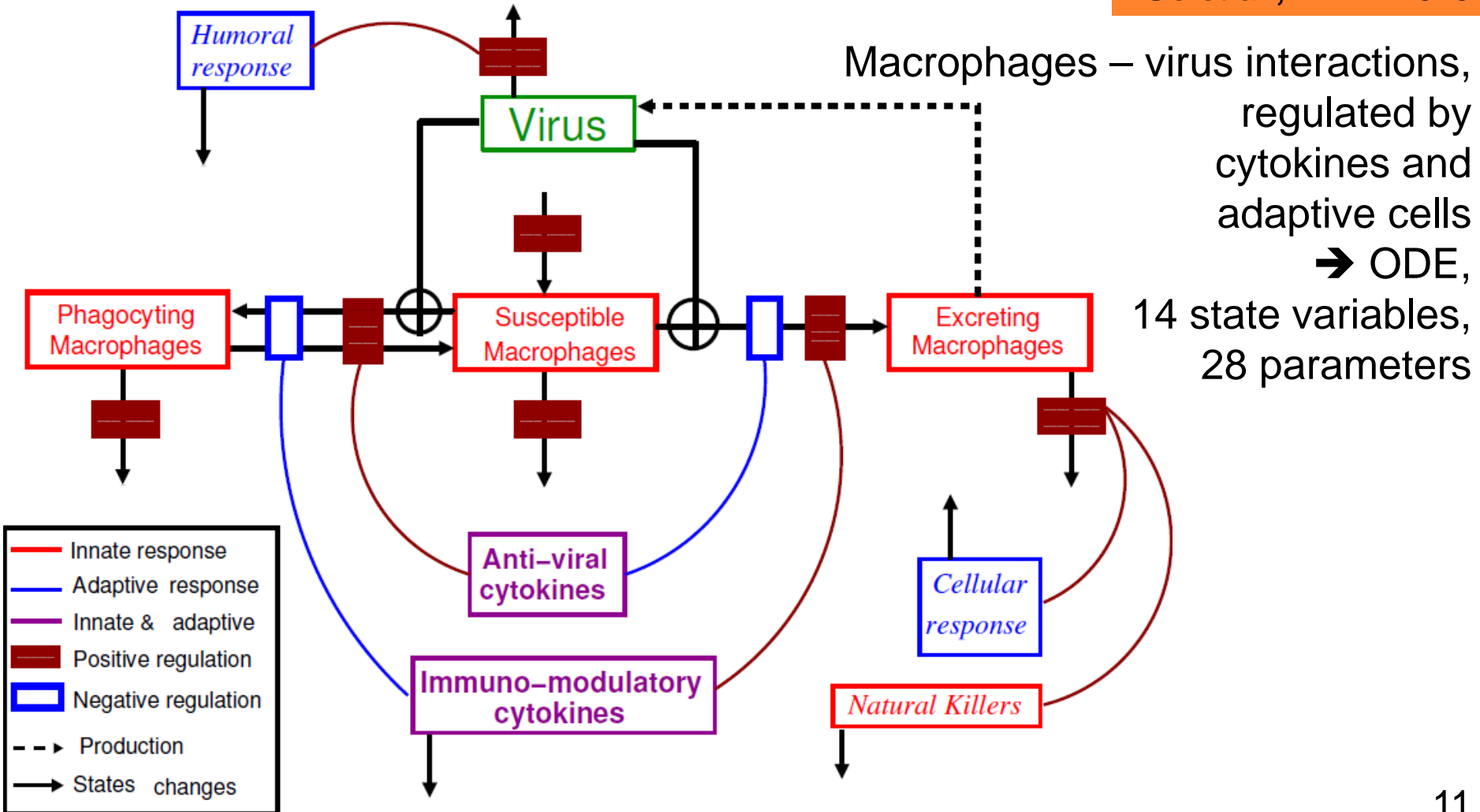


PRRSv infection dynamics within a pig

- Immune response to PRRSv infection poorly understood
 - ➔ control measures not efficient enough
 - ➔ major concern for the swine industry
- Aim: to identify the immune mechanisms determining host recovery considering the high variability of the immune response among pigs
- Method: a modeling approach

PRRSv infection dynamics in pig lung

Go et al., EAAP 2013



14 state variables,
28 parameters

PRRSv infection dynamics in pig lung

Go et al., EAAP 2013

Data: few experimental data, high variability and uncertainty.

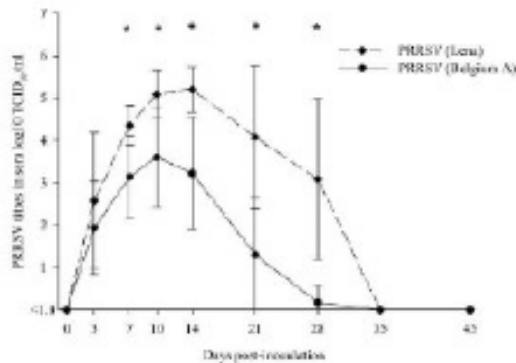
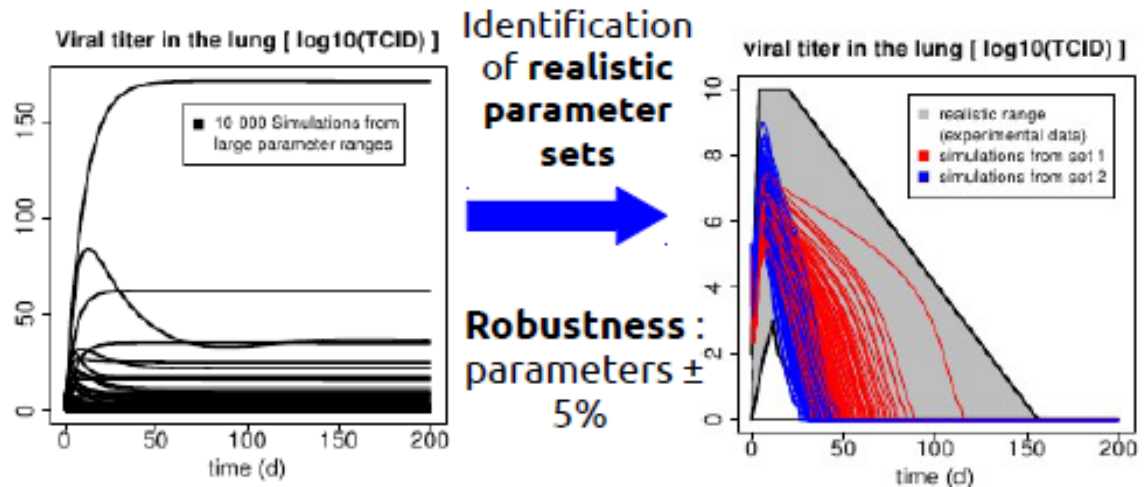


Figure 2 Virus titres in sera of pigs at different days post-inoculation with PRRSV (Lena) and PRRSV (Belgium A).

Parameter range estimation

Aim: Estimate parameter ranges resulting in simulations covering the response variability => **ad hoc method:**



PRRSv infection dynamics

■ Model extension

Lung model

- Contrasted viral dynamics
- Relationships between the immune mechanisms, infection duration, susceptibility & virulence



Pig model

A better representation of the adaptive response

Population model

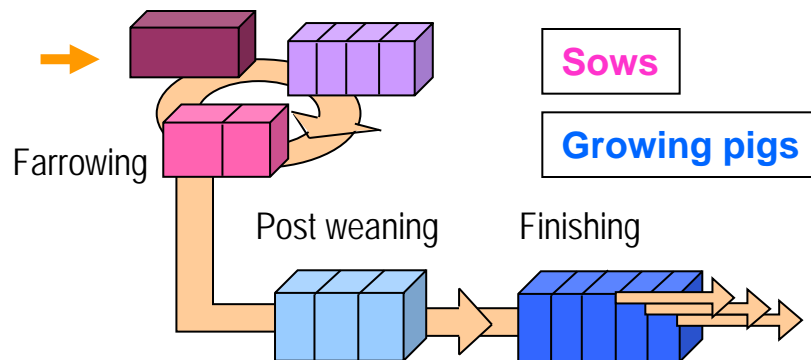
The individual model will be integrated in an immuno-epidemiological model

Pathogen infection dynamics in a herd

■ At the within-herd scale

■ Accounting for herd dynamics

- animal populations are structured
- Herds are heterogeneous in size, structure, composition, etc.



➔ Impact on infection spread, on potential target for control actions and on control efficiency

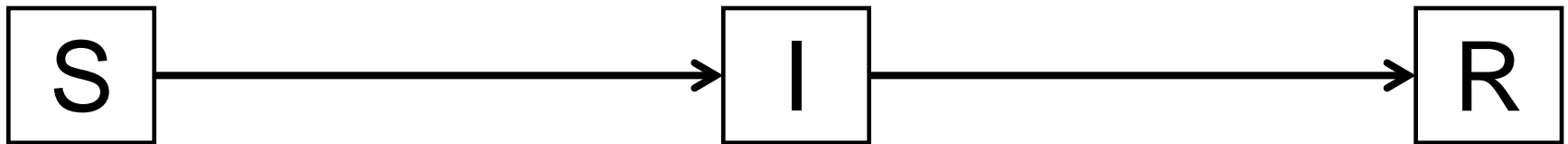
Pathogen infection dynamics in cattle

- At the within-herd scale

Courcoul et al., Vet. Res. 2011

- Q fever in a dairy herd

- Heterogeneity in shedding: level and routes (milk, mucus, faeces)



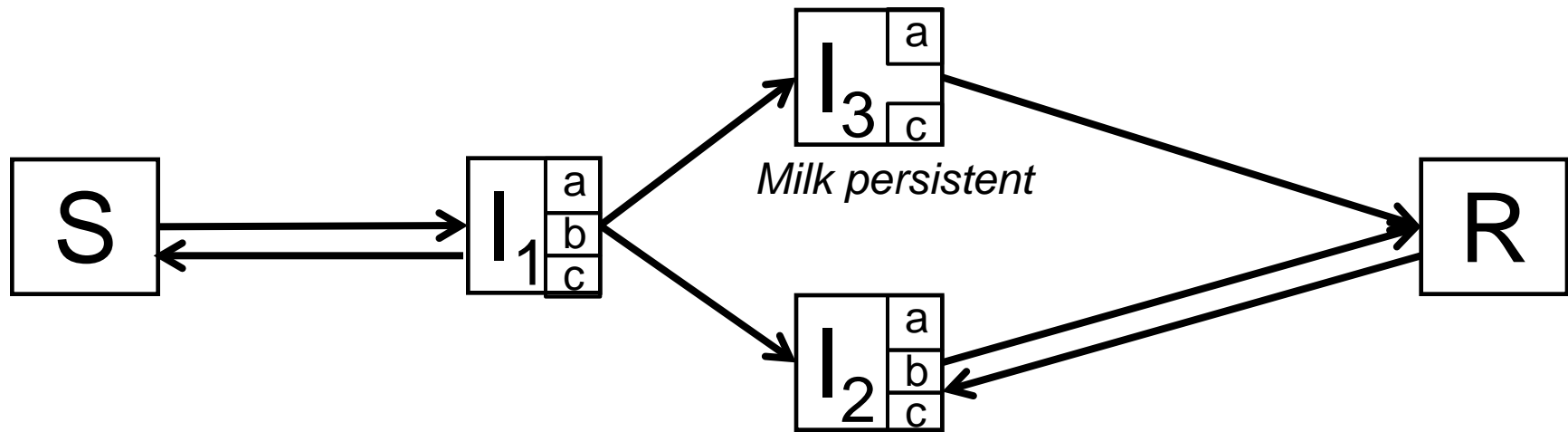
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Q fever in a dairy herd

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3 categories: milk only, mucus/faeces or both, milk + mucus/faeces or the 3 routes



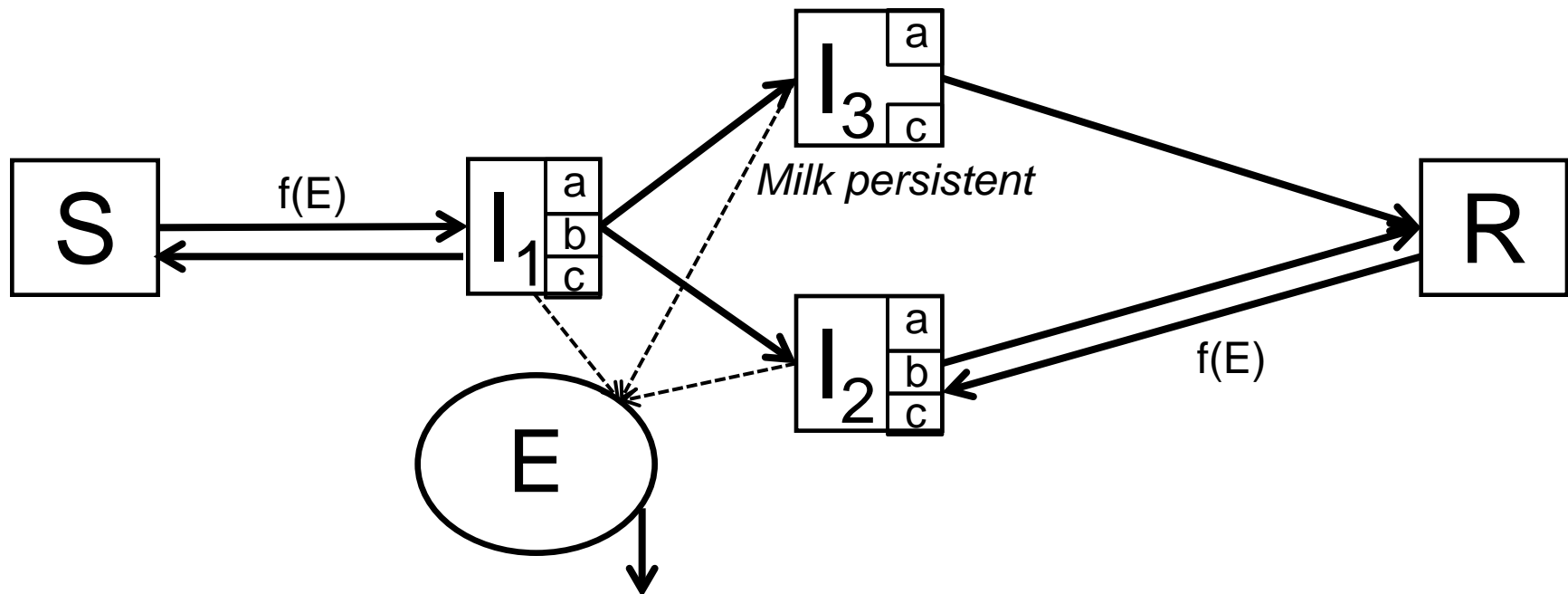
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Pathogen infection dynamics in cattle

Marcé et al., Vet. Res. 2011

At the within-herd scale

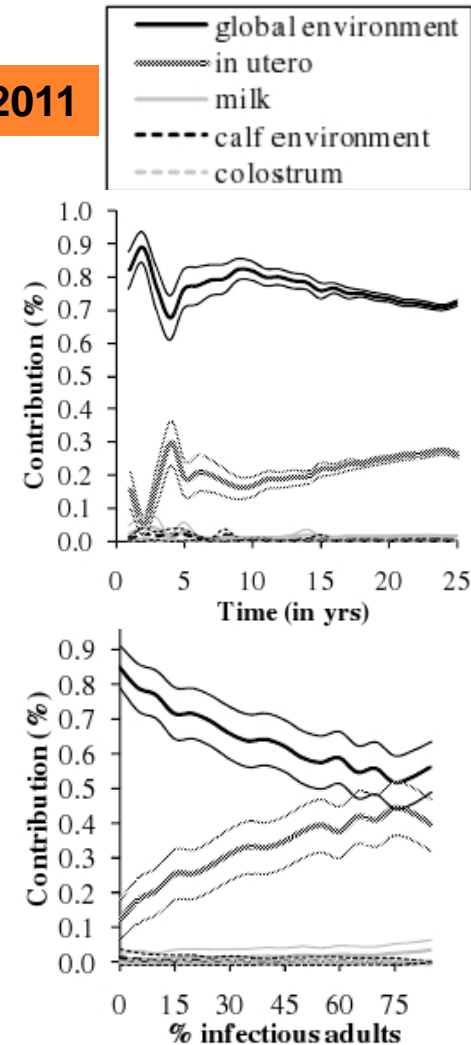
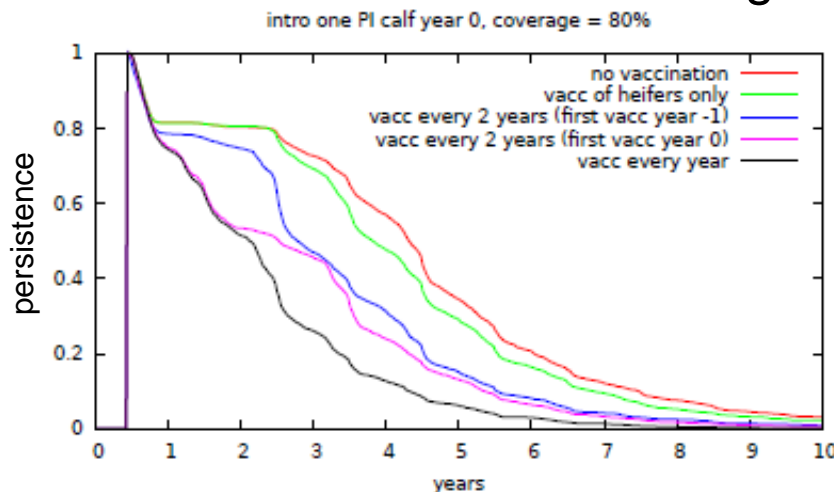
Paratuberculosis in dairy herd

- To understand the infection spread in a herd: relative contribution of transmission routes

BVD in beef herd

Damman et al., EAAP 2013

- To evaluate vaccination strategies



Pathogen infection dynamics in cattle

■ At the between-herd scale

■ Network of cattle movements between herds

Dutta et al., EAAP 2013

- Temporal and spatial variations in contacts
- Large range of herd types: size, production type
- Regional particularities: density, herd types, movement types

→ Coupling within-herd models through a contact network

- Animal movements
- Neighboring relationships

Summary



- Complementary skills to be mobilised
 - Epidemiology, immunology, infectiology, modelling, mathematics, computer sciences, health economics
 - To answer new research questions
 - To model multi-scale biological processes
 - To understand and formalised farmers' behaviours
- Viet et al., EAAP 2013**
- To guide management decision at different scales
 - After evaluating models' predictive capability

Thank you for listening!

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