



THE UNIVERSITY of EDINBURGH
Global Academy of
Agriculture and Food Systems

Modelling the AMR jump from farm animals to humans

Joao Gabriel Oliveira Marques

Dominic Moran

INTRODUCTION

1. AMU in humans vs. AMU in animals

- Primary focus is QALYs
 - Human health :: welfare
 - Goals:
 - Reduce hospital LOS (length of stay)
 - Reduce SD (sick days)
 - Control:
 - NHS
 - UK Health Security Agency
 - Department of Health and Social Care
- Primary focus is production
 - Animal health :: productivity
 - Goals:
 - Reduce financial losses
 - Reduce animal weight loss
 - Control:
 - Veterinary Medicines Directorate
 - Department for Environment, Food & Rural Affairs

1. AMU in humans vs. AMU in animals

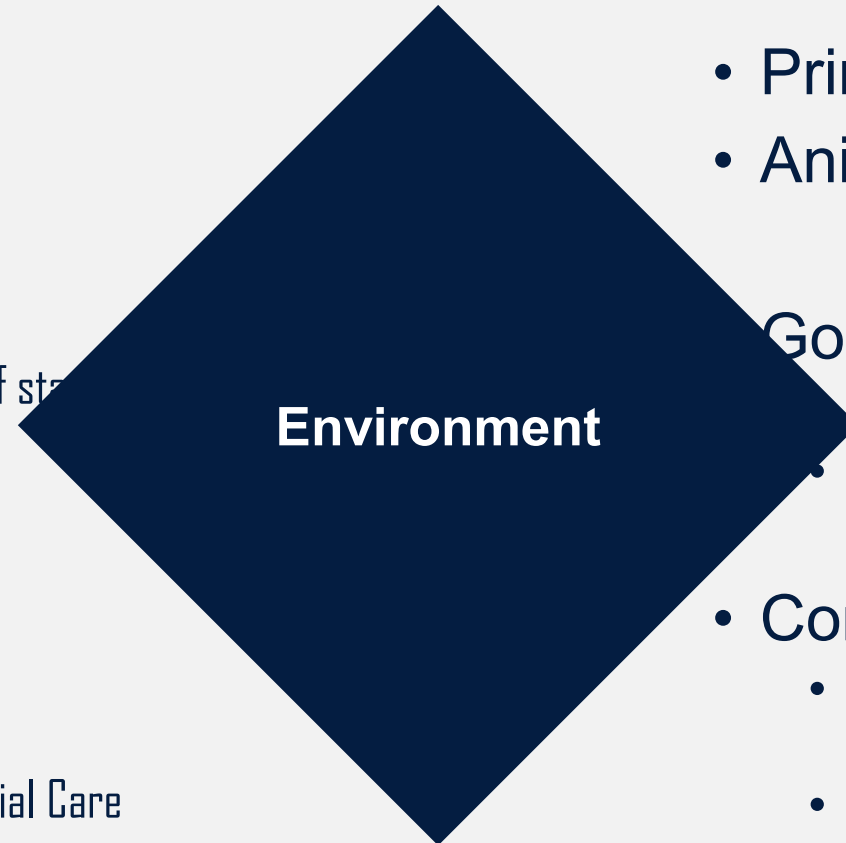
- Primary focus is QALYs
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- Goals:

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- NHS
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- Primary focus is production
- Animal health :: productivity

- Goals:

- Reduce financial losses
- Reduce animal weight loss

- Control:

- Veterinary Medicines Directorate
- Department for Environment, Food & Rural Affairs

2. Not a new a idea

- Goal: connect AMU in farm animals to AMR in humans
- Environment as a reservoir of bacteria (AMR and "Simple")

Lanzas 2011



Research article

Modelling the impact of curtailing antibiotic usage in food animals on antibiotic resistance in humans

B. A. D. van Bunnik  and M. E. J. Woolhouse

Van Bunnik 2017



3. What can be improved?

$$\frac{dR_H}{dt} = \Lambda_H(1 - R_H) + \beta_{HH}R_H(1 - R_H) + \beta_{AH}R_A(1 - R_H) - \mu_H R_H;$$

$$\frac{dR_A}{dt} = \Lambda_A(1 - R_A) + \beta_{AA}R_A(1 - R_A) + \beta_{HA}R_H(1 - R_A) - \mu_A R_A,$$

parameter	high impact
β_{AA}	0.1
β_{HH}	0.1
β_{AH}	0.1
β_{HA}	0.001
Λ_H	0.1
Λ_A	0.1
μ_A	0.1
μ_H	0.1

Van Bunnik 2017



3. What can be improved?

$$\frac{dR_H}{dt} = \Lambda_H(1 - R_H) + \beta_{HH}R_H(1 - R_H) + \beta_{AH}R_A(1 - R_H) - \mu_H R_H;$$

$$\frac{dR_A}{dt} = \Lambda_A(1 - R_A) + \beta_{AA}R_A(1 - R_A) + \beta_{HA}R_H(1 - R_A) - \mu_A R_A,$$

However, **the current state** of mathematical modelling of ABR **has both conceptual and empirical gaps**, which urgently need to be filled given the importance of having good models. Model results tell us that details matter [...]. However, without being able to routinely **inform and calibrate** these **models with** comprehensive **epidemiologic data**, we currently lack confidence in model predictions, most notably at the larger regional and national scale.

Knight 2019



Van Bunnik 2017

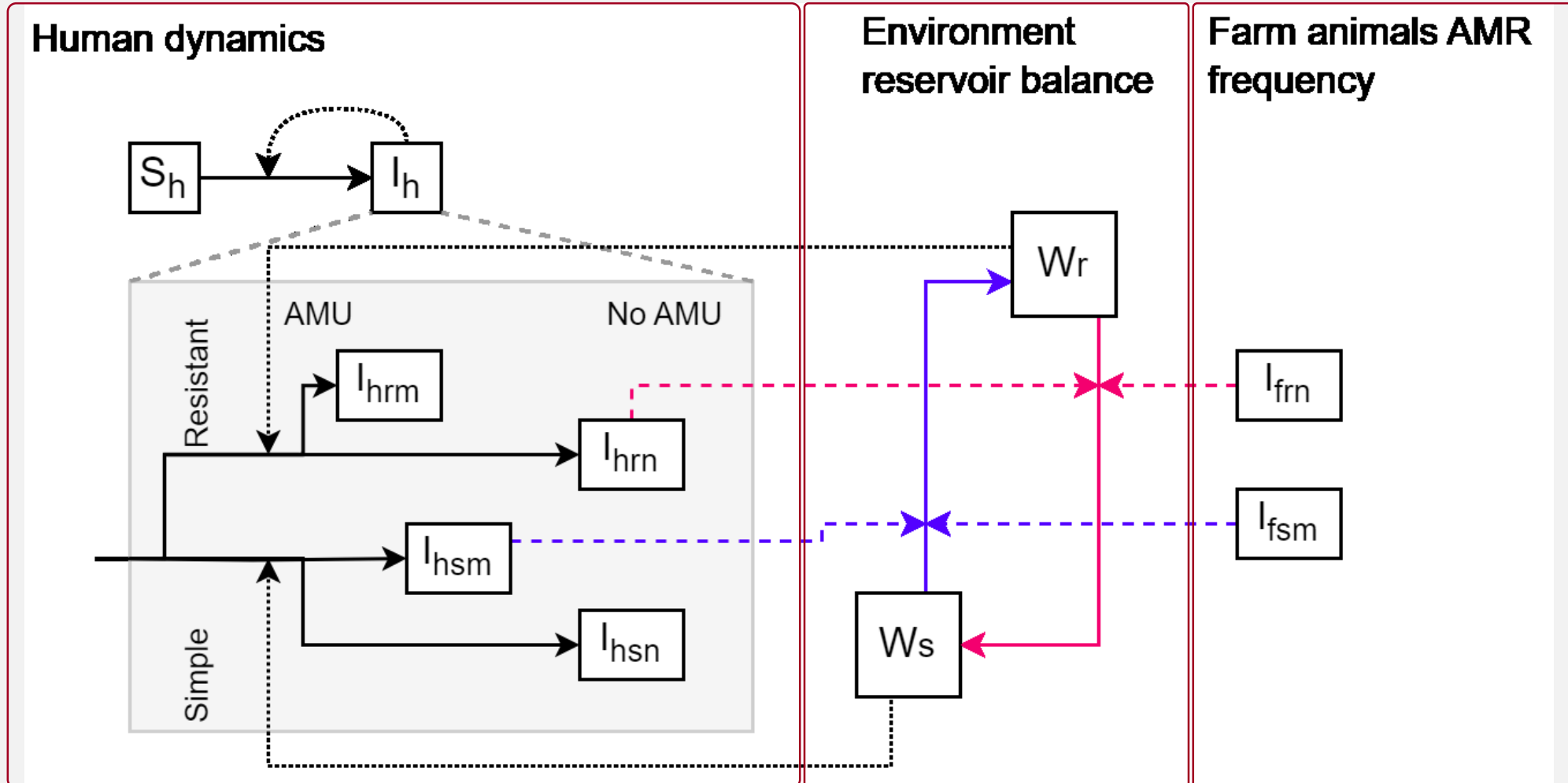


4. Narrowing the problem

- E. coli infection in humans
- Generalized AMR prevalence in farm animals
- Proxy by antimicrobials sales
- Indirect effect of farm animal on humans (environment reservoir)
- Environment reservoir "self regulates".

METHODOLOGY

5. Dynamic system



Quick recap 1

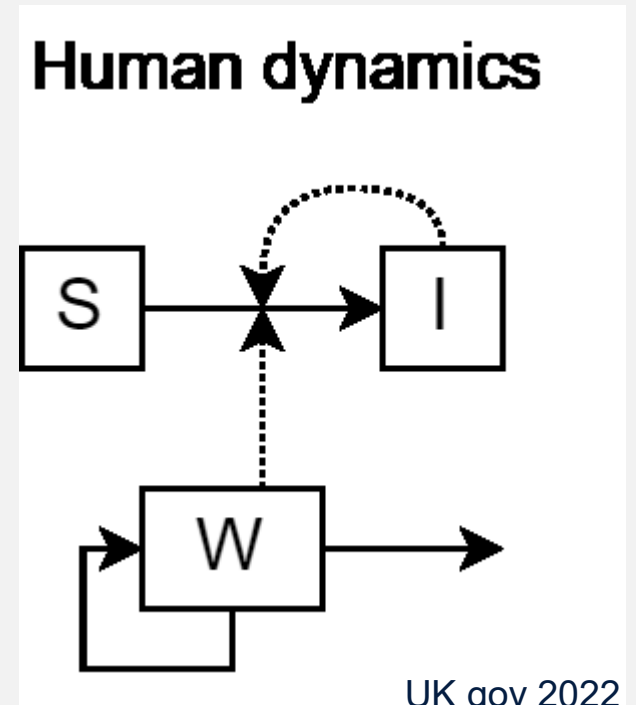
- S – susceptible proportion
- I – infected proportion
- $I = S+I$
- X – susceptible (total)
- Y – infected (total)
- $N = X+Y$

$$\frac{dX}{dt} \times \frac{1}{N} = \frac{dI}{dt}$$

5.1 Modelling AMR humans

- Data for England
- E. coli yearly infections
- Divided by hospital/community
- Sapronotic pathogen
- Estimation of:

$$\frac{dI_h}{dt}, \frac{dS_h}{dt}$$



Lanzas 2019



5.1 Modelling AMR humans

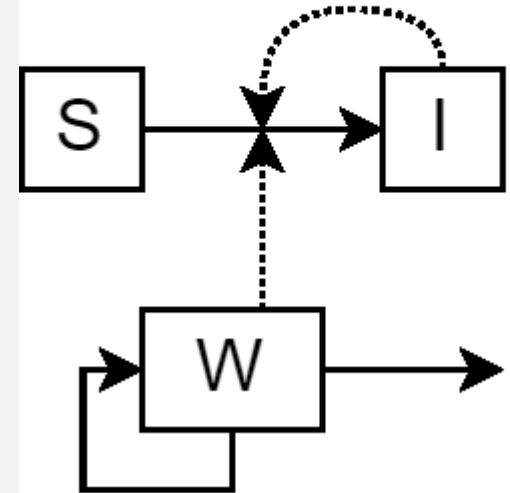
$$\frac{dI_h}{dt} = (1 - \alpha_{hW}) \frac{dI_{h\beta}}{dt} + \alpha_{hW} \frac{dI_{hW}}{dt} - \gamma_h I_h$$

$$\frac{dI_{h\beta}}{dt} = \beta IS$$

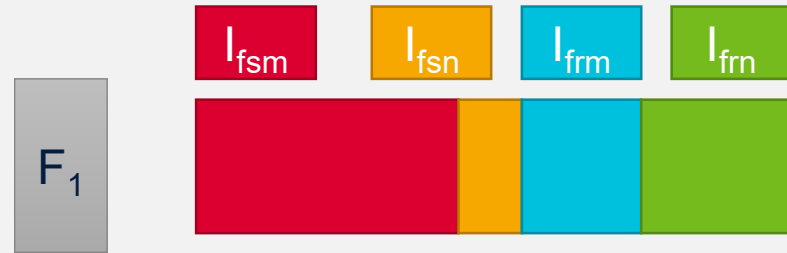
$$\frac{dI_{hW}}{dt} = \omega WS$$

$$\frac{dS_h}{dt} = \gamma_h I_h$$

Human dynamics



5.2 Modelling AMU on animals



$$I_{fsm} = \frac{Y_{fsm}}{Y_f}$$

$$Y_f = N_f \sum_j I_j$$

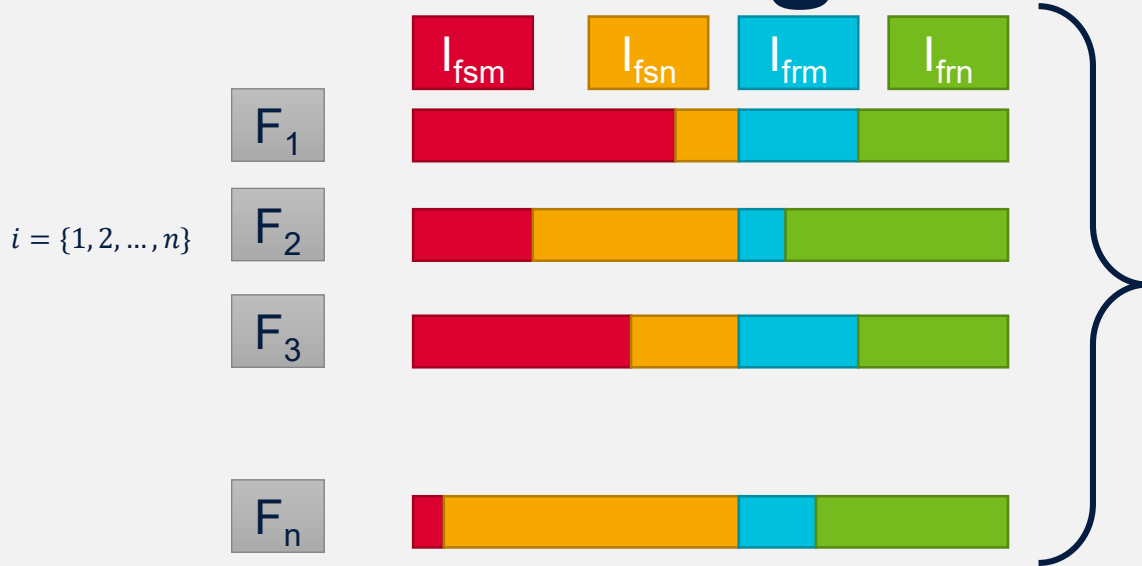
	Simple	Resistant	
AMU	I_{fsm}	I_{frm}	m
No AMU	I_{fsn}	I_{frn}	n
	s	r	

$j = \{fsm, fsn, frm, frn\}$

$$I_f = \frac{Y_f}{S_f}$$

- Typically density dependent model

5.2 Modelling AMU on animals



$$I_{i, fsm} = \frac{Y_{i, fsm}}{Y_i}$$

$$Y_i = N_i \sum_j I_{f, j}$$

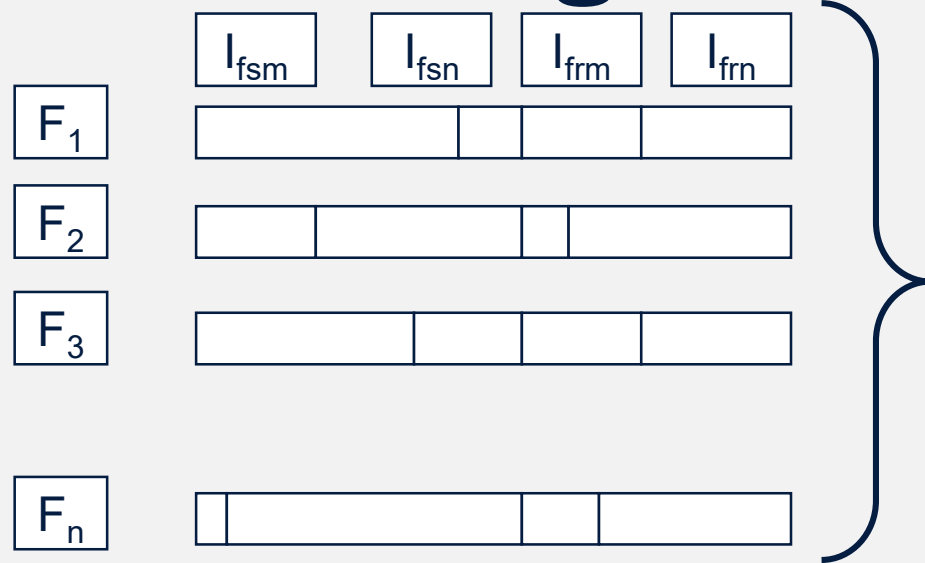
$$Y_f = \sum_i Y_i$$

$$I_f = \frac{Y_f}{S_f}$$

	Simple	Resistant	
AMU	I_{fsm}	I_{frn}	m
No AMU	I_{fsn}	I_{frn}	n
	s	r	

- Frequency dependent model

5.2 Modelling AMU on animals



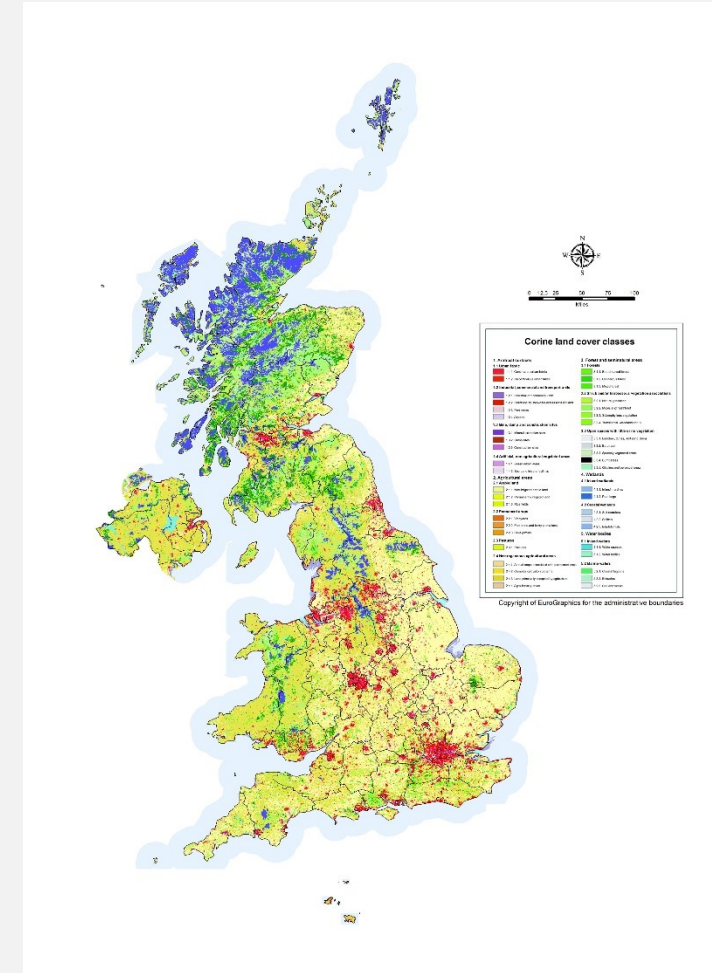
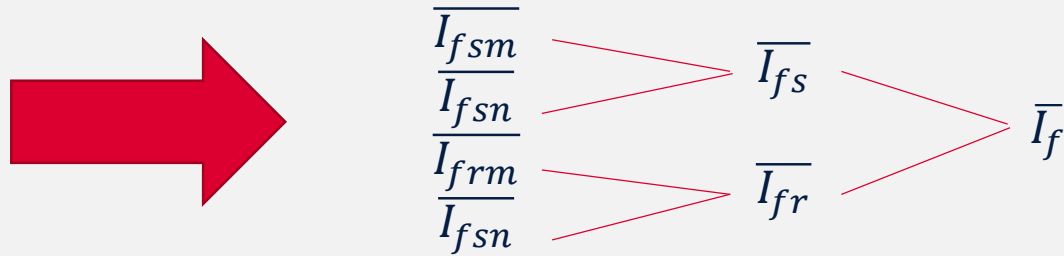
$$I_{i, fsm} = \frac{Y_{i, fsm}}{Y_i}$$

$$Y_i = N_i \sum_j I_{f, j}$$

$$Y_f = \sum_i Y_i$$

$$I_f = \frac{Y_f}{S_f}$$

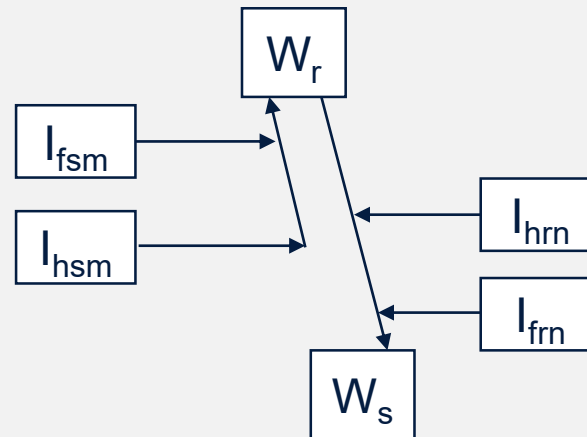
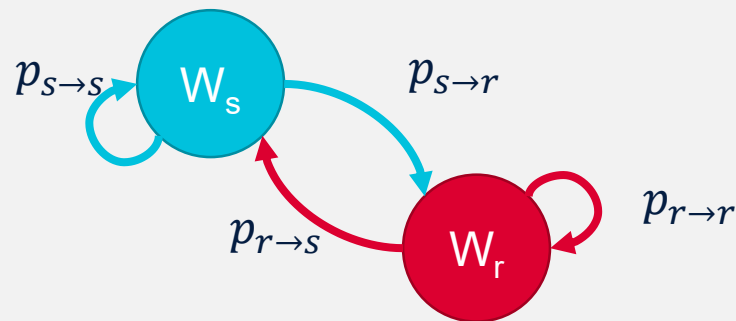
- Frequency dependent model



5.3 Modelling AMR environment

- Markov chain \equiv dynamic system

	From W_s	From W_r
To W_s	$p_{s \rightarrow s}$	$p_{r \rightarrow s}$
To W_r	$p_{s \rightarrow r}$	$p_{r \rightarrow r}$



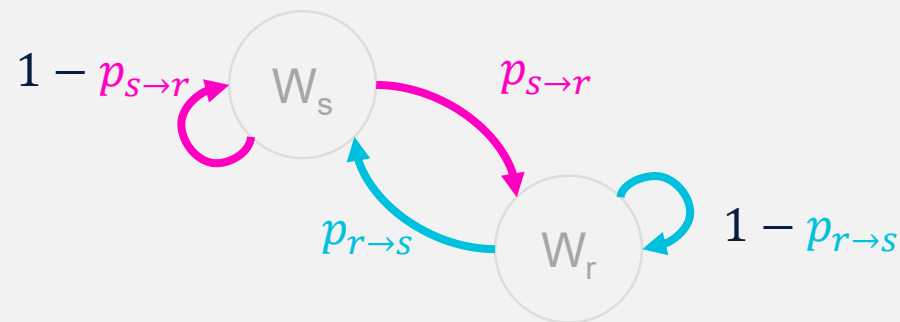
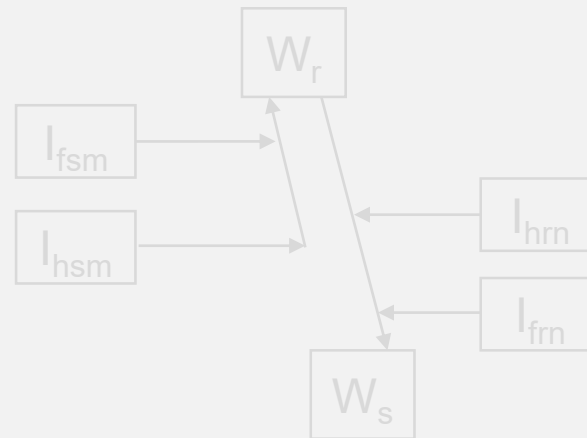
$$\frac{dW_r}{dt} = \alpha_{Wh}(\lambda_{hsm}I_{hsm} - \lambda_{hrn}I_{hrn}) + \alpha_{Wf}(\lambda_{fsm}I_{fsm} - \lambda_{frn}I_{frn}) - \tau W_r$$

$$\frac{dW_s}{dt} = \alpha_h(-\lambda_{hsm}I_{hsm} + \lambda_{hrn}I_{hrn}) + \alpha_f(-\lambda_{fsm}I_{fsm} + \lambda_{frn}I_{frn}) + \tau W_r$$

5.3 Modelling AMR environment

- Markov chain \equiv dynamic system

	From W_s	From W_r
To W_s	$p_{s \rightarrow s}$	$p_{r \rightarrow s}$
To W_r	$p_{s \rightarrow r}$	$p_{r \rightarrow r}$



$$\frac{dW_r}{dt} = \alpha_{Wh}(\lambda_{hsm}I_{hsm} - \lambda_{hrn}I_{hrn}) + \alpha_{Wf}(\lambda_{fsm}I_{fsm} - \lambda_{frn}I_{frn}) - \tau W_r$$

$$p_{r \rightarrow s} = \frac{\alpha_{Wh}\lambda_{hrn}I_{hrn} + \alpha_{Wf}\lambda_{frn}I_{frn} + \tau W_r}{W_r}$$

$$p_{s \rightarrow r} = \frac{\alpha_{Wh}\lambda_{hsm}I_{hsm} + \alpha_{Wf}\lambda_{fsm}I_{fsm} + \tau W_s}{W_s}$$

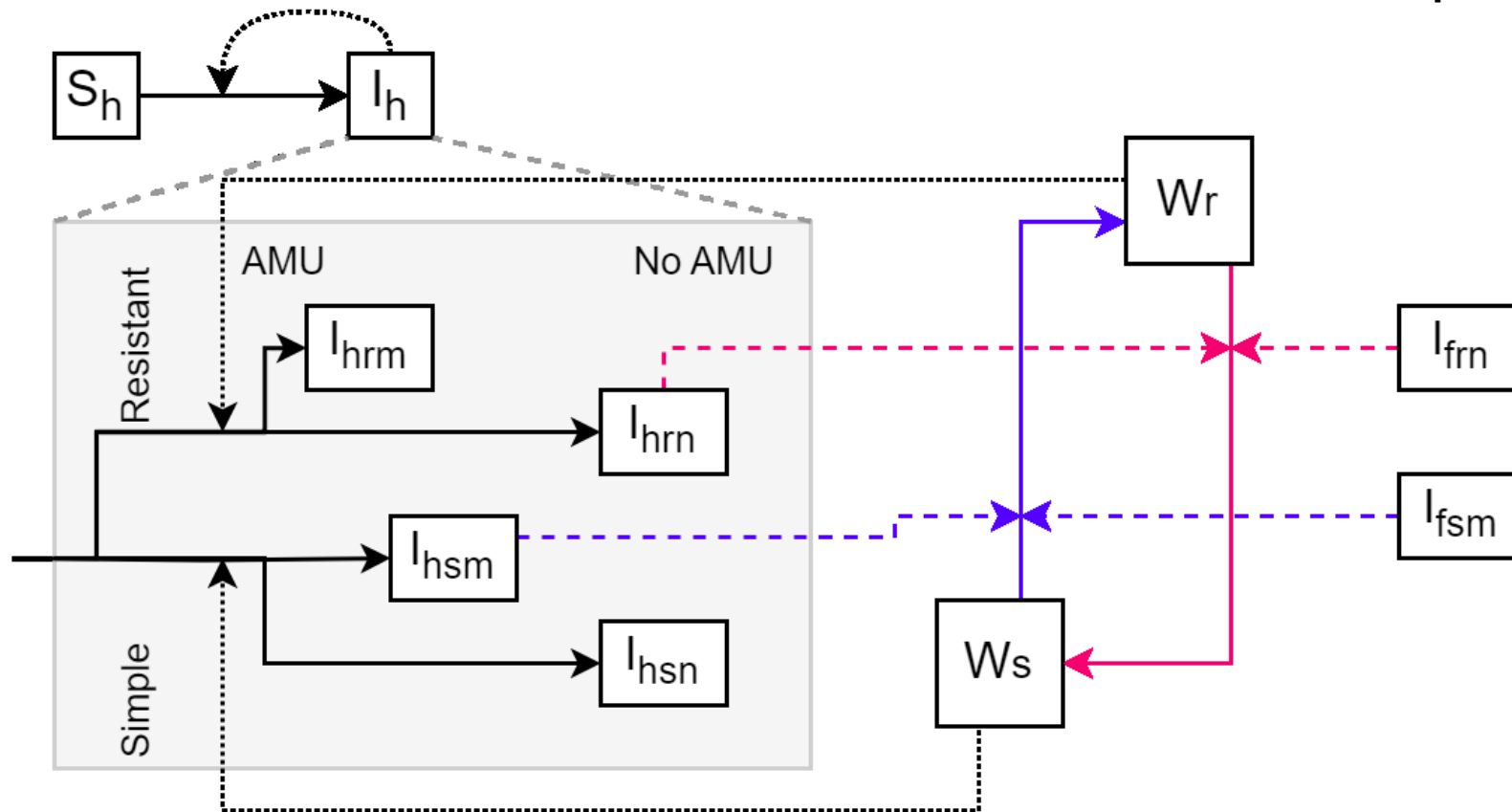
$$\frac{dW_s}{dt} = \alpha_h(-\lambda_{hsm}I_{hsm} + \lambda_{hrn}I_{hrn}) + \alpha_f(-\lambda_{fsm}I_{fsm} + \lambda_{frn}I_{frn}) + \tau W_r$$

6. Dynamic system

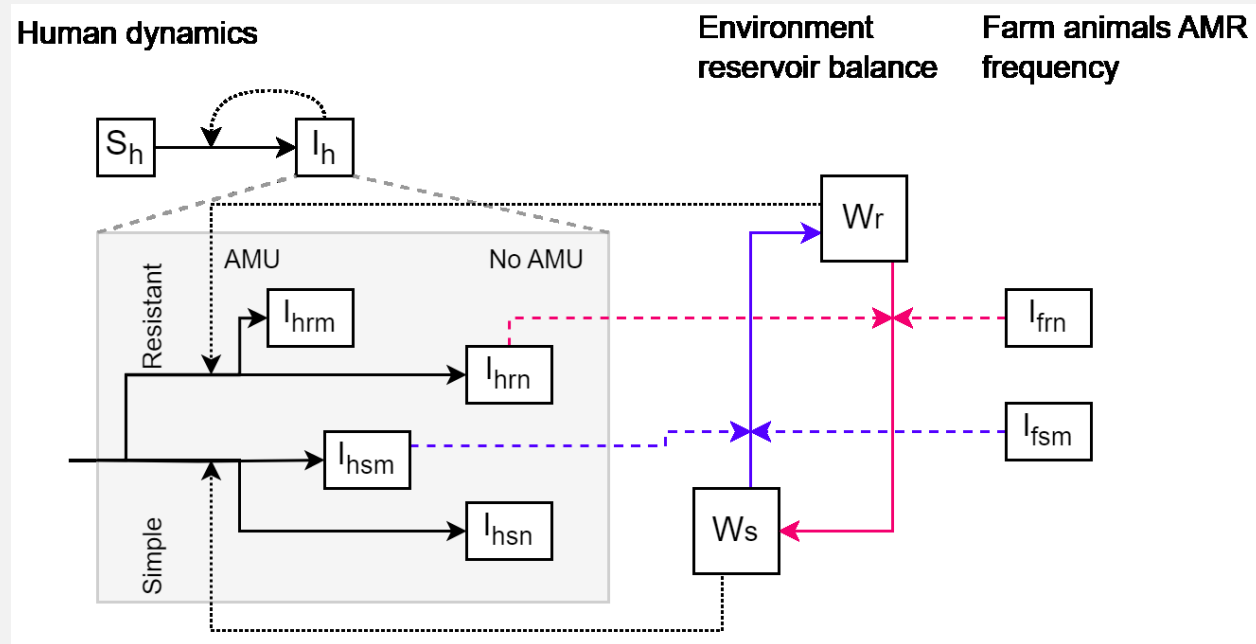
Human dynamics

Environment
reservoir balance

Farm animals AMR
frequency



6. Dynamic system



Data values / calibrated equilibrium:

- Human: S_h, I_h, β, ω

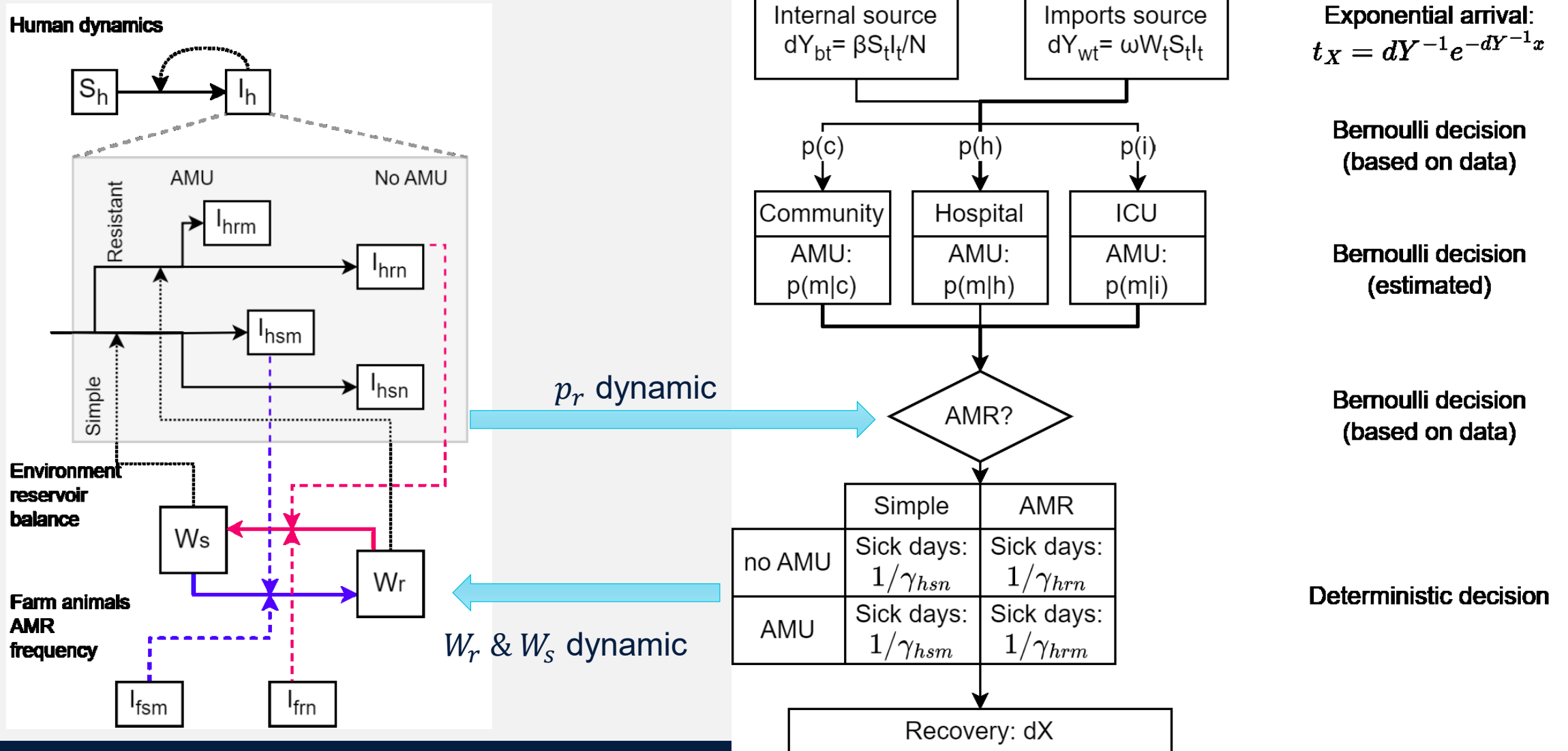
Estimated:

- Environment: $W_s, W_r, \alpha_{Wh}, \alpha_{Wf}, \lambda$
- Human: α_w
- Farm: I_f, I_{frm}, \dots

Output:

- Environment: W_s, W_r
- Human: S_h, I_h

7. Discrete event simulation



PRELIMINARY RESULTS



Sassi 2006

Quick recap 2

- DALY and QALY:

$$D = 1 - Q$$

- QALYs gained:

$$Q_g = Q_i \frac{1 - e^{-rL_i}}{r} - Q \frac{1 - e^{-rL}}{r}$$

L: duration of the disease (i : with treatment)

Q: health-related quality of life weights predicted (or observed)

r: discount rate

- DALYs from infection:

$$D_g = \frac{D}{r} (1 - e^{-rL_i})$$

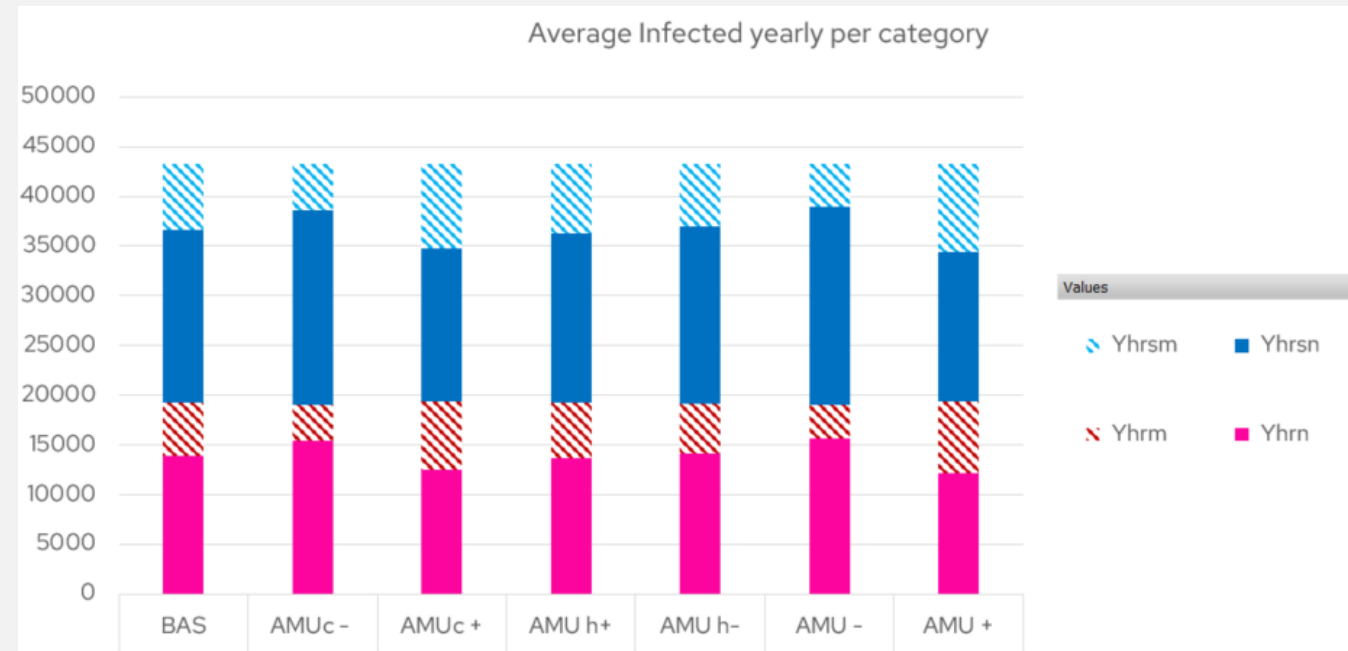
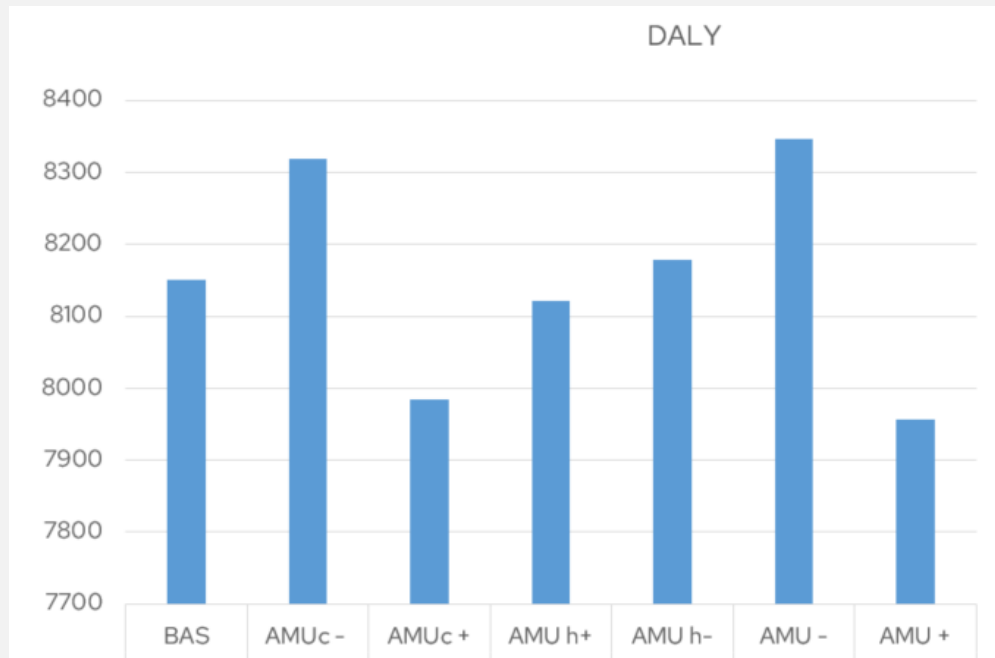
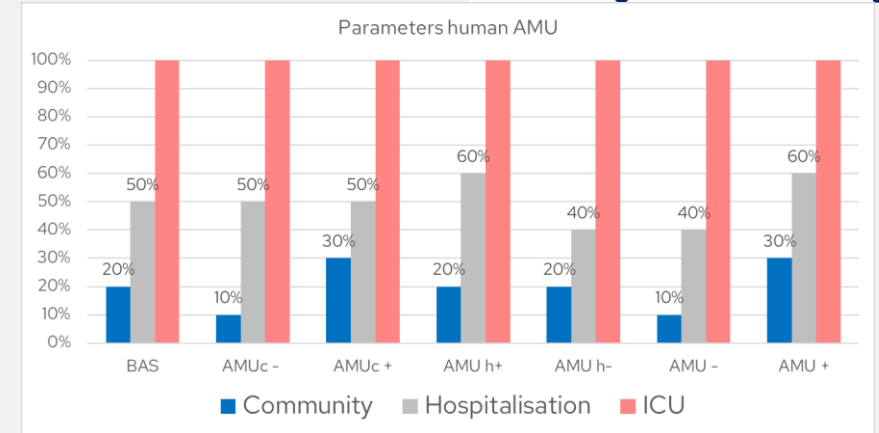
- Infectious disease, acute episode moderate $D = 0.051$
- $r = 3\%$

Haagsma 2015



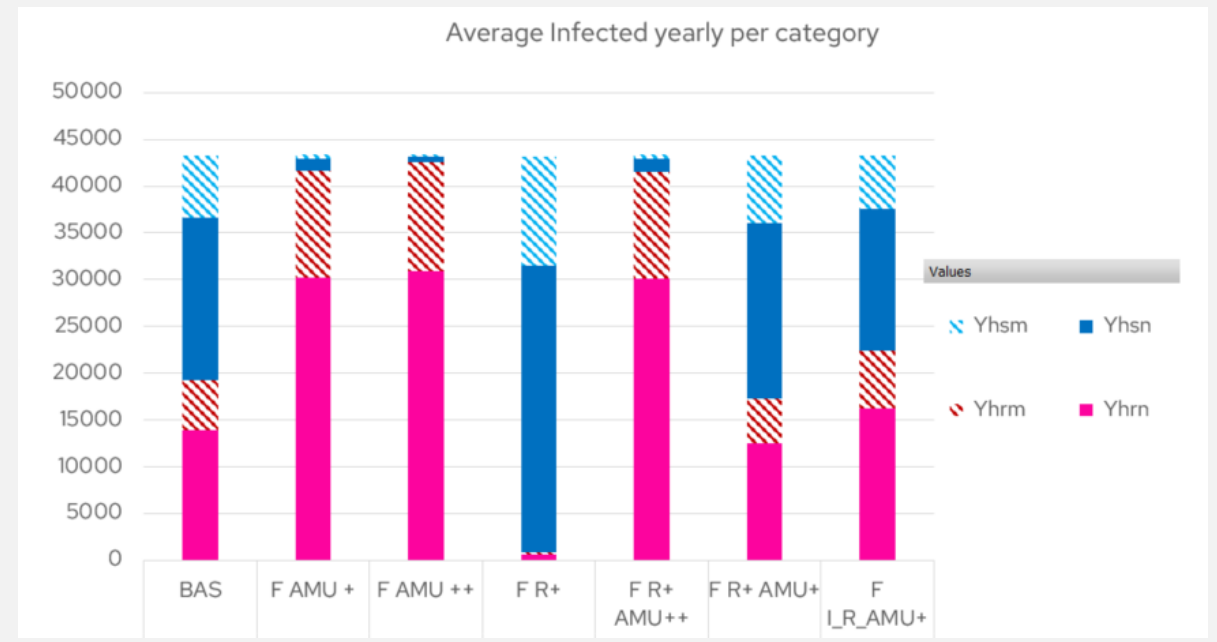
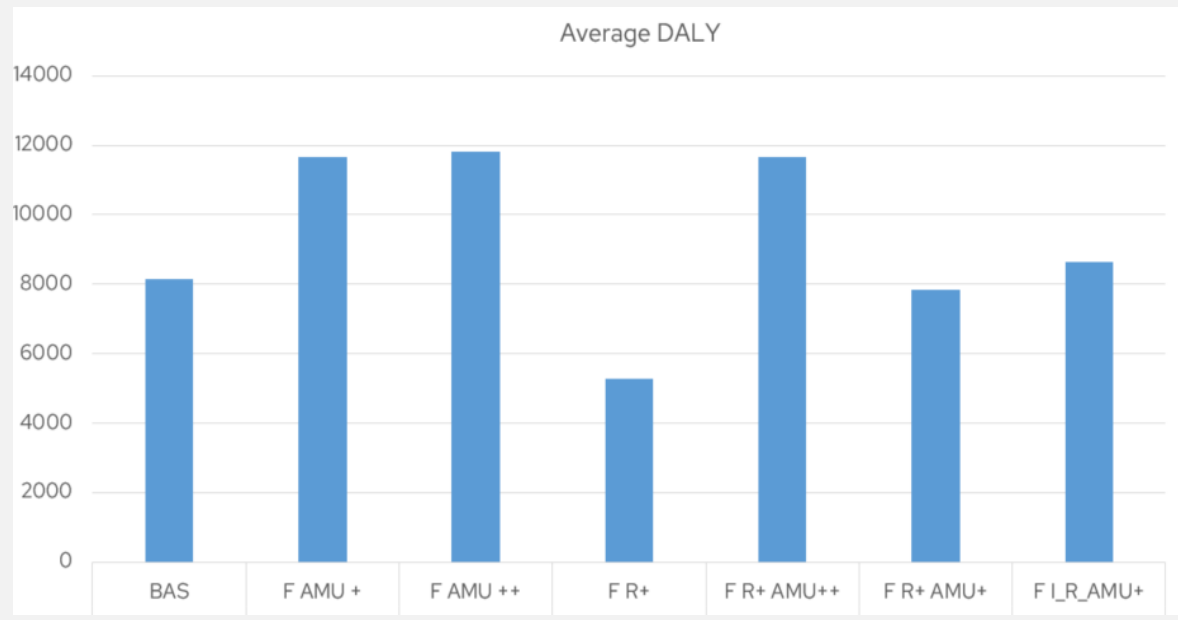
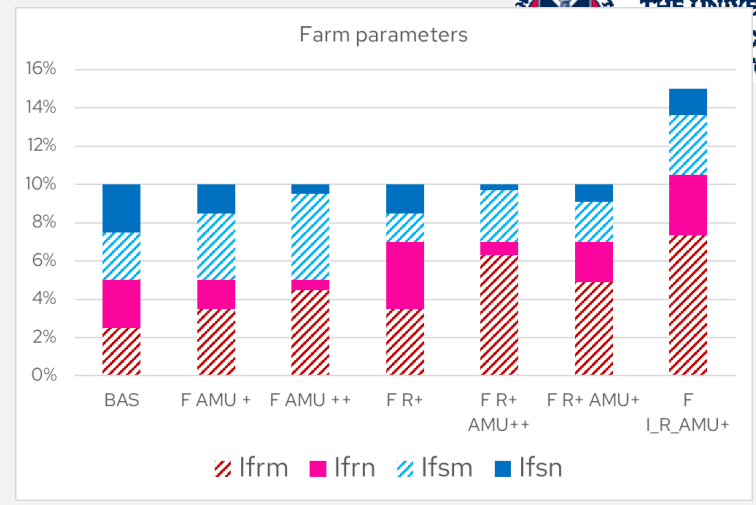
8.1 Scenarios Human

- Changing AMU prescription policy



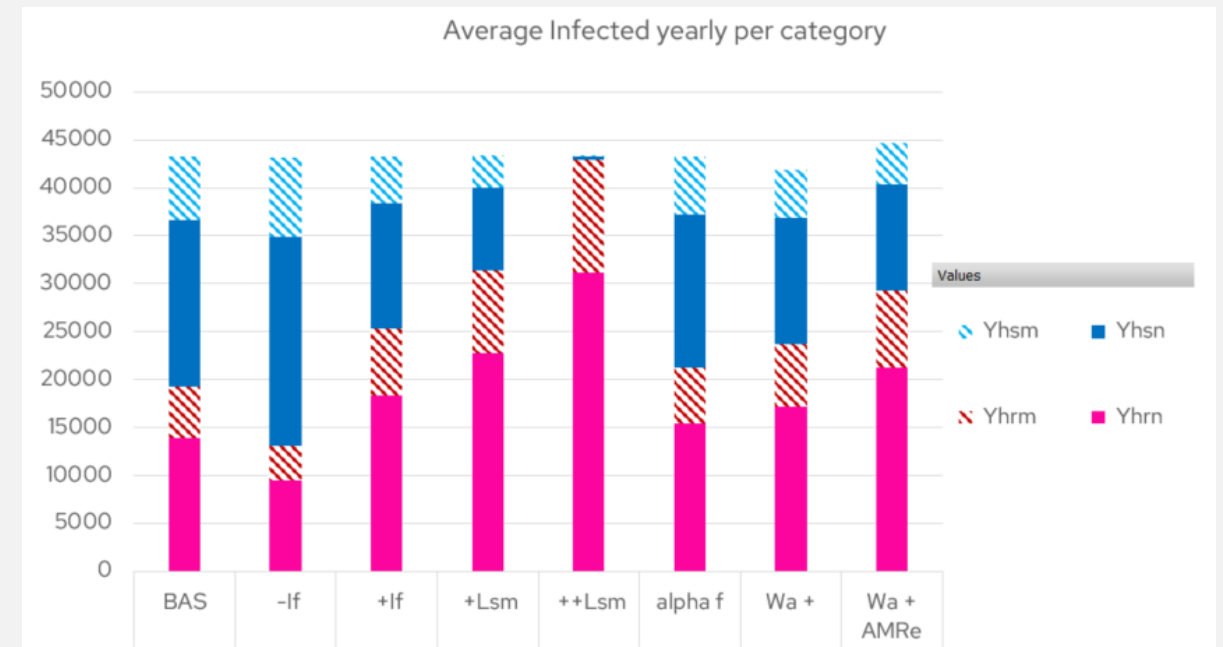
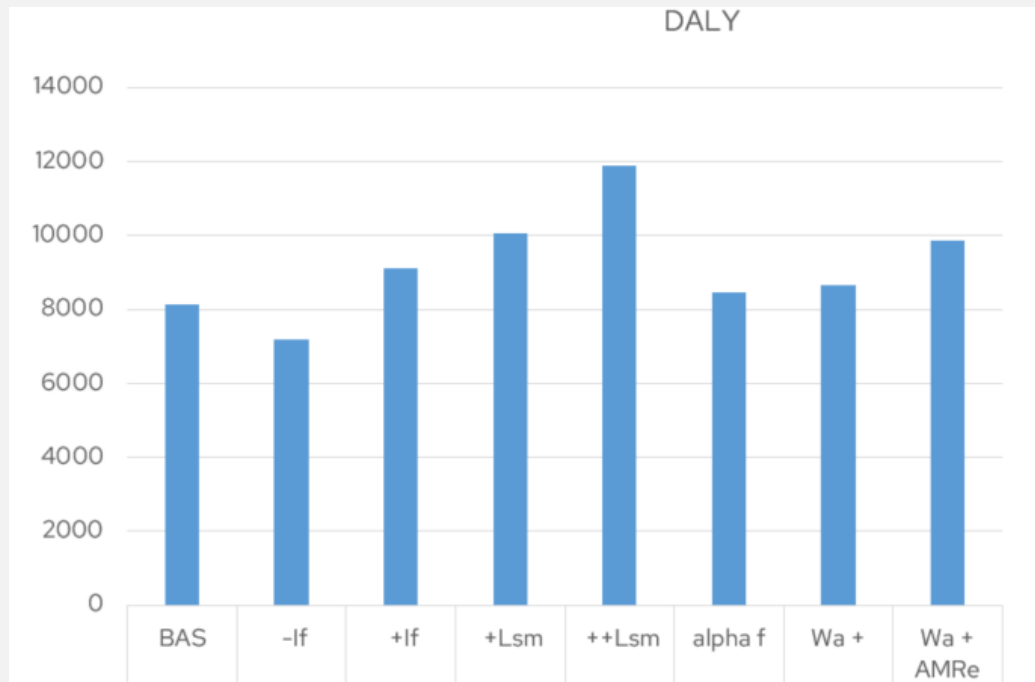
8.2 Scenarios Farm

- Farm parameters analysis



8.3 Scenarios Meta

Scenario Name	If	Lhsm	Lfsm	alphaW	Wa0	Ws0	alpha_fw	alpha_hw
BAS	0.1	0.6	0.6	0.4	0.4	0.6	0.01	0.99
-If	0.05	0.6	0.6	0.4	0.4	0.6	0.01	0.99
+If	0.15	0.6	0.6	0.4	0.4	0.6	0.01	0.99
+Lsm	0.1	0.7	0.7	0.4	0.4	0.6	0.01	0.99
++Lsm	0.1	0.9	0.9	0.4	0.4	0.6	0.05	0.95
alpha f	0.1	0.6	0.6	0.4	0.4	0.6	0.05	0.95
Wa +	0.1	0.6	0.6	0.4	0.45	0.55	0.01	0.99
Wa + AMRe	0.1	0.6	0.6	0.5	0.45	0.55	0.01	0.99



9. Next steps

- Calibrate different years will produce information on
 - W_s and W_r dynamics
 - α_{Wh} and α_{Wf} : effects of human and farm animals in the environmental equilibrium
 - λ : effect of each infected proportion in the environmental equilibrium
 - α_{hW} : proportion of human infections coming from the environment



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Thank you!

gabriel.marques@ed.ac.uk