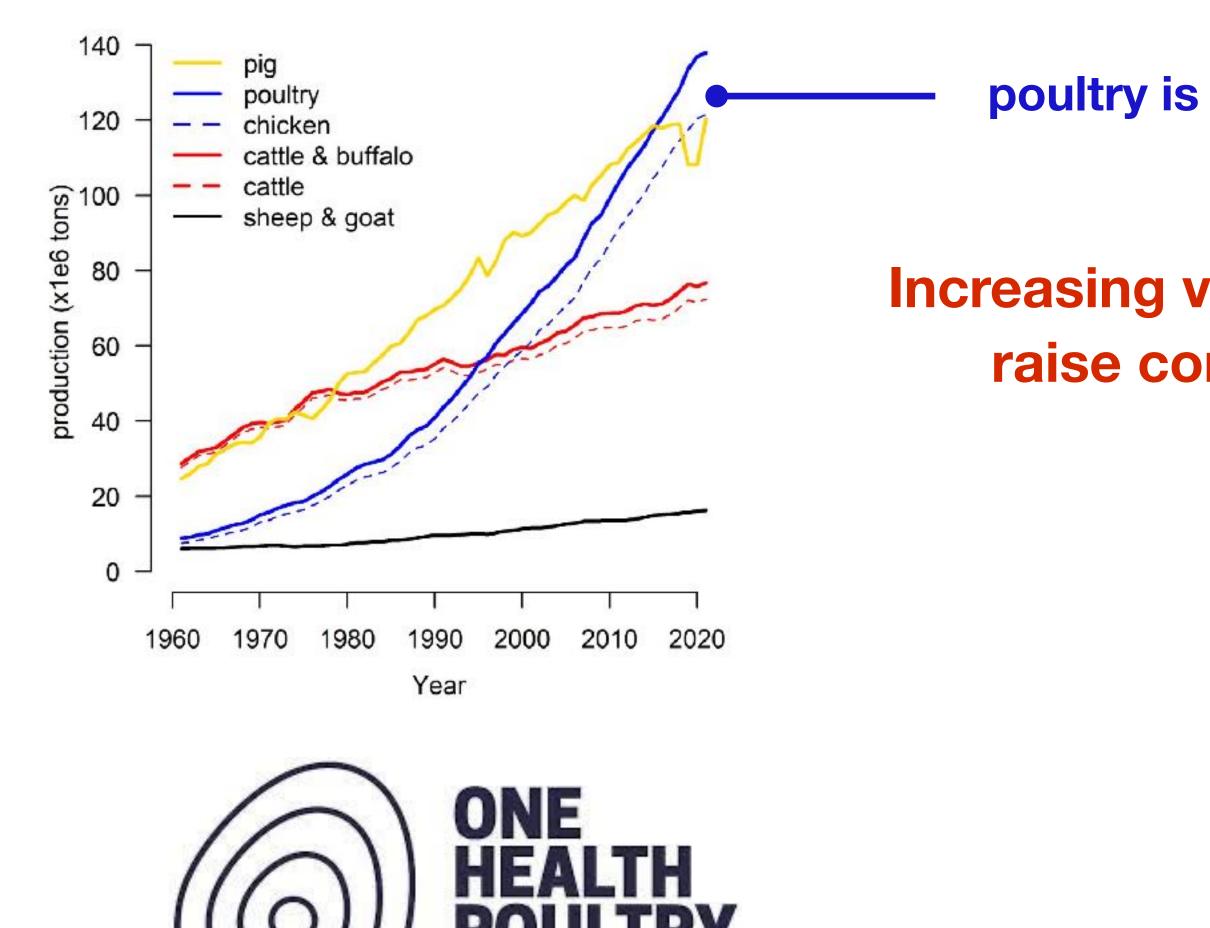
Modelling avian influenza transmission in poultry production and distribution networks in Bangladesh

Francesco Pinotti, ModStatSAP, 16/05/2024





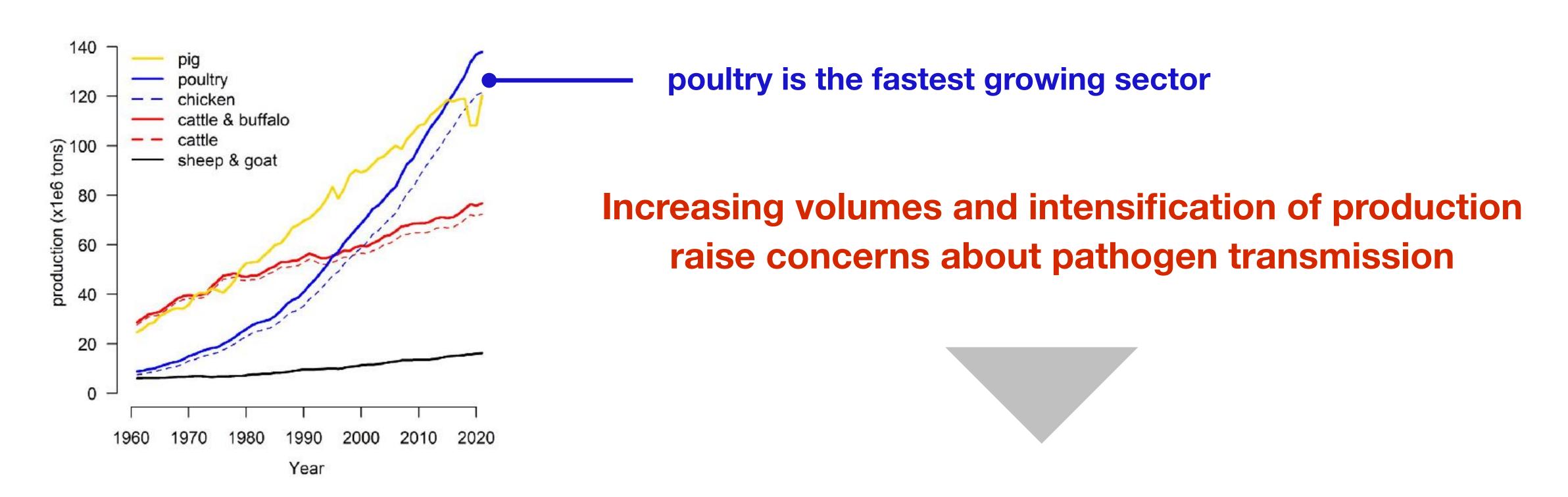
Livestock production is increasing globally



poultry is the fastest growing sector

Increasing volumes and intensification of production raise concerns about pathogen transmission

Livestock production is increasing globally



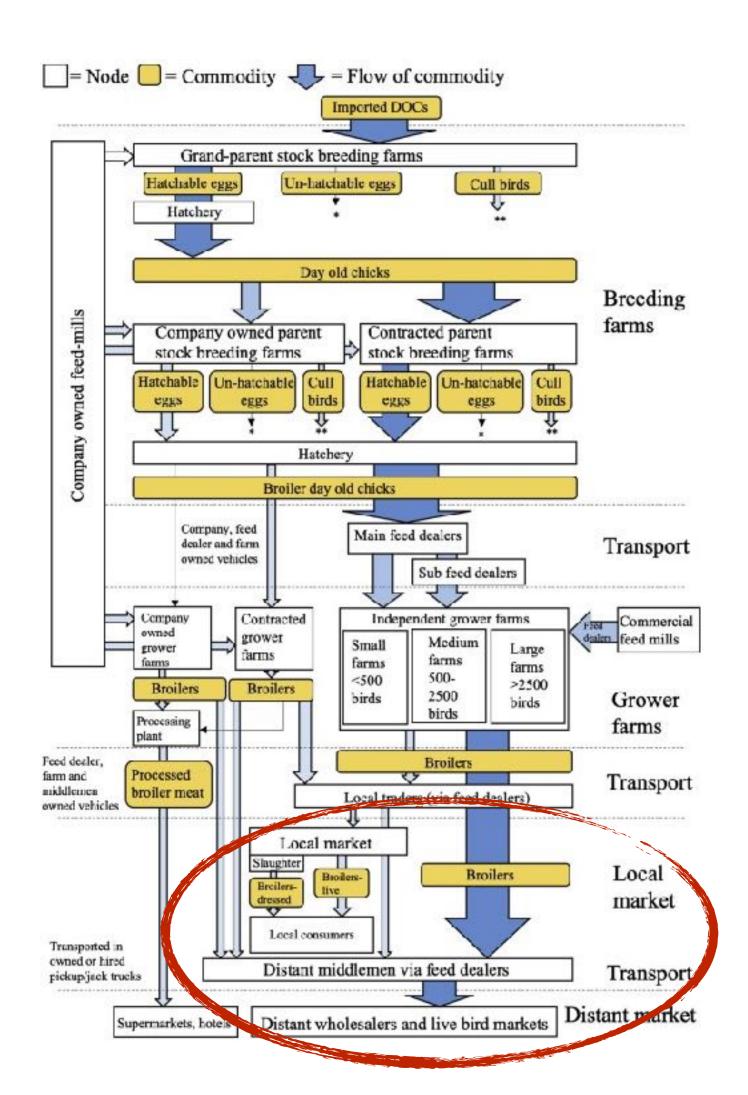


Characterise the networks through which chickens are produced and distributed (PDNs)

Poultry production and distribution in Bangladesh

Poultry moves along complex and heterogeneous networks before reaching consumers

(LBMs)



of birds

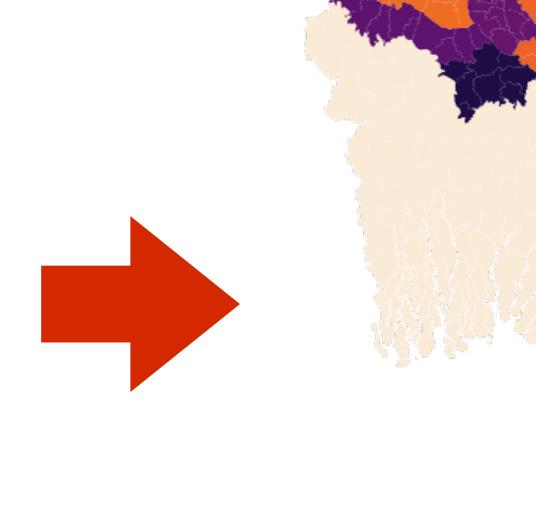


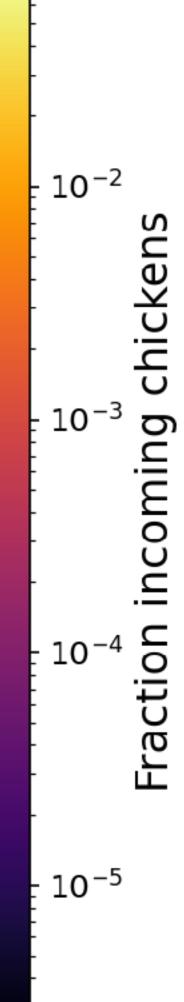
Live Bird Markets in Bangladesh

LBMs sustain AIV transmission

- High (H9N2 AIV) viral prevalence
 Negovetich et al, *PLOS ONE*, 2011; Kim et al, *Emerging Infect Dis*, 2018
- High mixing of birds and viruses
 Moyen et al, *Sci Rep*, 2021

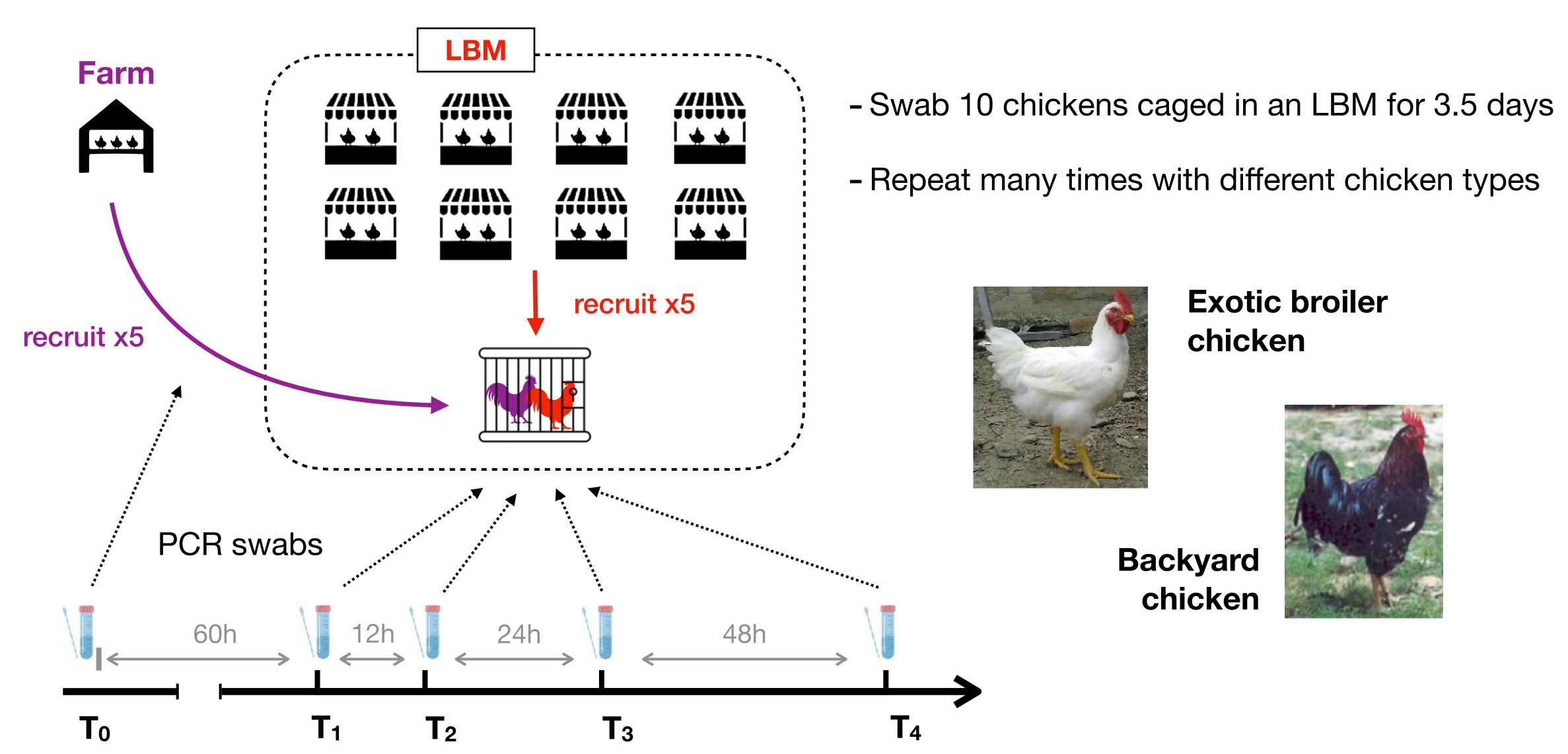
Catchment area for a single LBM (where do chickens come from?)





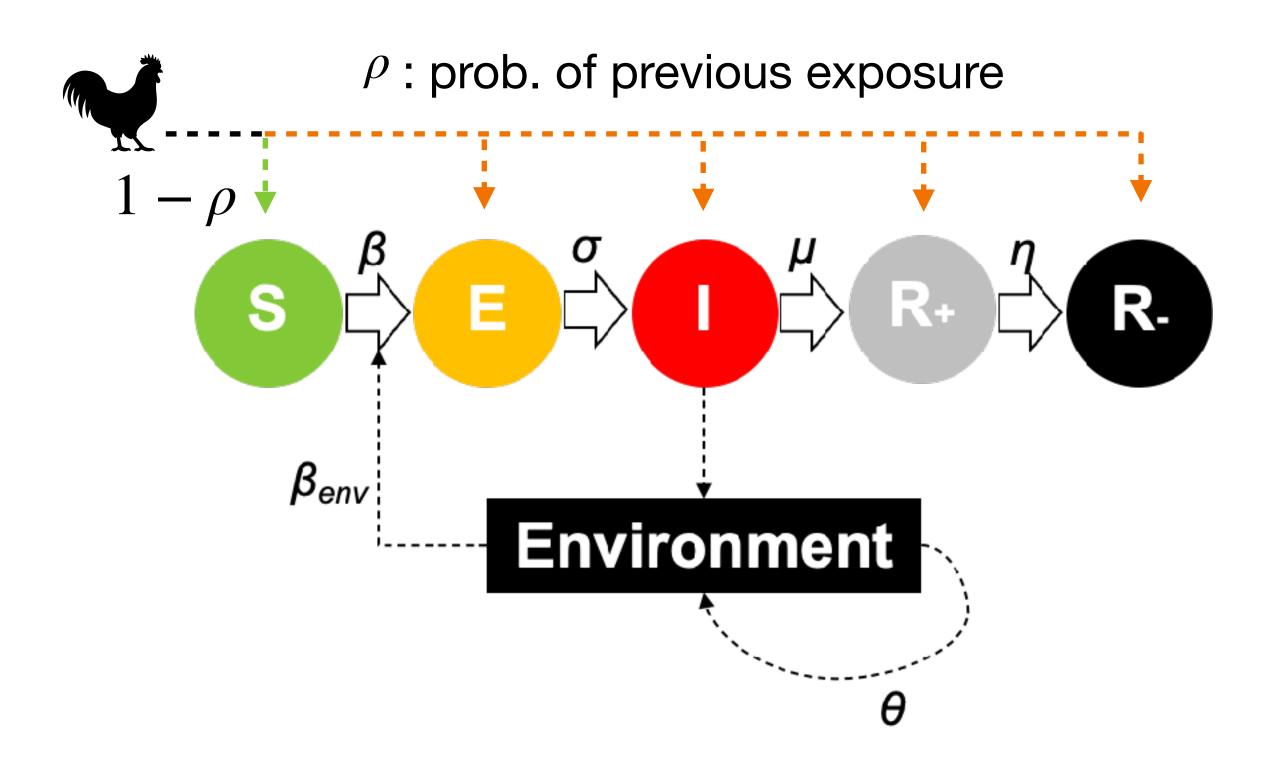
H9N2 AIV transmission in an LBM - a field experiment

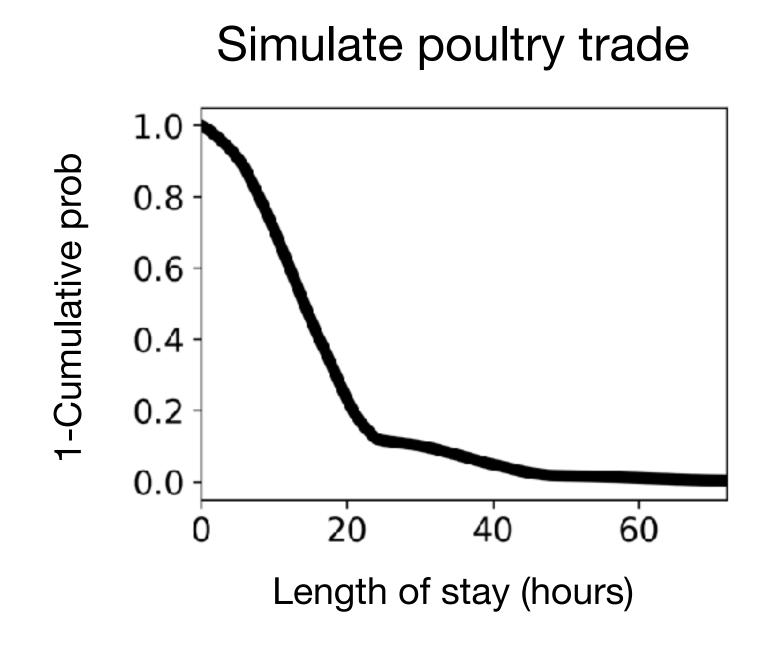
Credits: Lisa Kohnle



Fitting a mechanistic model

SEEIRR compartmental model for H9N2 AIV

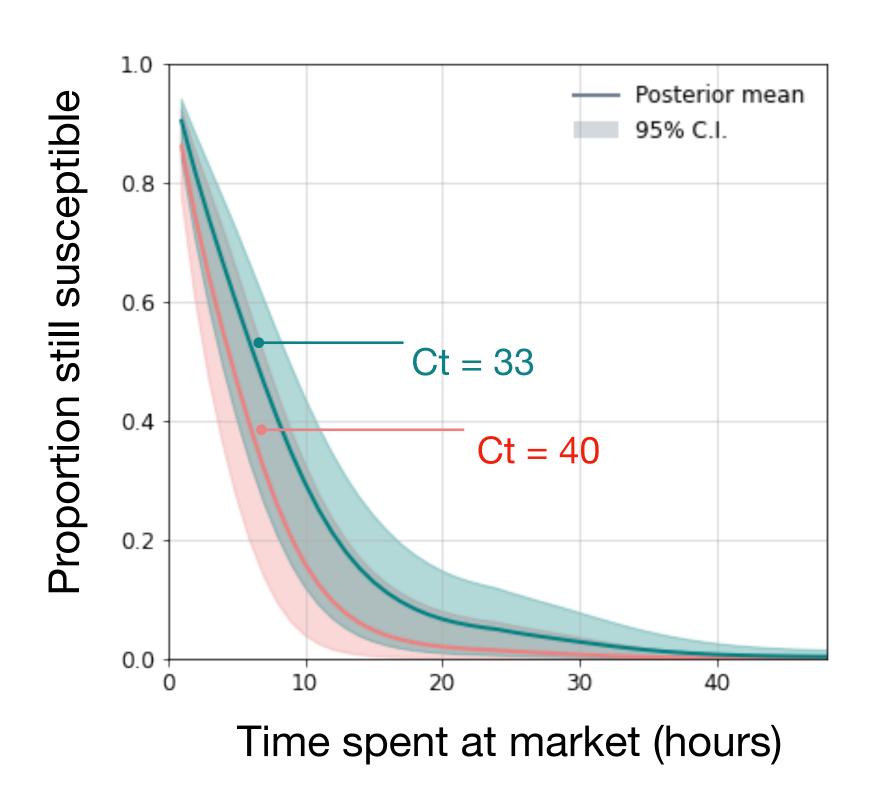




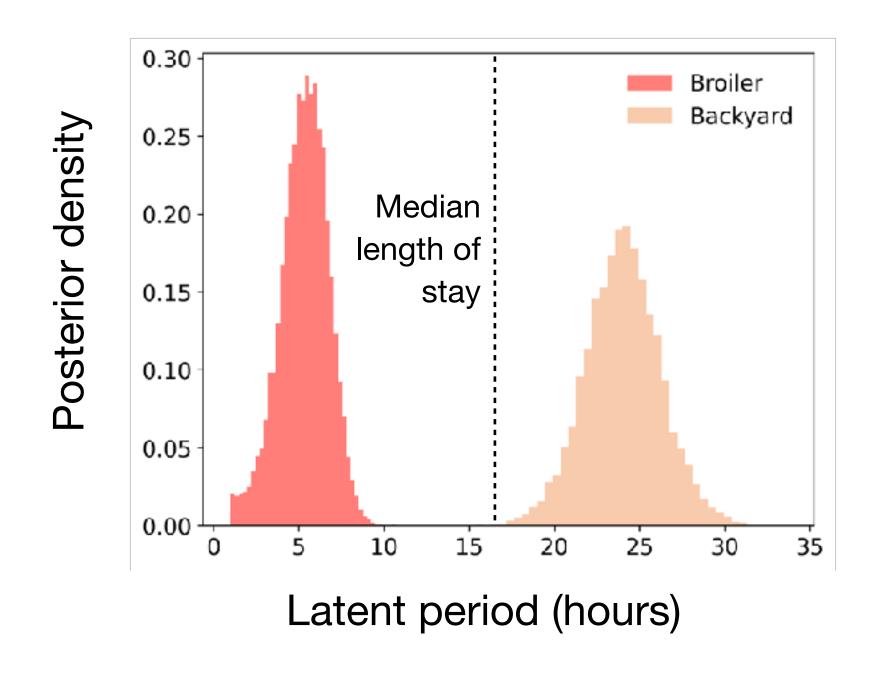
Infer parameters about transmission, disease progression, viral introduction

High H9N2 AIV transmission within LBM

(1) More than 80% of chickens become infected within 24 hours in the LBM

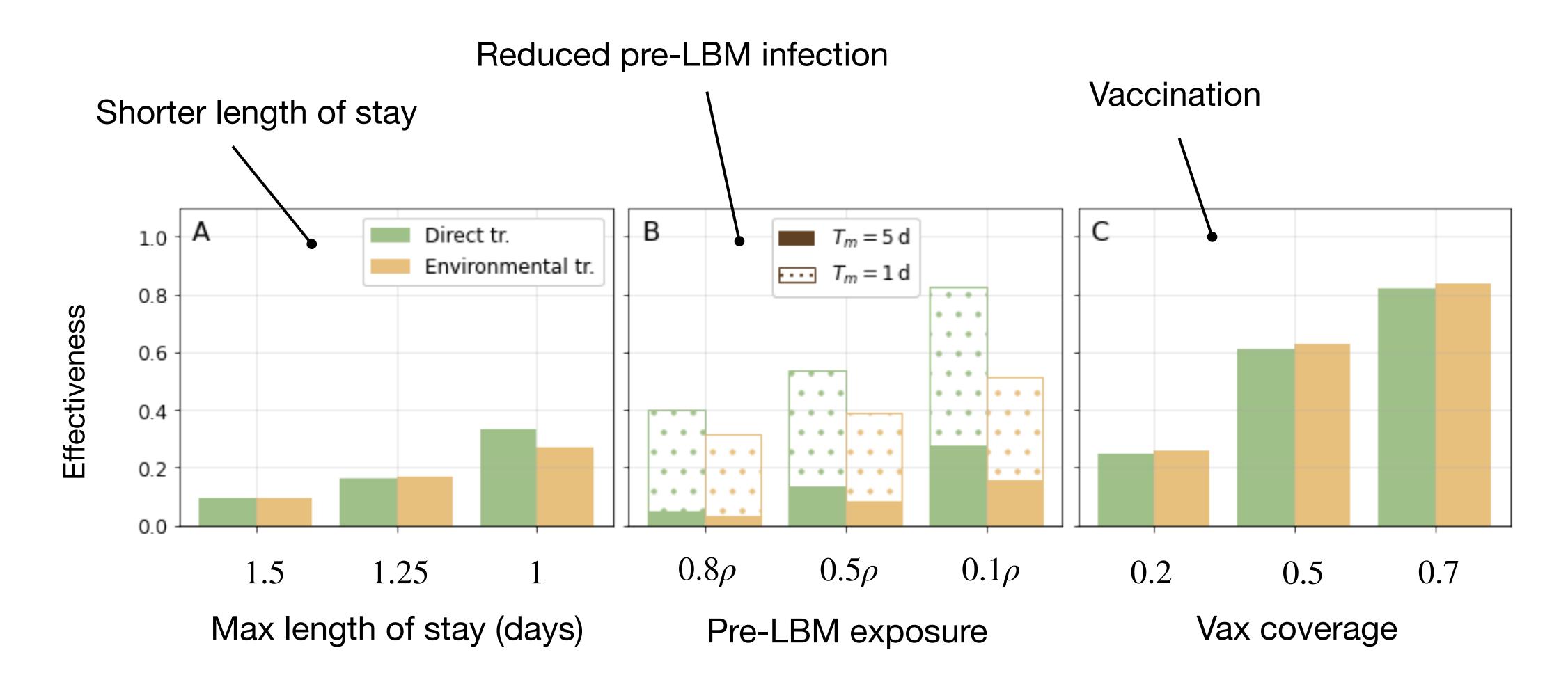


(2) broilers become infectious in ~6 hours



(3) ~10% of broilers enter the LBM as latent/infectious

Interventions



We should consider not only markets but the entire production & distribution system

EPINEST (EPIdemic NEtwork Simulation in poultry Transportation systems)

- Generate synthetic PDNs
- Data-driven
- Modular & customisable

LBMs Farms Mobile traders Stats/Modelling Geography **Poultry production** 1e+04 1e+05 Outgoing poultry flux

Available data



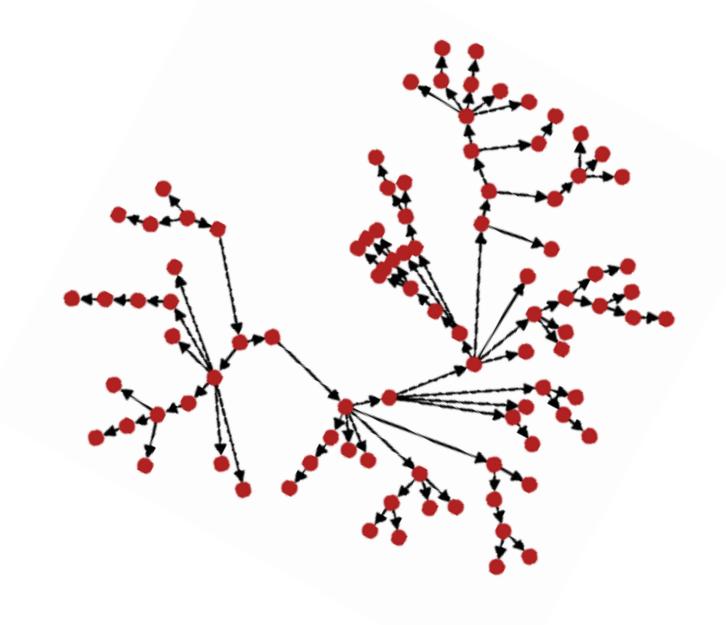
Questionnaires



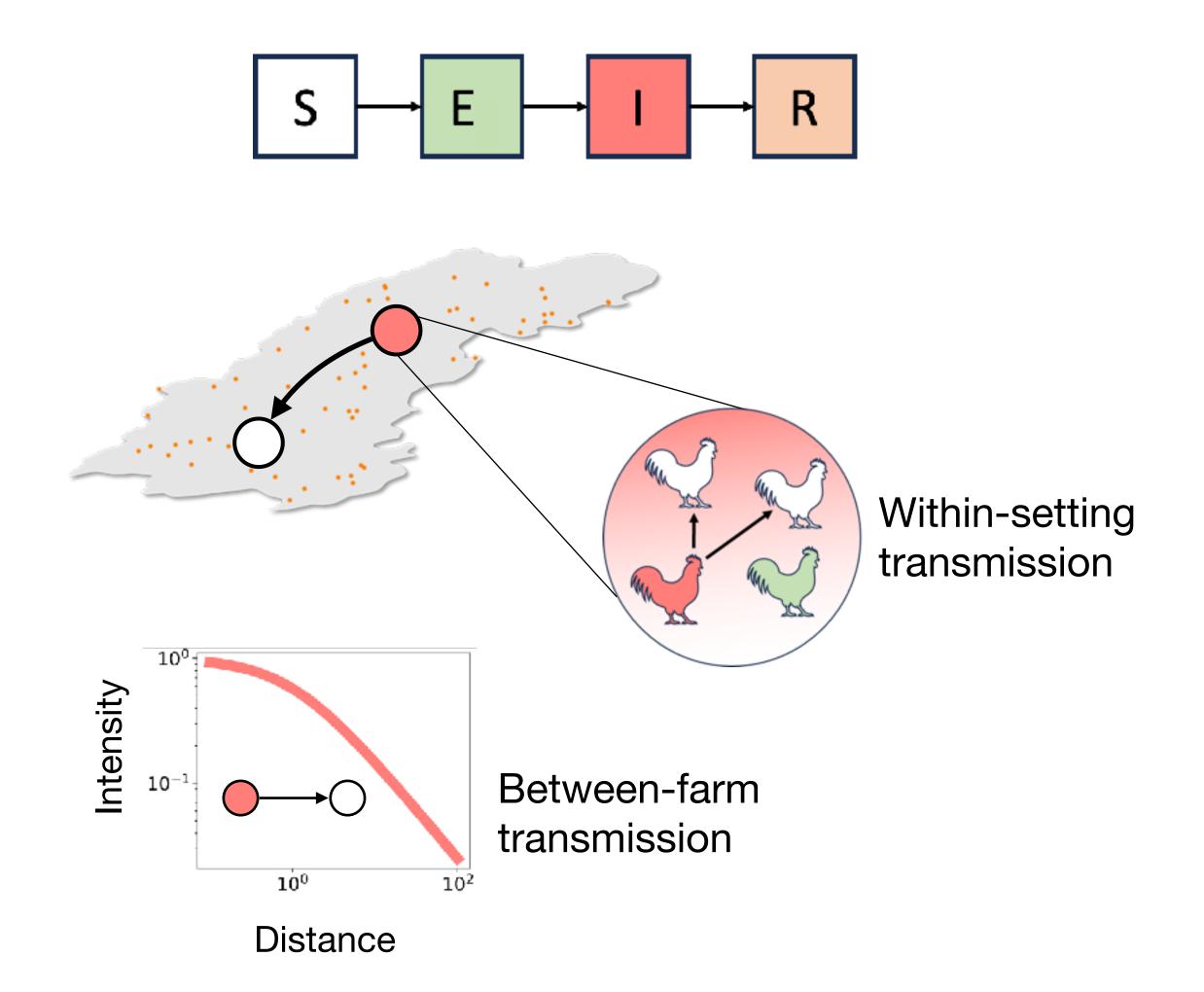
Pinotti et Al, PLOS Comp Biol, 2024

Epidemic simulations

- Explore different modelling assumptions
- Single/Multi strain simulations
- Detailed output (e.g. transmission chains)

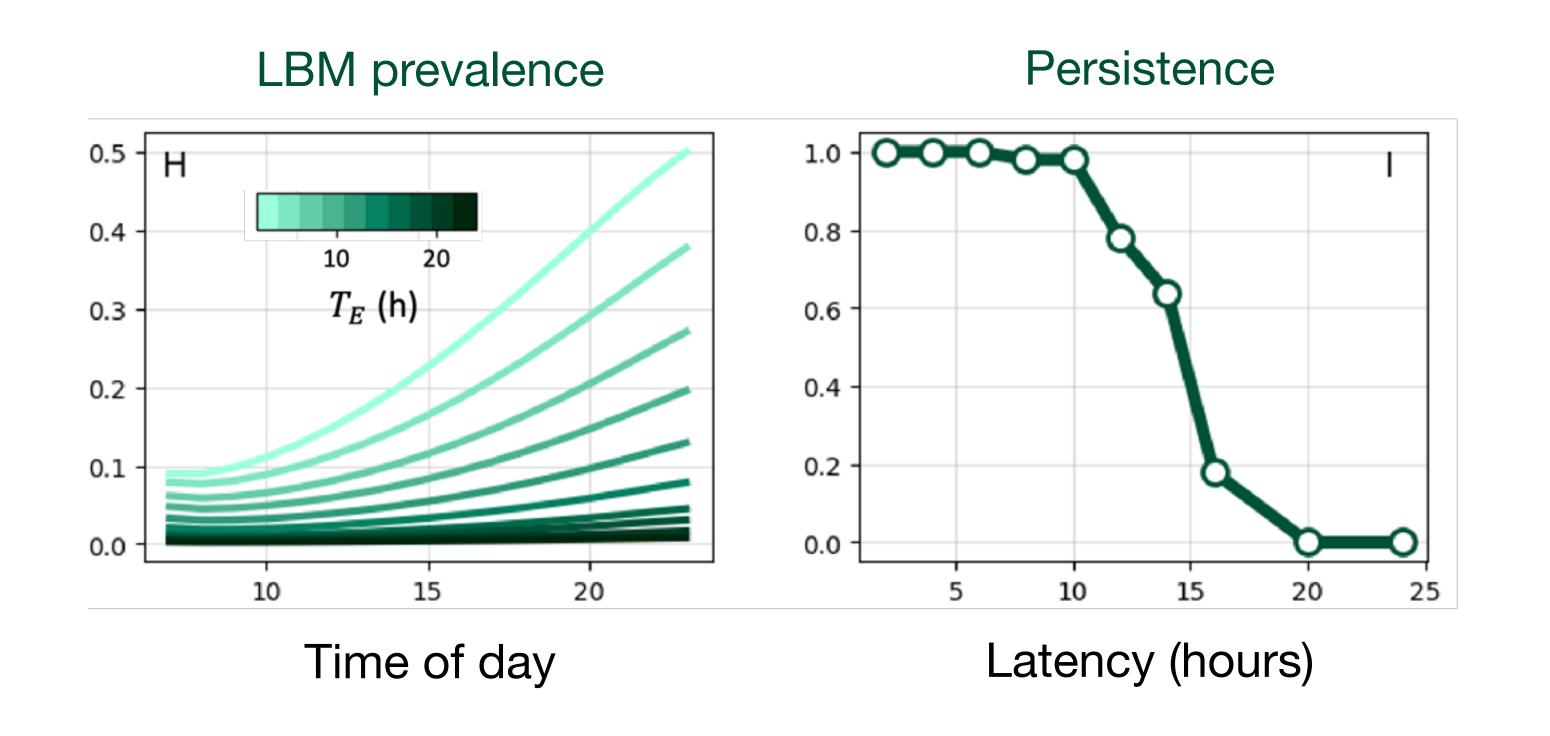


Pinotti et Al, PLOS Comp Biol, 2024



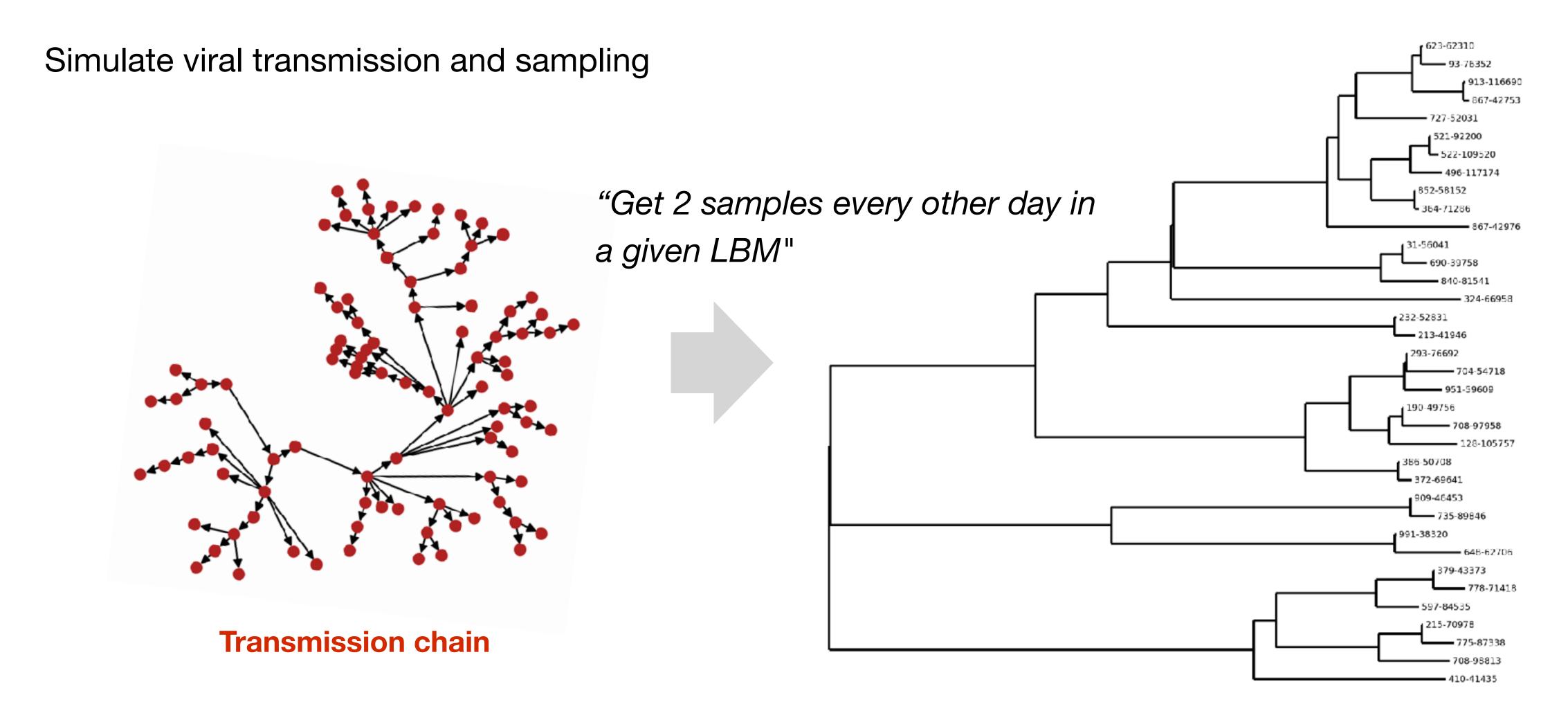
Example: characterise AIV persistence in LBMs

We vary the duration of latency T_E (infection to infectiousness)



Persistence = Probability of a long-lived transmission chain

Outlook: making sense of genetic data



Ancestry of sampled lineages

Pinotti et Al, PLOS Comp Biol, 2024

Conclusions

- Useful epidemiological insights from field experiments.
- LBMs support high transmission of H9N2 AIV.
- Veterinary public health interventions should consider the entire PDN.
- Implemented an ABM to simulate realistic poultry movements and pathogen transmission in PDNs.

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