

Koi herpesvirus disease spread among carp populations: a modeling approach

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Background

- KHVD caused by aquatic herpesvirus
 - Cyprinid herpesvirus 3 (CyHV-3)
- Affect cyprinids
 - Principally *Cyprinus carpio* – wild and ornamental
- Distributed worldwide – except Australia
- Detected in wild carps in Minnesota in 2017

Klafack et al. 2017; Bergmann *et al.*, 2020



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Objectives

- The effect of KHV in the carp population is in overall unknown
- Modeling the KHV dynamic
 - Agent-based model
 - Theoretical population



Modeling workflow

Research question

Identify outcome

Identify key process that affect outcome

Identify relevant characteristic of individuals

Stablish relationships and build model

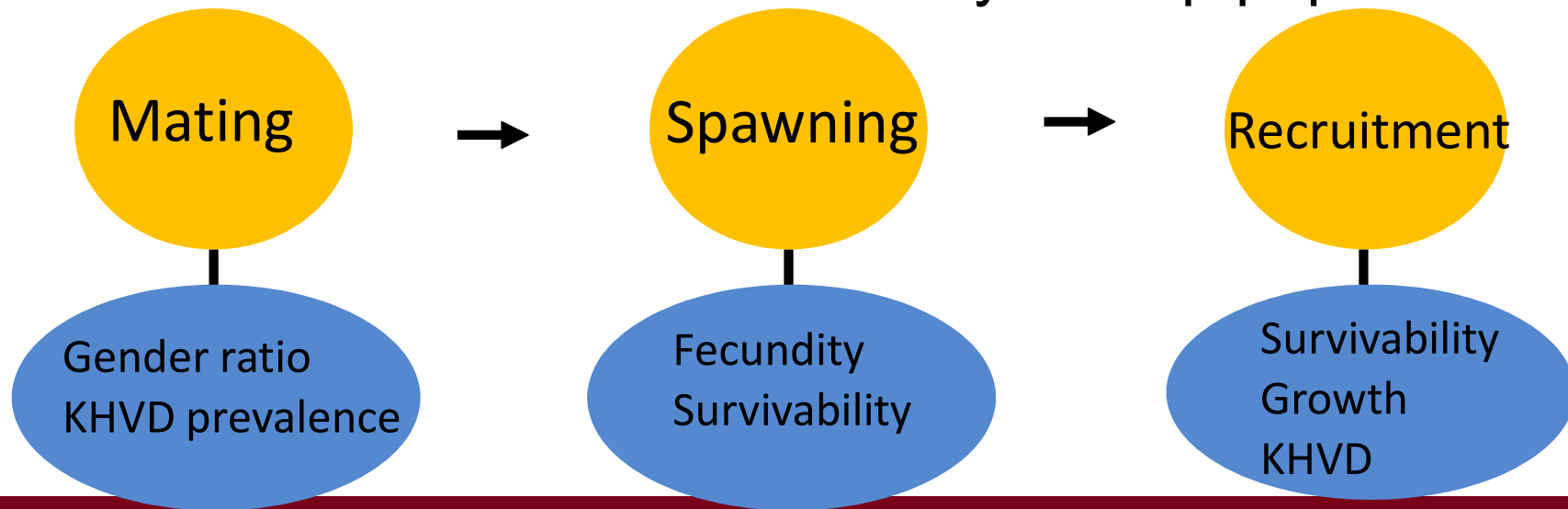
Pulliam, 2017



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Model building

- *“Impact of KHVD in the carp mortality”*
- Outcome: KHV-related mortality in carp population



Model world

- Parameters

β transmission coef.

ϕ incubation period

σ KHV mortality rate

λ Latency

κ Spawning effectivity

μ Natural mortality

T° Temperature

- State variables

S Susceptible

E Exposed

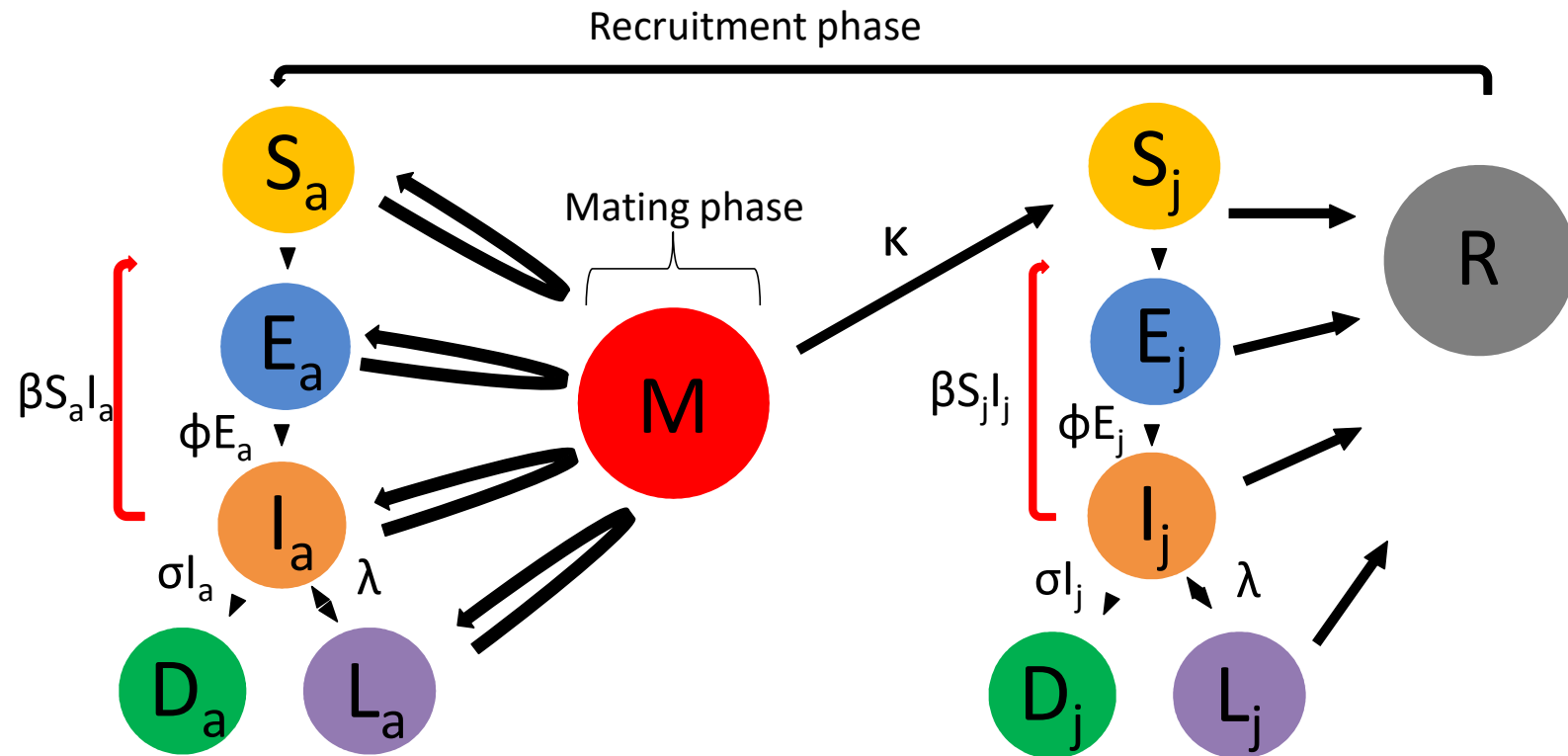
I Infected

D Dead

L Latency



Model building



References

- Klafack, S. *et al.* Genetic variability of koi herpesvirus in vitro-a natural event? *Front. Microbiol.* **8**, 1–8 (2017).
- Bergmann, S. M. *et al.* Koi herpesvirus (KHV) and KHV disease (KHVD) – a recently updated overview. *J. Appl. Microbiol.* **129**, 98–103 (2020).
- Pulliam, Juliet Creating a Model World to Address a Research Question. figshare. Media. (2015). <https://doi.org/10.6084/m9.figshare.5080390.v3>

