



Abstract

- We create and analyze reduced models for the spread of a *Wolbachia* bacteria infection in mosquitoes.
- Mosquitoes that are infected with *Wolbachia* are much less effective at transmitting zoonotic diseases. The infection persists in wild mosquitoes only if the fraction of infection exceeds a minimum threshold.
- Although the large system of differential equations capture the detailed transmission dynamics, they are difficult to analyze, which prevents practical extensions.
- The reduced models cut the analysis burden, yet capture the important properties of the original system.
- **The parameters for the reduced models are defined in terms of the ones in full model.**

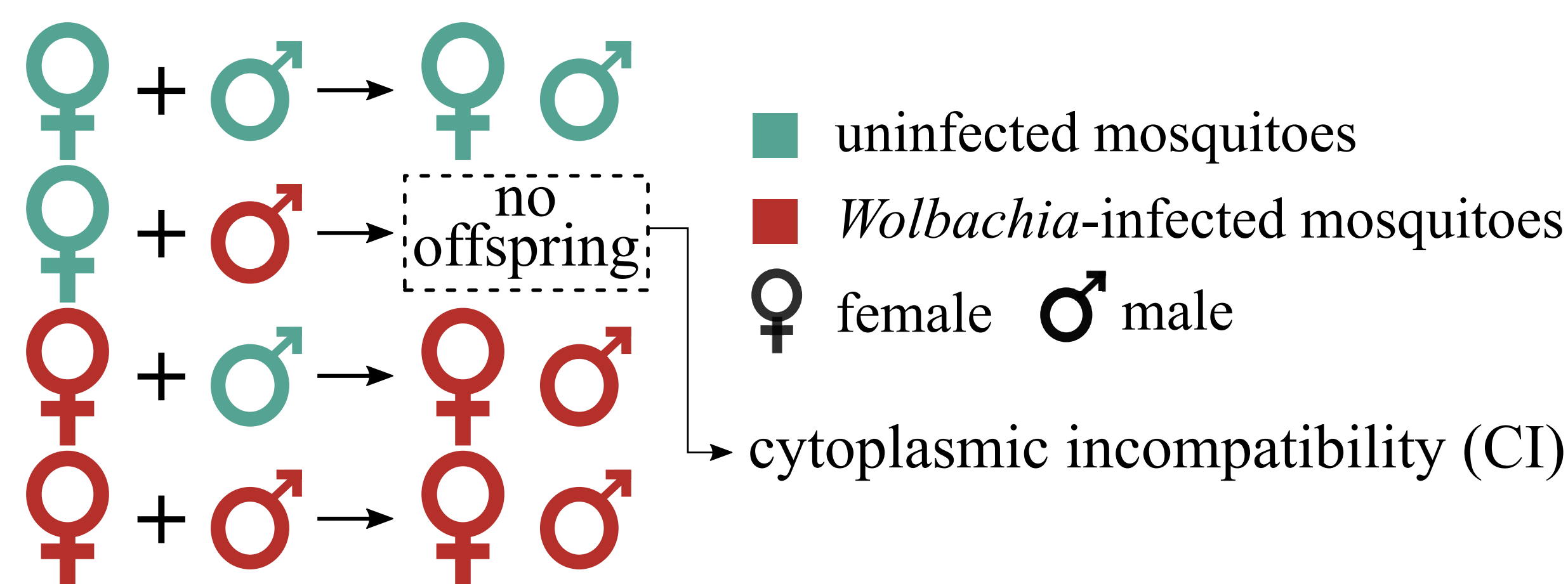
Mosquito-born Diseases v.s. *Wolbachia*

- nearly 700 million people get a mosquito-borne disease each year resulting in greater than one million deaths
- *Aedes aegypti* mosquito: the primary vector for dengue fever, chikungunya and Zika
- *Wolbachia* A promising strategy to stop diseases at source.
- a natural parasitic microbe, found in 60% insects species
- inhibits the proliferation of viruses inside the mosquito
⇒ blocks the disease transmission
- is not found in the wild *Aedes aegypti* mosquitoes

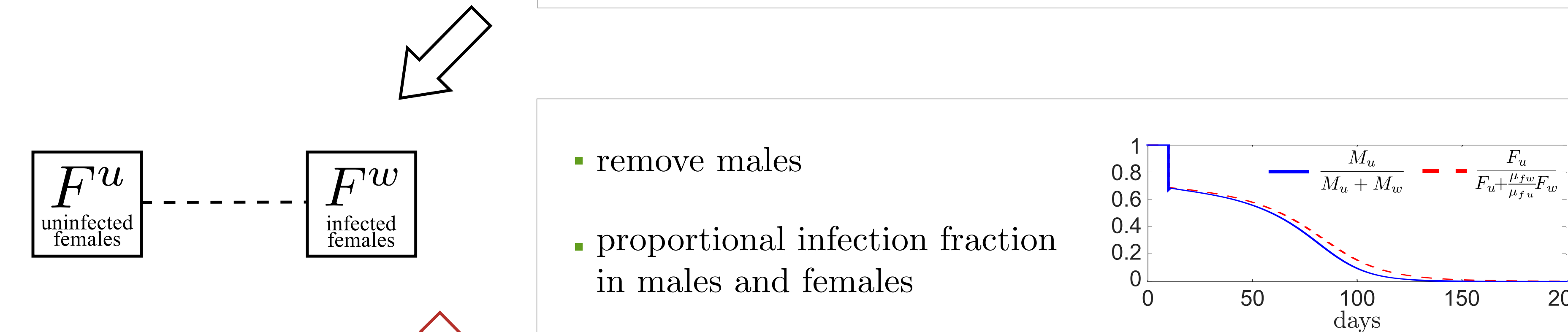
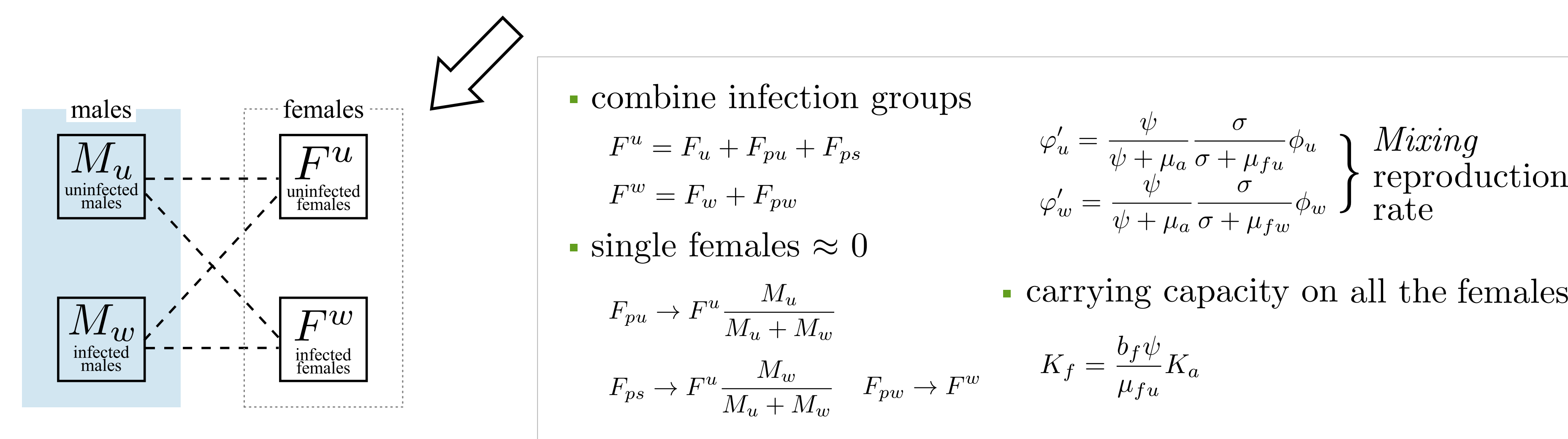
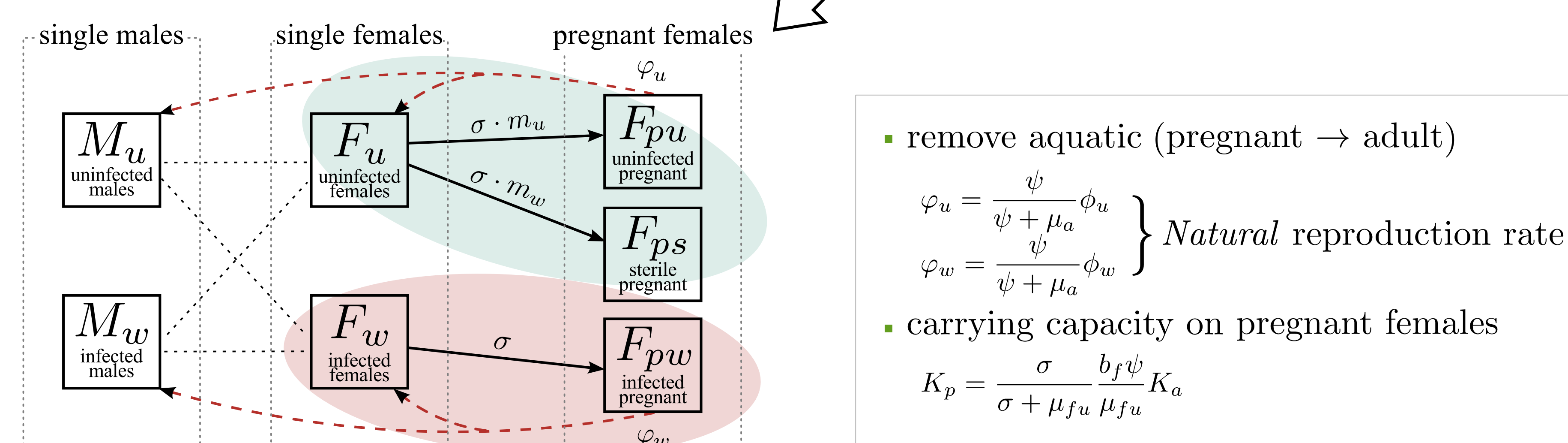
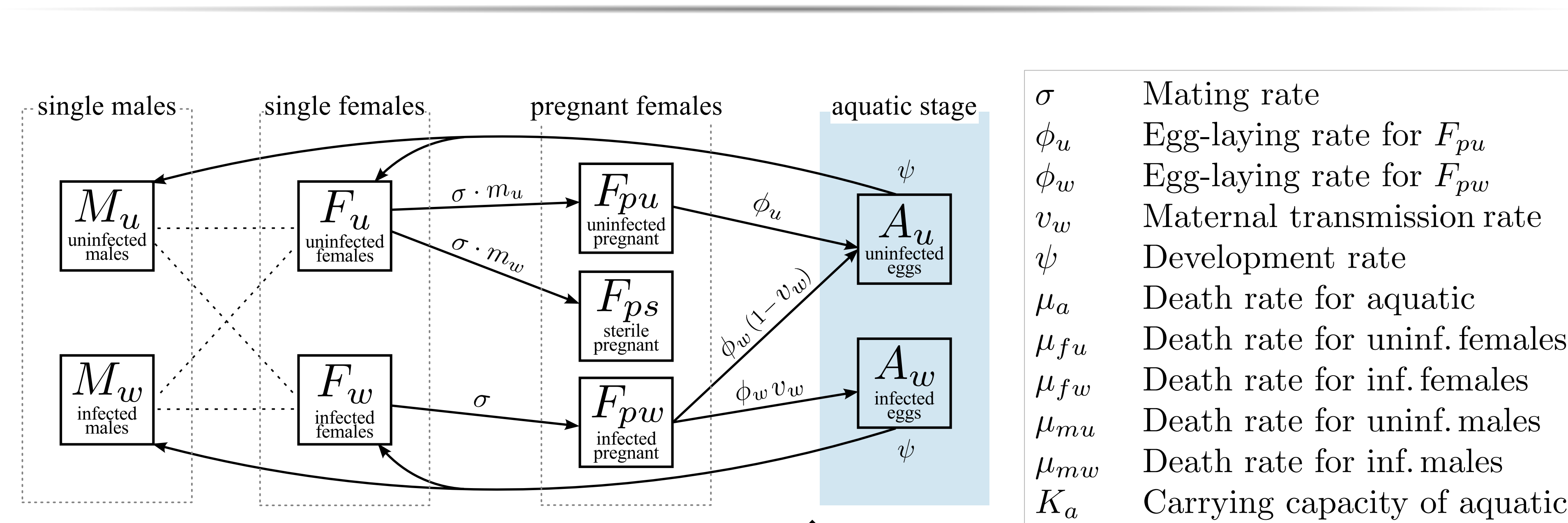
We need to create a *stable* infection in wild mosquitoes.

- field trials in Australia, Brazil, Colombia, Indonesia, Vietnam to control dengue/Zika epidemics
- Fraction of infection has to exceed a critical threshold.

Maternal transmission *Wolbachia* is maternally transmitted from infected mothers to offspring.



Model Reductions 9-ODEs → 2-ODEs



Reduced model

$$(F^u)_t = b_f \varphi'_u \frac{F^u}{F^u + \frac{\mu_{fw}}{\mu_{fu}} F^w} \left(1 - \frac{F^u + F^w}{K_f}\right) F^u + b_f (1 - v_w) \varphi'_w \left(1 - \frac{F^u + F^w}{K_f}\right) F^w - \mu_{fu} F^u + d(F^u_{xx} + F^u_{yy})$$

$$(F^w)_t = b_f v_w \varphi'_w \left(1 - \frac{F^u + F^w}{K_f}\right) F^w - \mu_{fw} F^w + d(F^w_{xx} + F^w_{yy})$$

Future work

- Include practical extensions
- spatial heterogeneity
- temporal variation

Full model

$$\dot{A}_u = (\phi_u F_{pu} + (1 - v_w) \phi_w F_{pw}) \left(1 - \frac{A_u + A_w}{K_a}\right) - (\mu_a + \psi) A_u$$

$$\dot{A}_w = v_w \phi_w F_{pw} \left(1 - \frac{A_u + A_w}{K_a}\right) - (\mu_a + \psi) A_w$$

$$\dot{F}_u = b_f \psi A_u - (\sigma + \mu_{fu}) F_u$$

$$\dot{F}_w = b_f \psi A_w - (\sigma + \mu_{fw}) F_w$$

$$\dot{F}_{pu} = \sigma m_u F_u - \mu_{fu} F_{pu}, \quad m_u = M_u / (M_u + M_w)$$

$$\dot{F}_{ps} = \sigma m_w F_u - \mu_{fu} F_{ps}, \quad m_w = M_w / (M_u + M_w)$$

$$\dot{F}_{pw} = \sigma F_w - \mu_{fw} F_{pw}$$

$$\dot{M}_u = b_m \psi A_u - \mu_{mu} M_u$$

$$\dot{M}_w = b_m \psi A_w - \mu_{mw} M_w$$

Model Comparisons

Reduced models preserve two important properties of the detailed 9-ODE model.

Basic reproductive number $\mathbb{R}_0$

- number of new infections created by a single *Wolbachia*-infected mosquito, given all the rest mosquitoes are fully susceptible
- often used as a threshold condition for disease outbreak (predict early epidemic)
- Full model: $\mathbb{R}_0^{(9)} = v_w \frac{\mu_{fu} \phi_w (\sigma + \mu_{fu})}{\mu_{fw} \phi_u (\sigma + \mu_{fw})} \approx 0.722$

$$\mathbb{R}_0^{(9)} = \mathbb{R}_0^{(7)} = \mathbb{R}_0^{(4)} = \mathbb{R}_0^{(2)}$$

Critical threshold for a stable *Wolbachia* infection

- a minimum threshold must be exceeded to establish a stable infection in wild mosquitoes
- backward bifurcation with an unstable steady-state

