

Remodeling selection to optimize disease forecasts and policies

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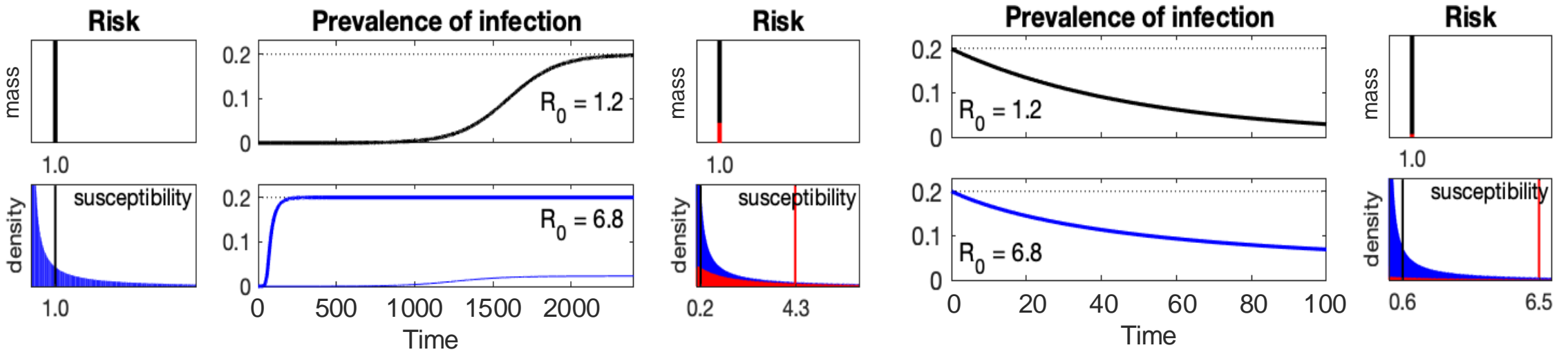
Background:

When variation in susceptibility is accounted for...

- estimated reproduction numbers are higher
- predicted impact of uniform control measures is lower

SI model and selective depletion bias:

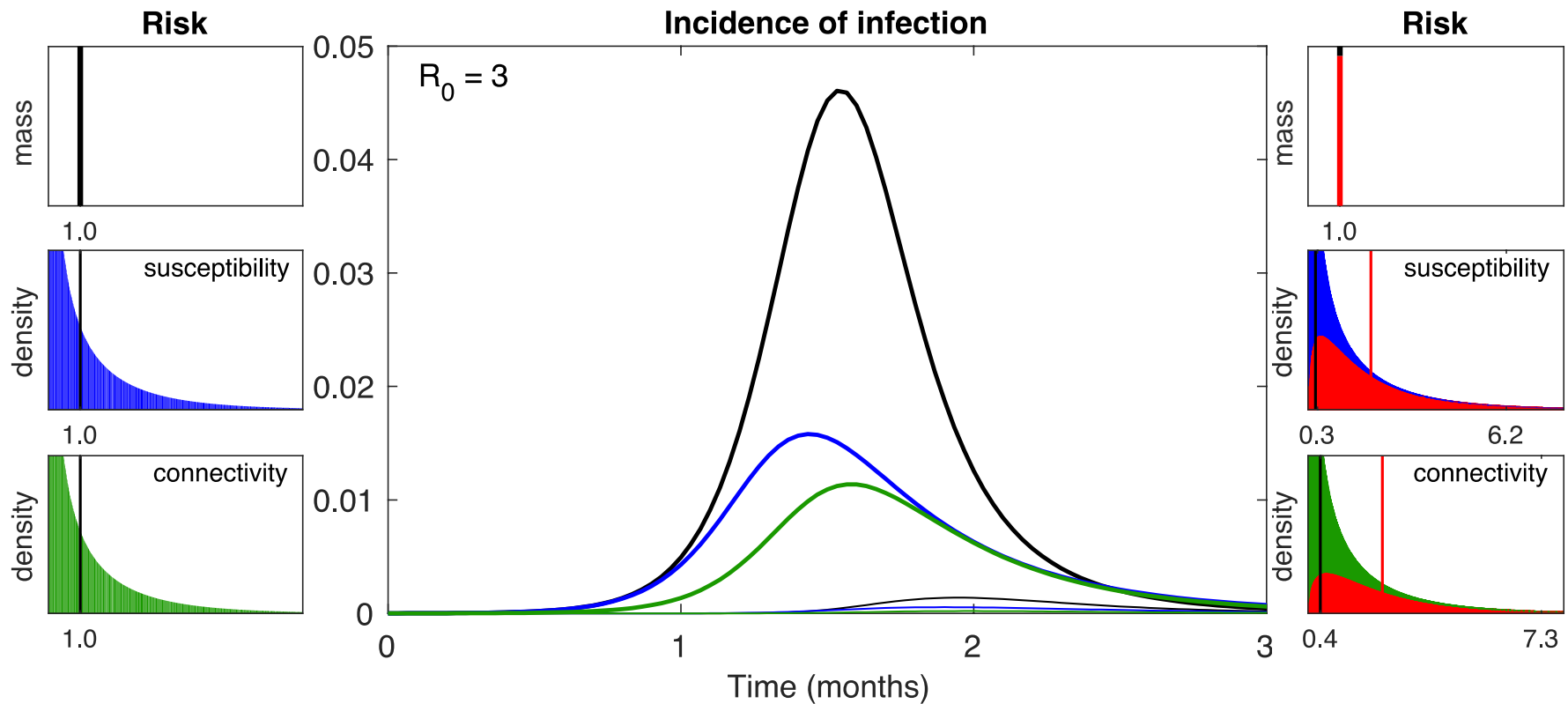
$$\frac{dS(x)}{dt} = q(x)\mu - \beta \int I(u)du xS(x) - \mu S(x), \quad \frac{dI(x)}{dt} = \beta \int I(u)du xS(x) - \mu I(x), \quad \mathcal{R}_0 = \frac{\beta}{\mu}$$



MGMG, AM Blagborough, KE Langwig, B Ringwald 2024 Remodelling selection to optimise disease forecasts and policies. *J Phys A: Math Theor* 57:103001.

SIR model and selective depletion bias:

$$\frac{dS(x)}{dt} = -\beta \int I(u) du xS(x), \quad \frac{dI(x)}{dt} = \beta \int I(u) du xS(x) - \gamma I(x), \quad \mathcal{R}_0 = \frac{\beta}{\mu}$$




This talk:

Variation in susceptibility can be estimated by remodeling selection when...

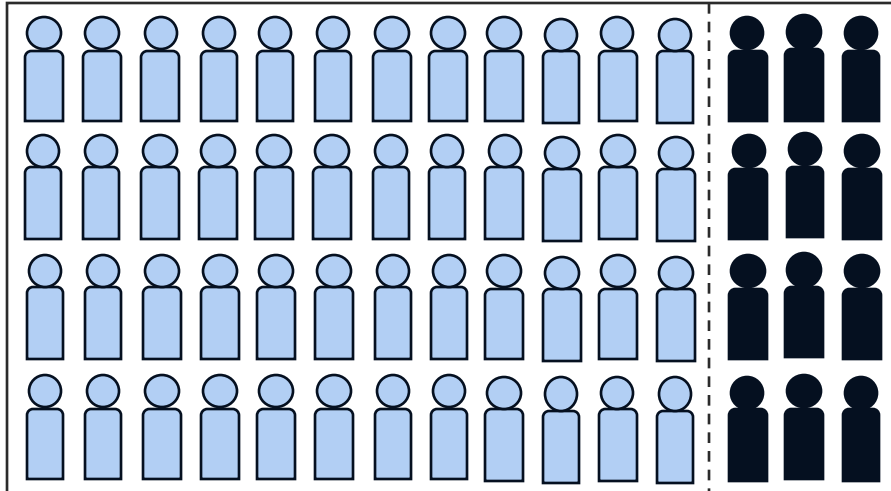
- observational data can be stratified over selection gradients
- experimental data can be generated over selection gradients

Observational data stratified by region:

 Low susceptibility (α_1)

 High susceptibility (α_2)

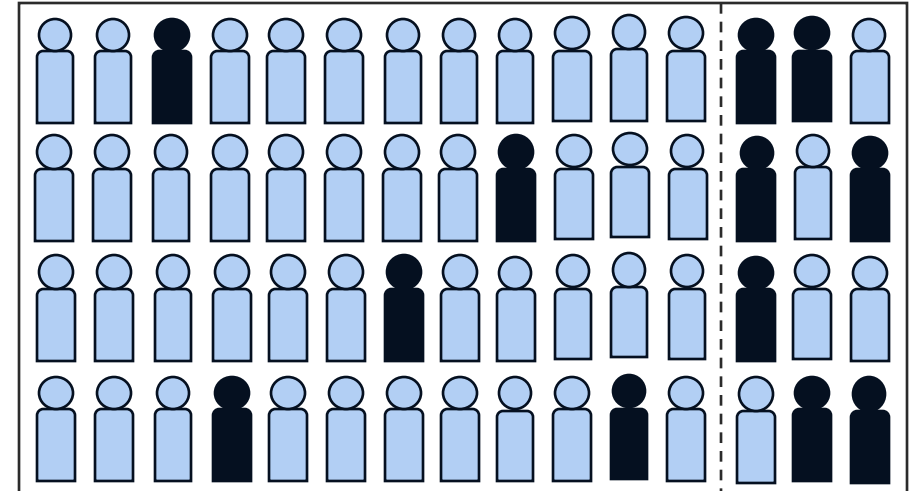
Metapopulation I



Region A

Region B

Metapopulation II



Region A

Region B

SI metapopulation model:

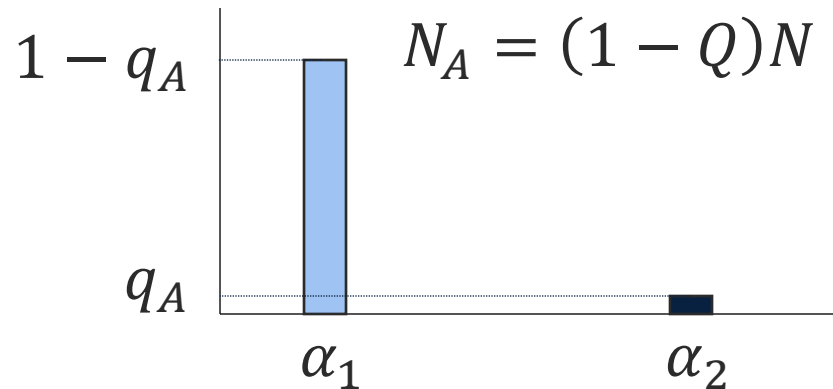
$$\frac{dS_{A1}}{dt} = (1 - q_A)\mu - \beta(I_{A1} + I_{A2})\alpha_1 S_{A1} - \mu S_{A1}$$

$$\frac{dI_{A1}}{dt} = \beta(I_{A1} + I_{A2})\alpha_1 S_{A1} - \mu I_{A1}$$

$$\frac{dS_{A2}}{dt} = q_A\mu - \beta(I_{A1} + I_{A2})\alpha_2 S_{A2} - \mu S_{A2}$$

$$\frac{dI_{A2}}{dt} = \beta(I_{A1} + I_{A2})\alpha_2 S_{A2} - \mu I_{A2}$$

$$\mathcal{R}_{0A} = [(1 - q_A)\alpha_1 + q_A\alpha_2] \frac{\beta}{\mu}$$



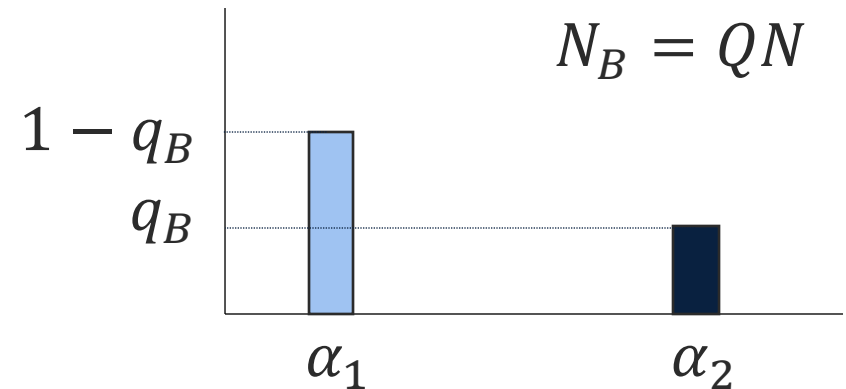
$$\frac{dS_{B1}}{dt} = (1 - q_B)\mu - \beta(I_{B1} + I_{B2})\alpha_1 S_{B1} - \mu S_{B1}$$

$$\frac{dI_{B1}}{dt} = \beta(I_{B1} + I_{B2})\alpha_1 S_{B1} - \mu I_{B1}$$

$$\frac{dS_{B2}}{dt} = q_B\mu - \beta(I_{B1} + I_{B2})\alpha_2 S_{B2} - \mu S_{B2}$$

$$\frac{dI_{B2}}{dt} = \beta(I_{B1} + I_{B2})\alpha_2 S_{B2} - \mu I_{B2}$$

$$\mathcal{R}_{0B} = [(1 - q_B)\alpha_1 + q_B\alpha_2] \frac{\beta}{\mu}$$



$$Q = (1 - Q)q_A + Qq_B$$

$$\mathcal{R}_0 = (1 - Q)\mathcal{R}_{0A} + Q\mathcal{R}_{0B} = \frac{\beta}{\mu}$$

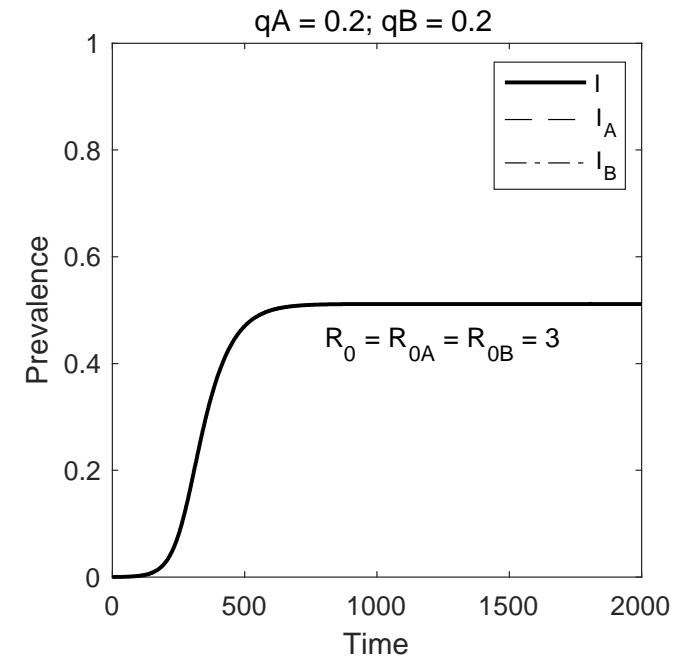
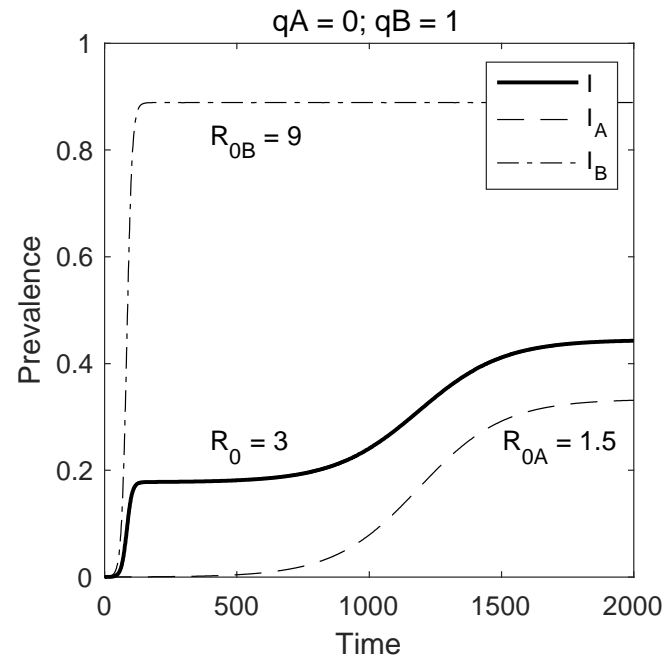
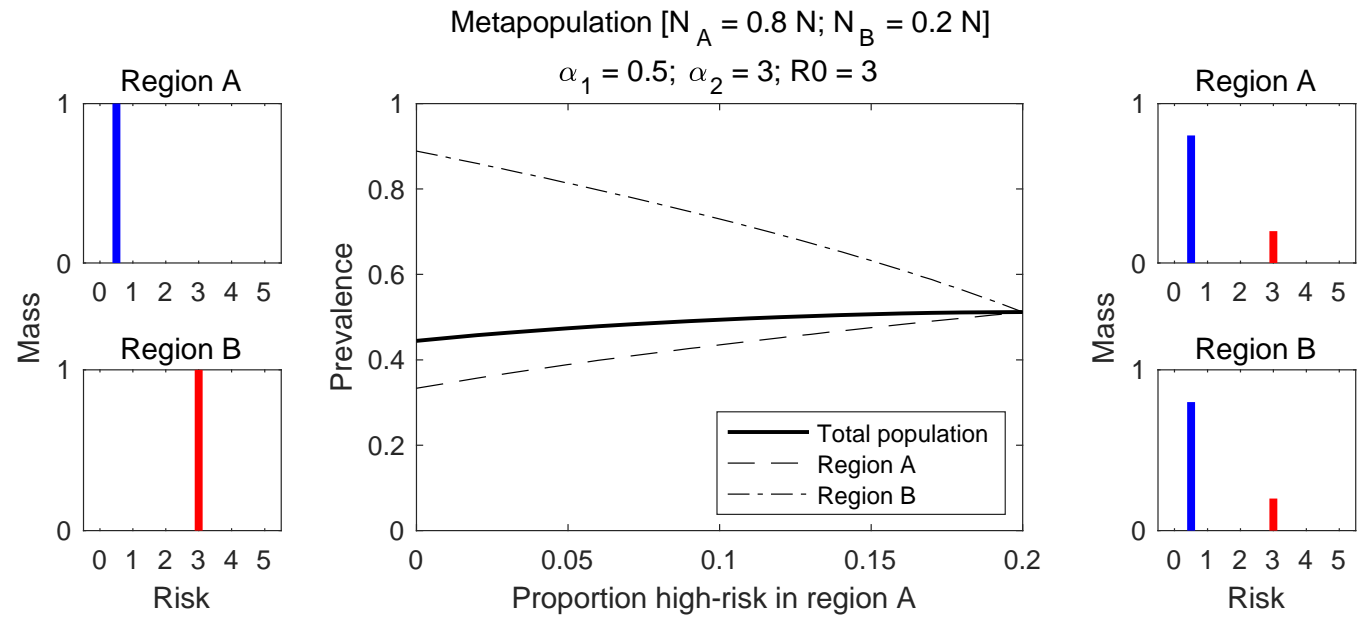
$$\text{Mean}(\alpha) = (1 - Q)\alpha_1 + Q\alpha_2 = 1$$

$$\text{Var}(\alpha) = (1 - Q)(\alpha_1 - 1)^2 + Q(\alpha_2 - 1)^2$$

Endemic equilibrium:

$$\frac{\alpha_1(1 - q_A)}{\beta I_A \alpha_1 + \mu} + \frac{\alpha_2 q_A}{\beta I_A \alpha_2 + \mu} = \frac{1}{\beta}$$

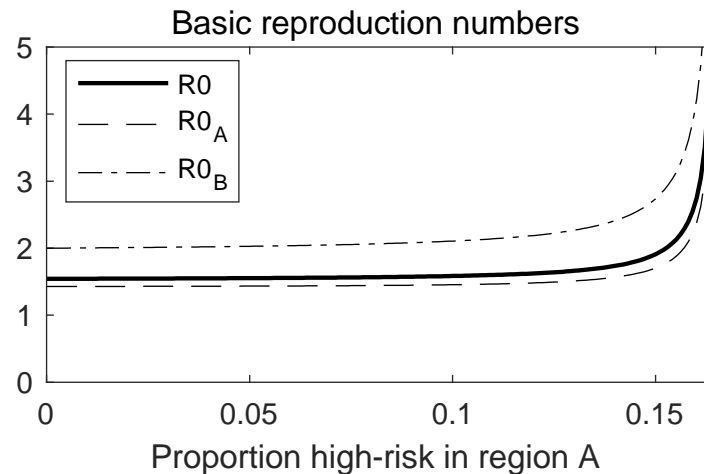
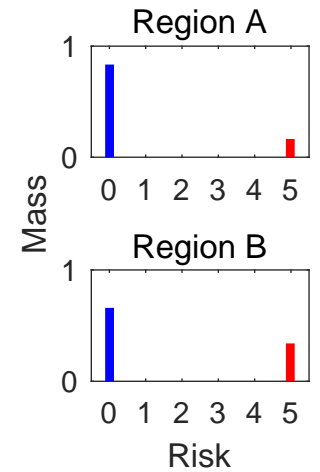
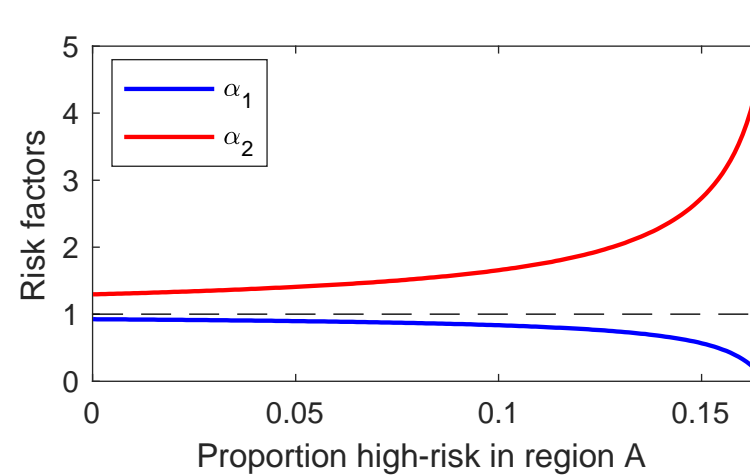
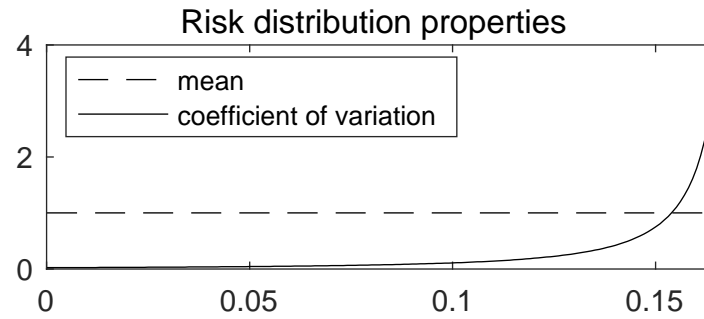
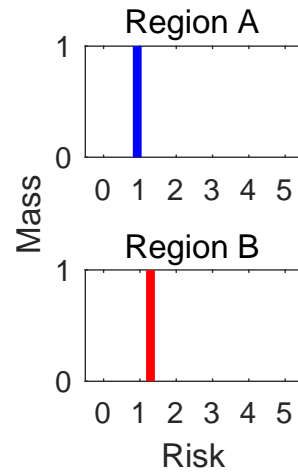
$$\frac{\alpha_1(1 - q_B)}{\beta I_B \alpha_1 + \mu} + \frac{\alpha_2 q_B}{\beta I_B \alpha_2 + \mu} = \frac{1}{\beta}$$

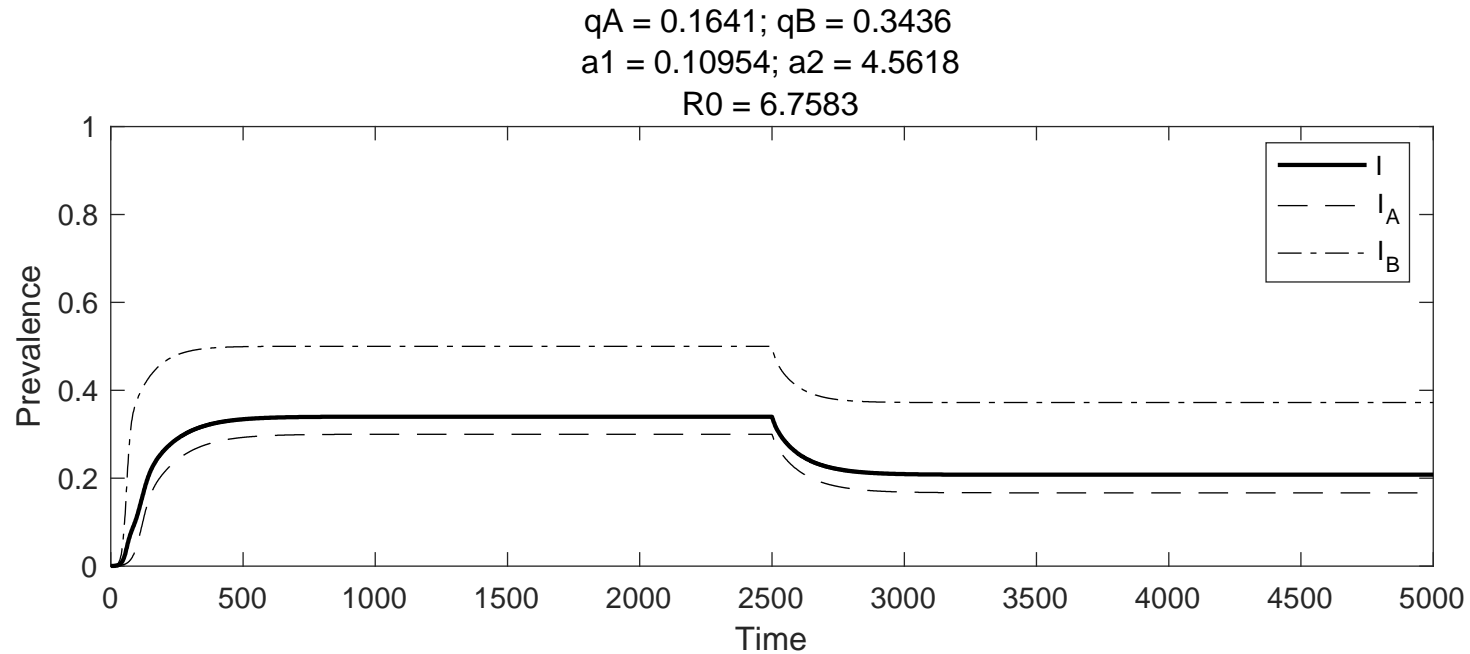
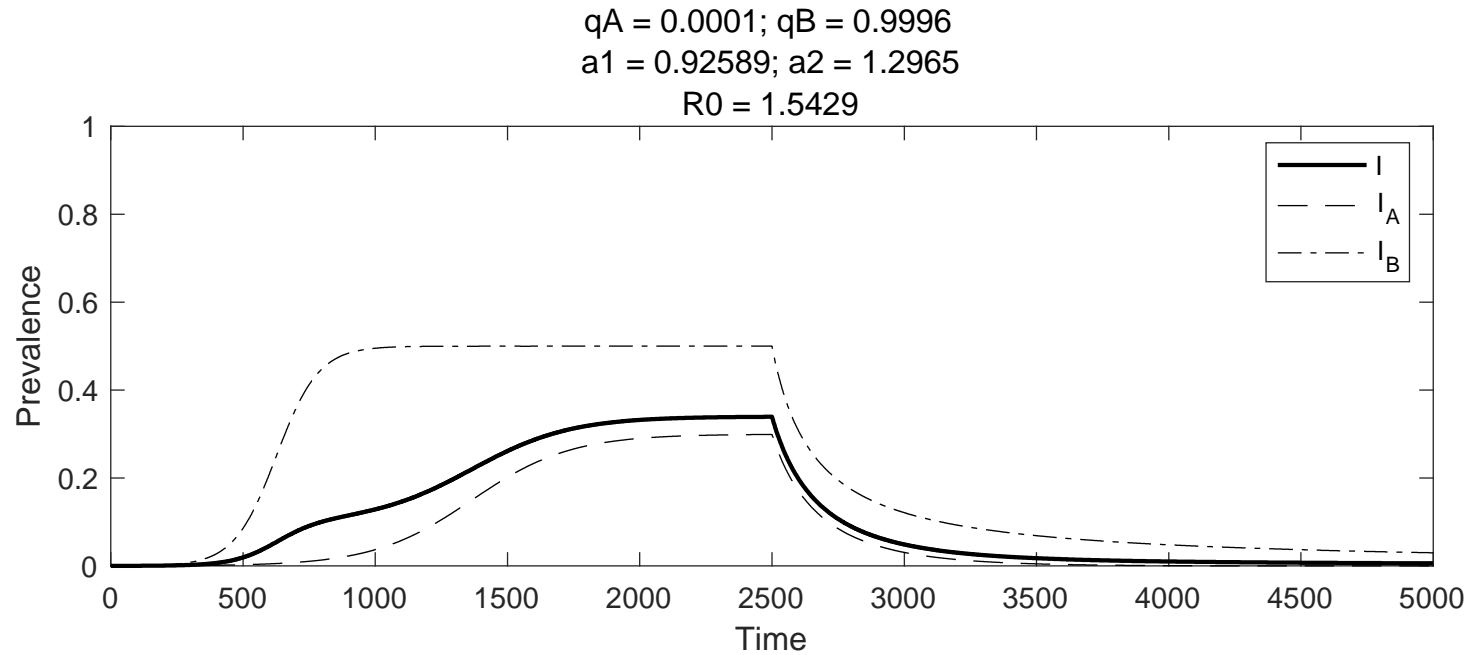


Given I_A and I_B the inverse problem can be solved exactly given the proportion high risk in one of the regions:

$$I_A = 0.3$$

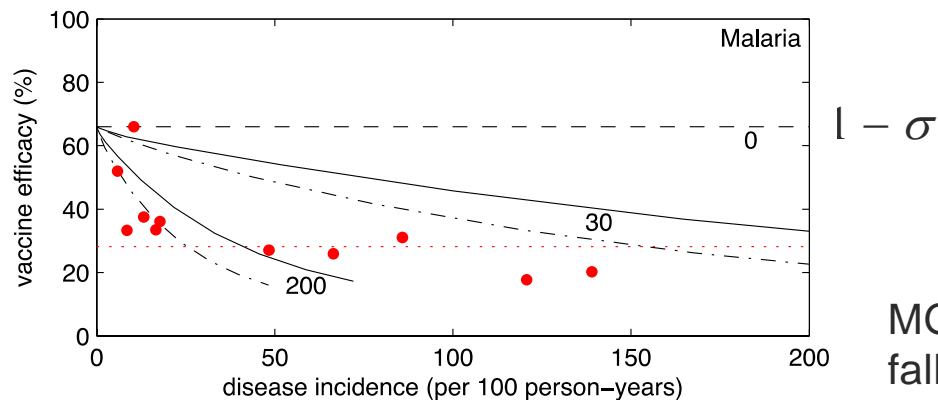
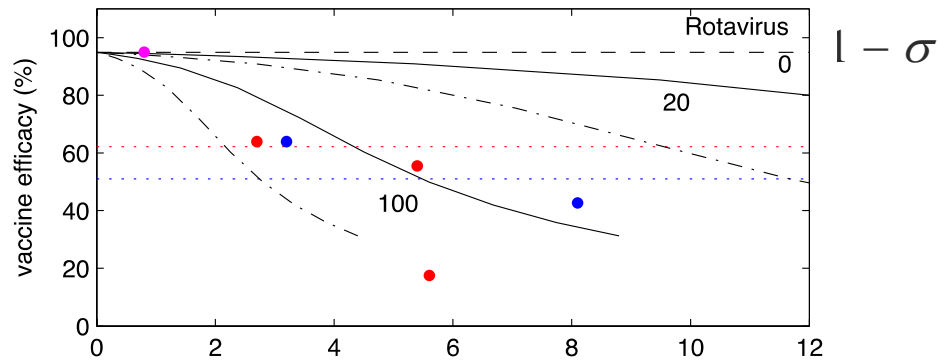
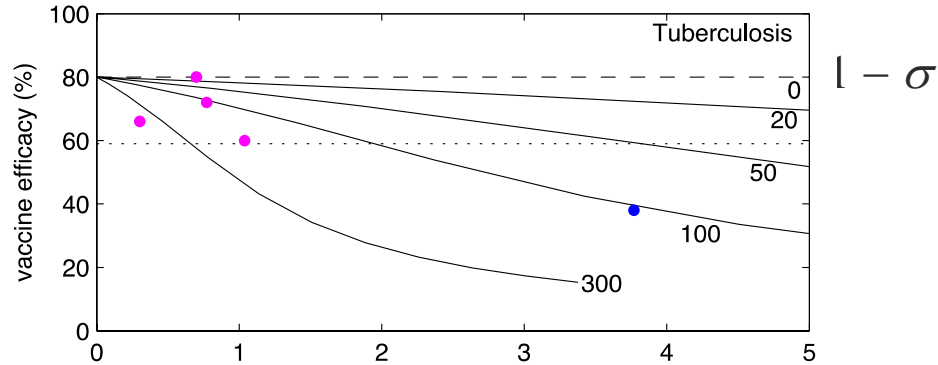
$$I_B = 0.5$$





A classical metapopulation model would overpredict the impact of an intervention that uniformly reduces susceptibility, such as a vaccine

Vaccine trials (low efficacy under high exposure):



Control group:

$$\frac{dS_c}{dt}(x) = -\lambda x S_c(x) + \gamma(q(x) - S_c(x)),$$

$$\frac{dI_c}{dt}(x) = \lambda x S_c(x),$$

Vaccine group:

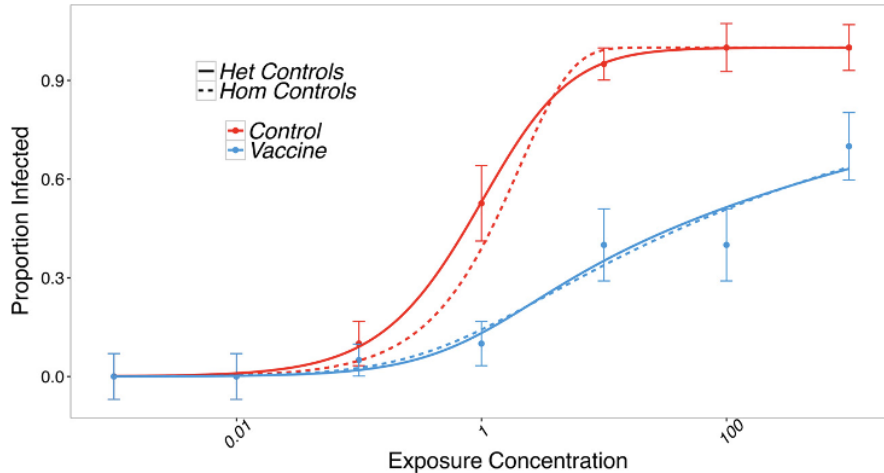
$$\frac{dS_v}{dt}(x) = -\sigma \lambda x S_v(x) + \gamma(q(x) - S_v(x)),$$

$$\frac{dI_v}{dt}(x) = \sigma \lambda x S_v(x).$$

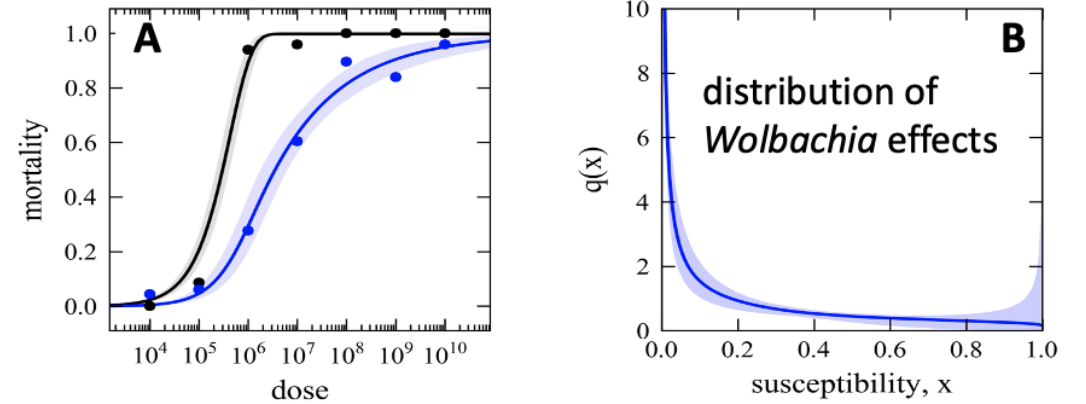
MGMG, SB Gordon, DG Lalloo 2016 Clinical Trials: The mathematics of falling vaccine efficacy with rising disease incidence. *Vaccine* 34:3007.

Dose-response infection experiments:

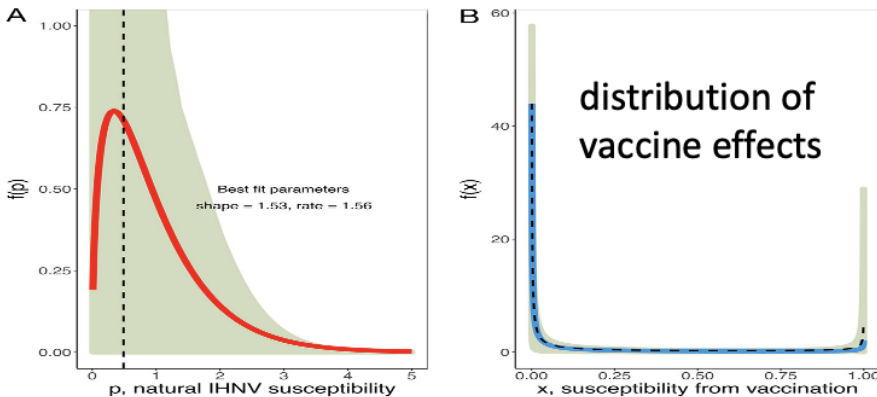
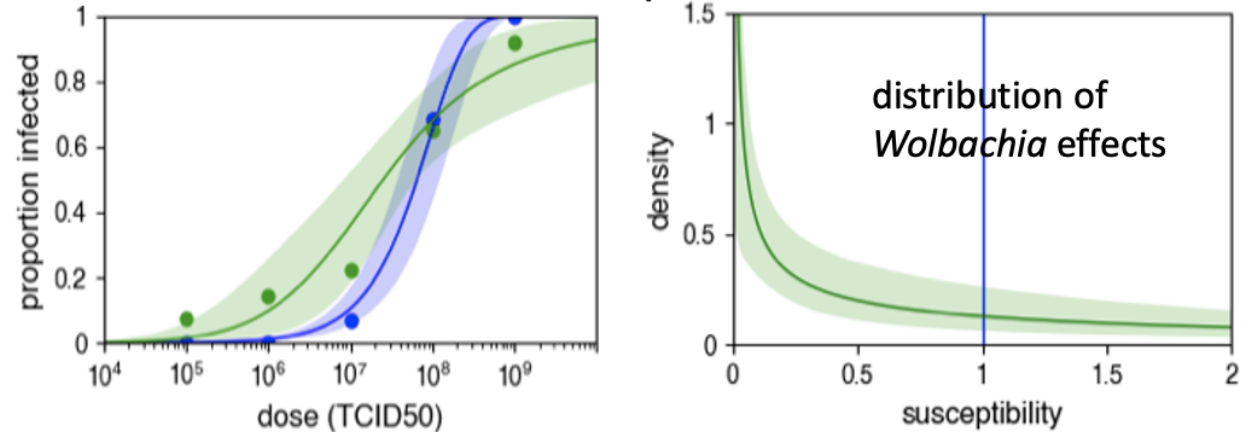
Fish (trout)



Fruit flies



Mosquitoes



KE Langwig *et al* 2017 *mBio* 8:e00796-17.
 KE Langwig *et al* 2019 *Sci Rep* 9:3203.

D Pessoa *et al* 2014 *PLOS Comput Biol* 10:e1003773.
 JG King *et al* 2018 *Nat Commun* 9:1483.

Summary:

- “Selective depletion bias” applies generally when the population mean value of a trait (such as susceptibility) changes over time or across environments due to selective depletion of the frailest (most susceptible) individuals, and this is misinterpreted as **individuals** (rather than **population** composition) changing.
- “Remodeling selection” is a novel approach (method) to infer unobserved variation from its effects on observational or experimental data collected over selection gradients.