

SMIDDY 2012

# Case and basic reproduction number for a curable sexually transmitted infection

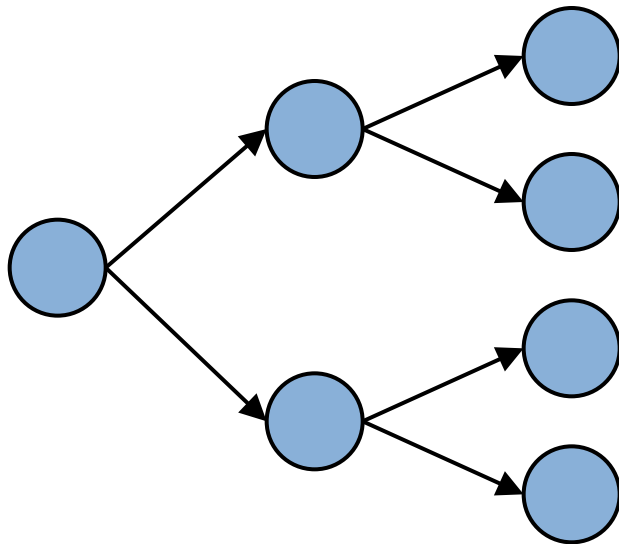
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# Basic reproduction number ( $R_0$ )

- >  $R_0$  is used to determine whether an infectious disease can invade in a susceptible population



$$R_0 = 2$$

# Frequency-de transmitted d transmission behaviour



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**Mathematical  
Biosciences**

[www.elsevier.com/locate/mbs](http://www.elsevier.com/locate/mbs)

James O. Lloyd-Sm

## Analysis and simulation of a stochastic, discrete-individual model of STD transmission with partnership concurrency

Stephen E. Chick <sup>a,\*</sup>, Andrew L. Adams <sup>b</sup>, James S. Koopman <sup>c</sup>

ment shown  
electronic Ap  
 $R_0$ , is the exp  
an infectious  
(Diekmann &

dition for dis  $\pi_u$  in the unpartnered population. If the individual is partnered with an uninfected individual, then  $R_0 = Y(0)$ . A reasonable assumption [3] is to define  $R_0 = Z(0)$  as the expected number of precise secondary cases, given that a newly infected individual was infected by a partner, but that the rest one in of the population is susceptible. This definition is used in the remainder of the paper. susceptible.

*Sex Transm Infect* 2002;**78**(5

Sexually Transmitted Diseases • December 2005

the potential for spread of infection as the number of new infections ca age when an infection enters an ent ble population.<sup>2</sup> Two things are worth noting from The basic reproduction number ( $R_0$ ) is fundamental to understanding infectious disease epidemiology.  $R_0$  describes the average number of secondary infections that an infected individual generates when entering a fully susceptible population, and when greater

## Secondary infections vs. secondarily infected individuals (cases)

- > Curable sexually transmitted infections (STI) → partners can re-infect each other



- > Number of secondary infections ( $R_0$ )  $\neq$  number of secondary cases ( $R_c$ )

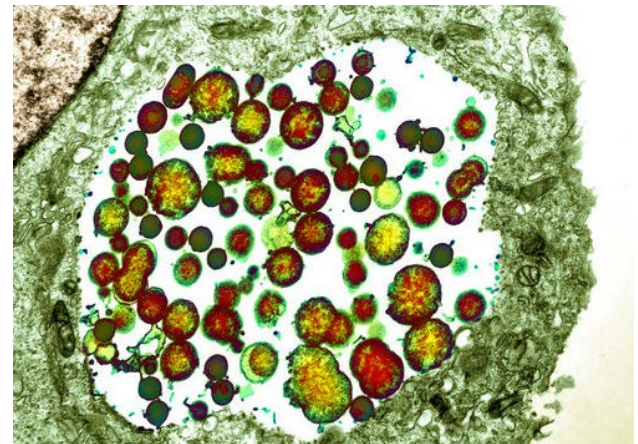


# Objectives

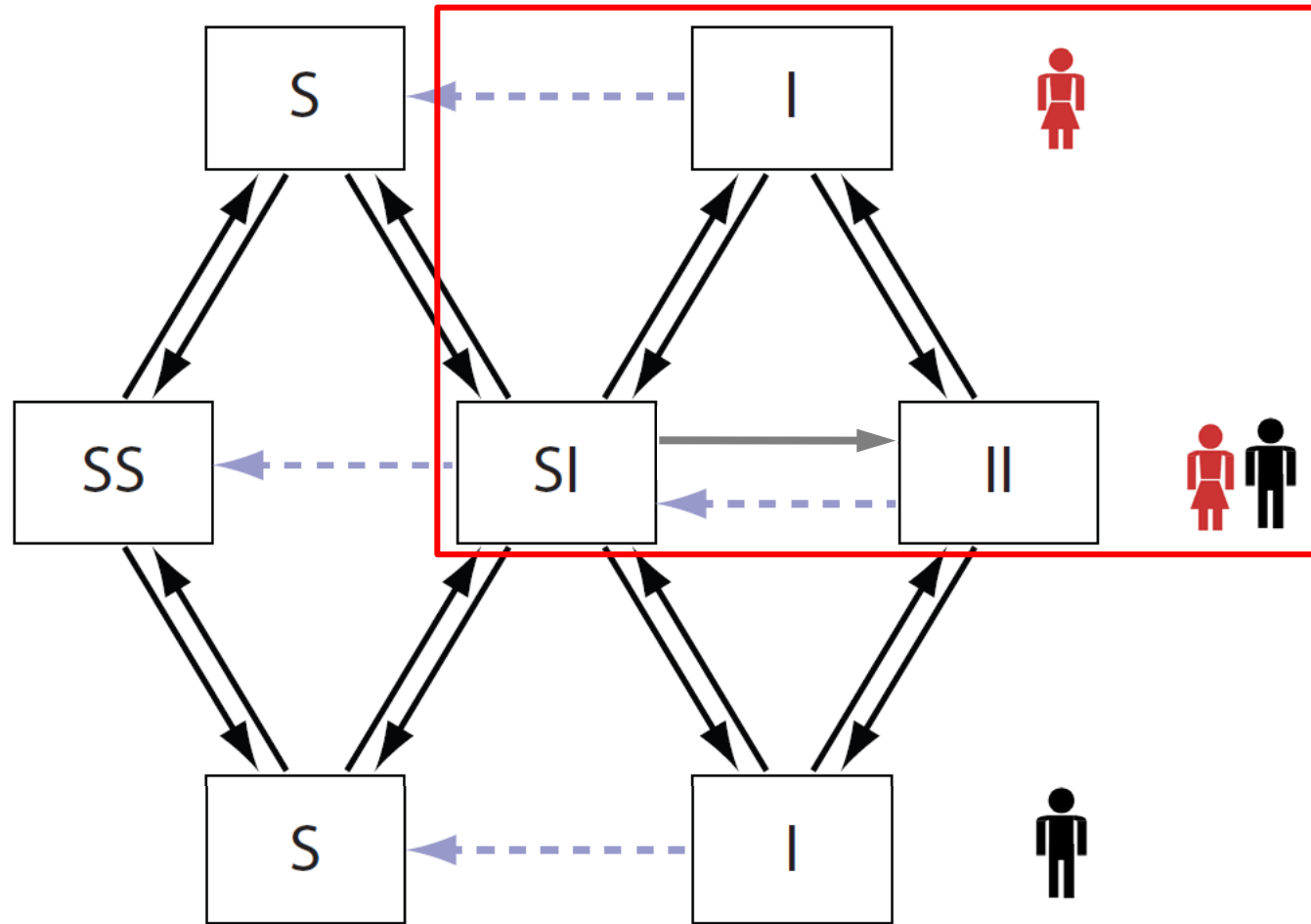
- > To obtain expressions for the number of secondary **infections** ( $R_0$ ) and the number of secondary **cases** ( $R_c$ ) for curable STI
- > To use  $R_0$  and  $R_c$  to investigate the contribution of re-infection within partnerships to endemicity of curable STI

# *Chlamydia trachomatis*

- > Most commonly notified STI in many developed countries
- > Duration of infection is long (on average one year)
- > Mostly asymptomatic
- > People can clear the infection naturally
- > Antibiotic treatment is effective
- > Long term complications

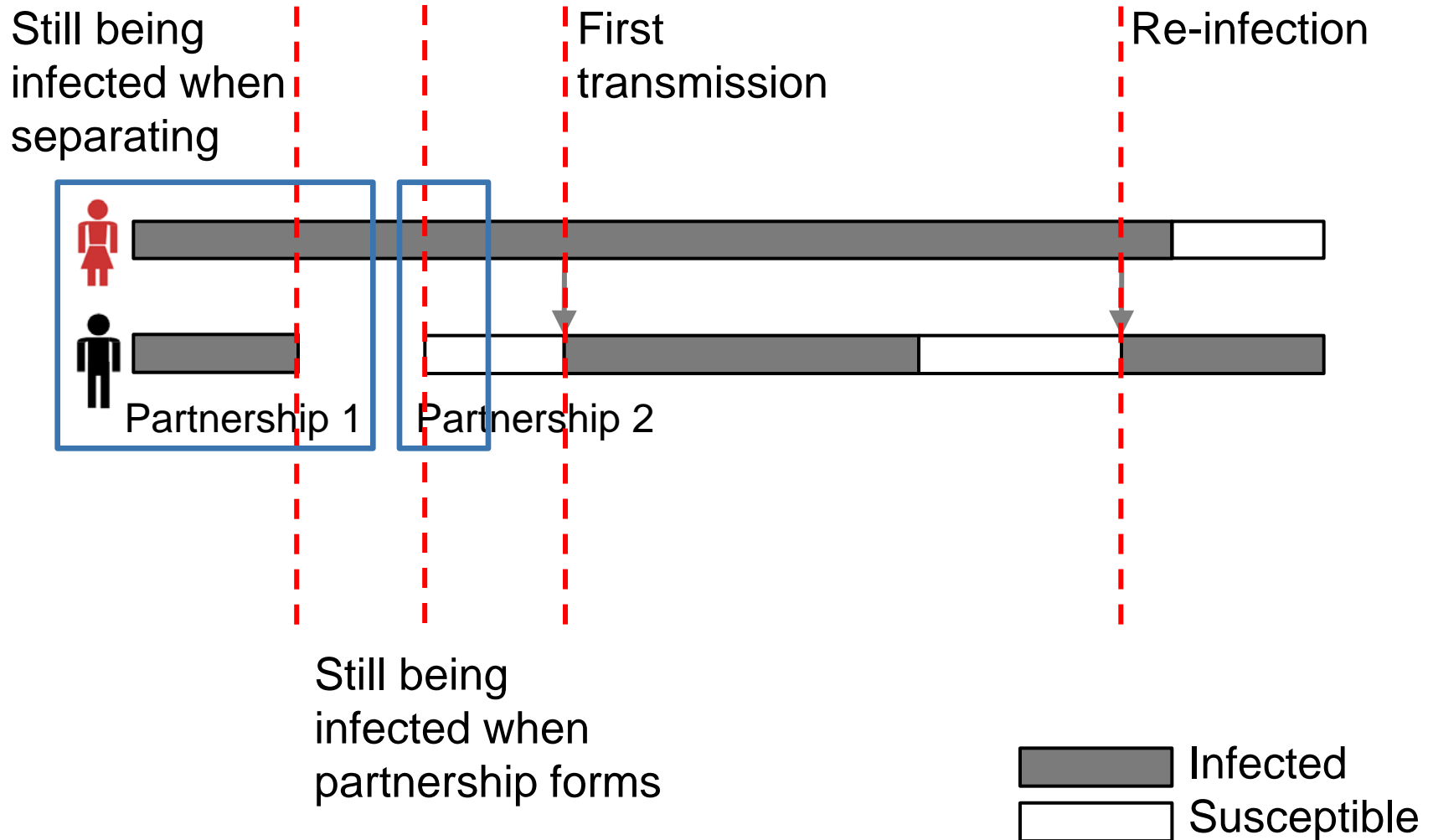


# Pair model



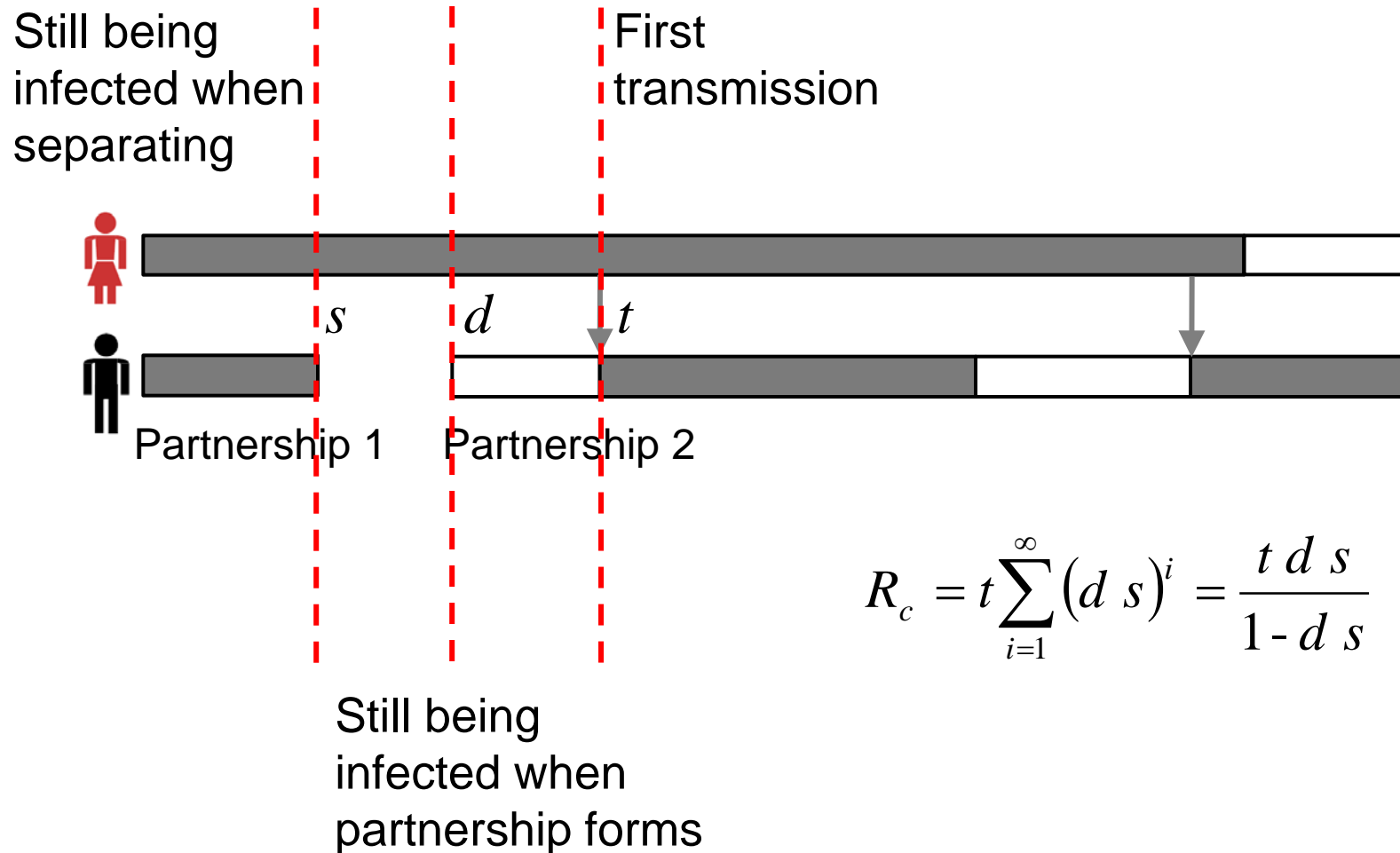
→ Pair formation/ separation rate      - - - - -> Recovery by natural clearance  
→ Transmission within partnership

# Assumptions and probabilities

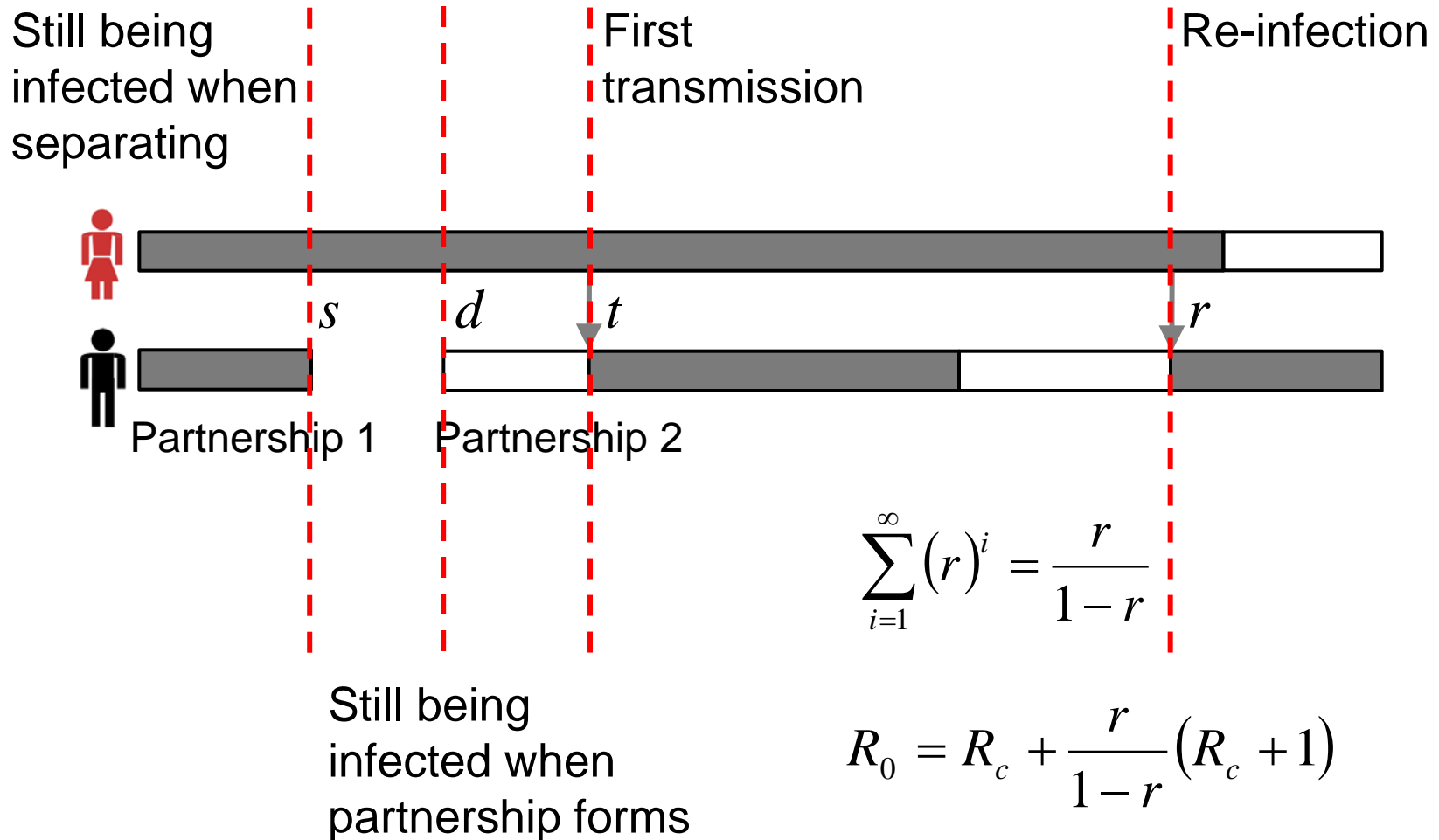




# Case reproduction number

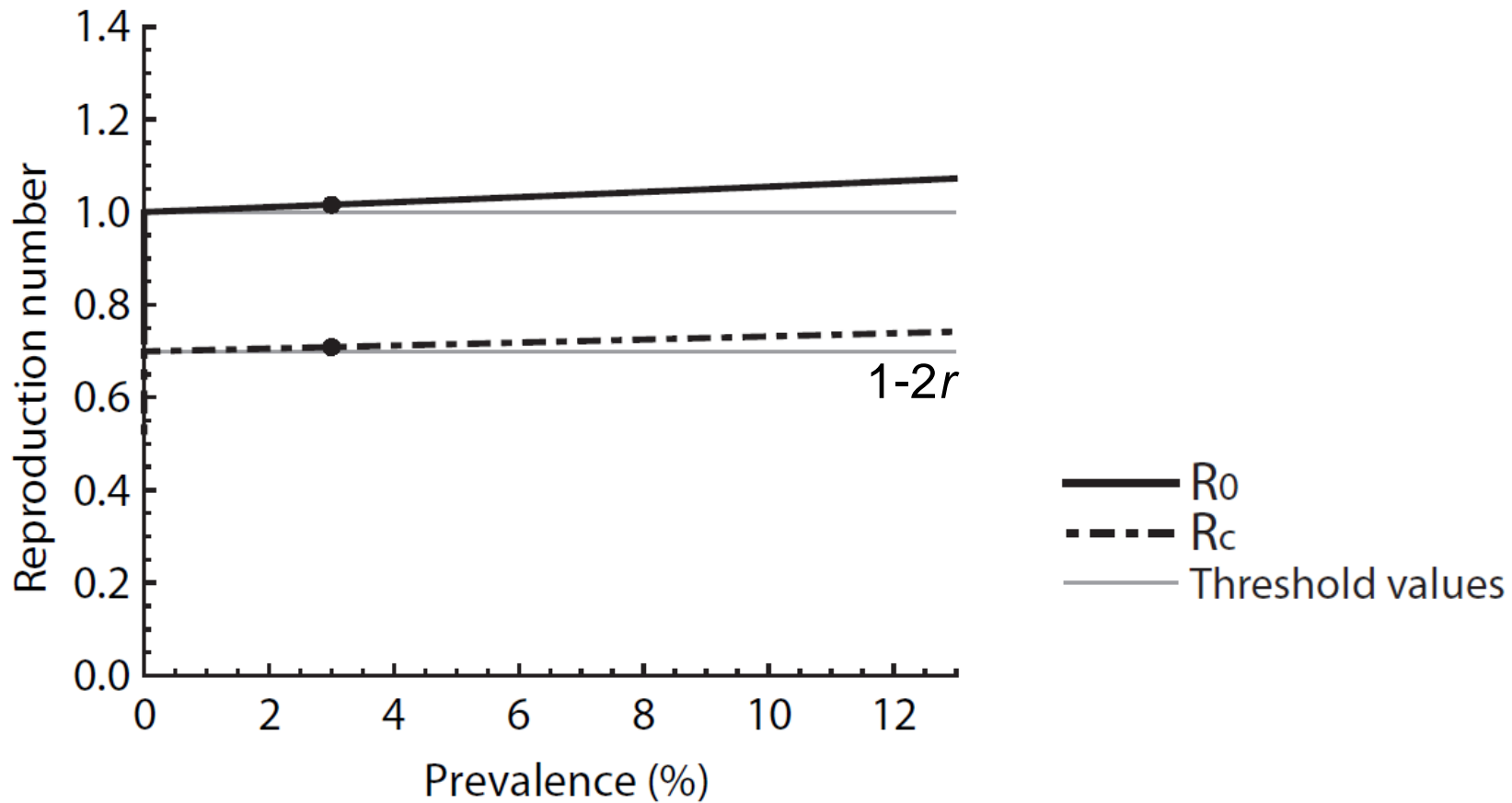


# Basic reproduction number



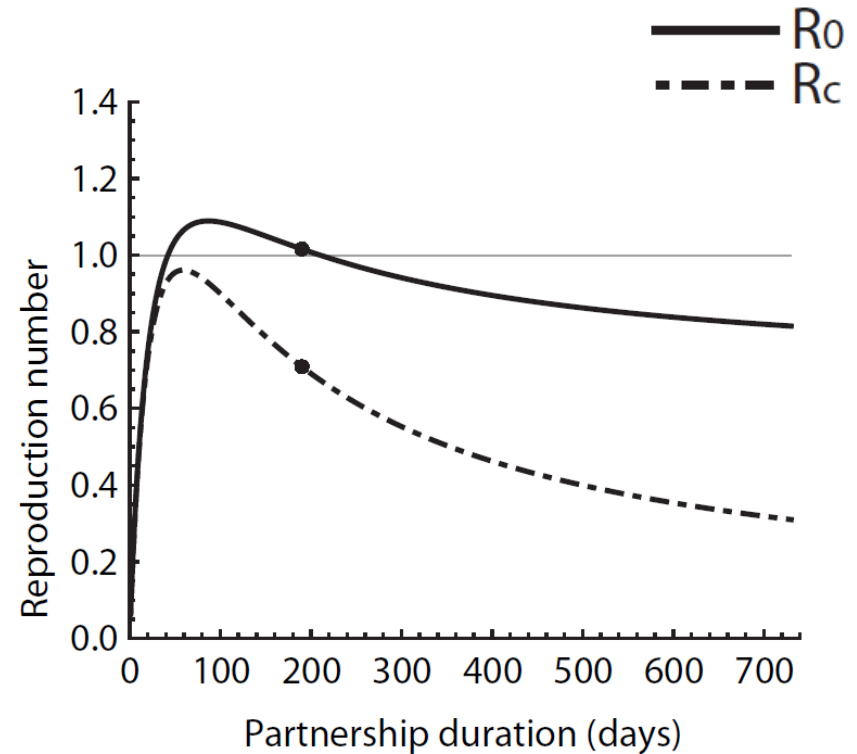
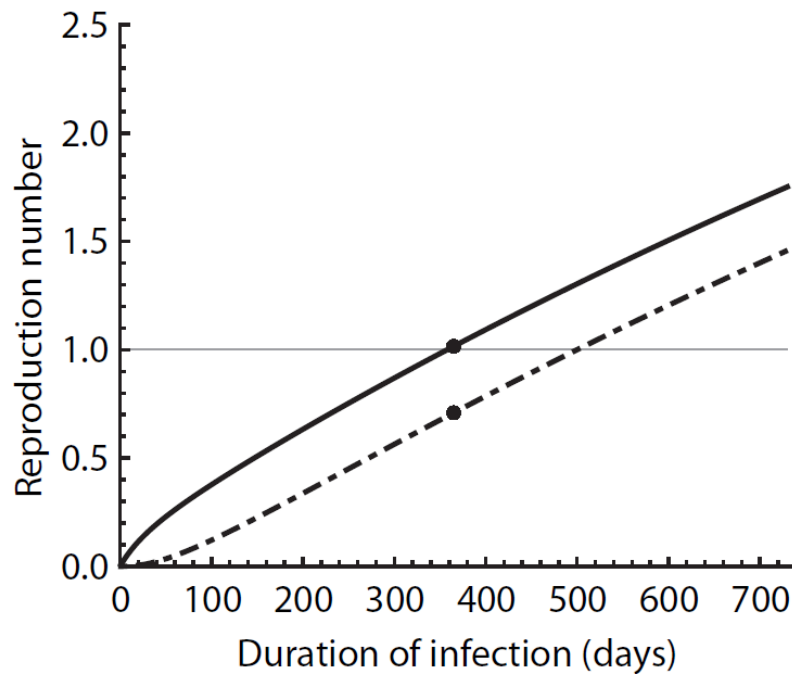
# Results

## Epidemic threshold levels



# Results

## Duration of infection and partnerships



# Strengths and limitations

- > Strength:
  - Simple Model
- > Limitations:
  - Homogeneous mixing
  - No immunity

# Discussion

- > Applicability to other infections:
  - Infections that are curable and transmitted within households
  - Hospital infections
- > Estimating reproduction numbers from other study designs such as contact tracing data

# Conclusions

- > The precise meaning of reproduction numbers and how they incorporate re-infection should be stated clearly by modellers
- > Re-infection within partnerships means that curable STI can be sustained endemically even when the number of secondary cases ( $R_c$ ) is below one

# Acknowledgments

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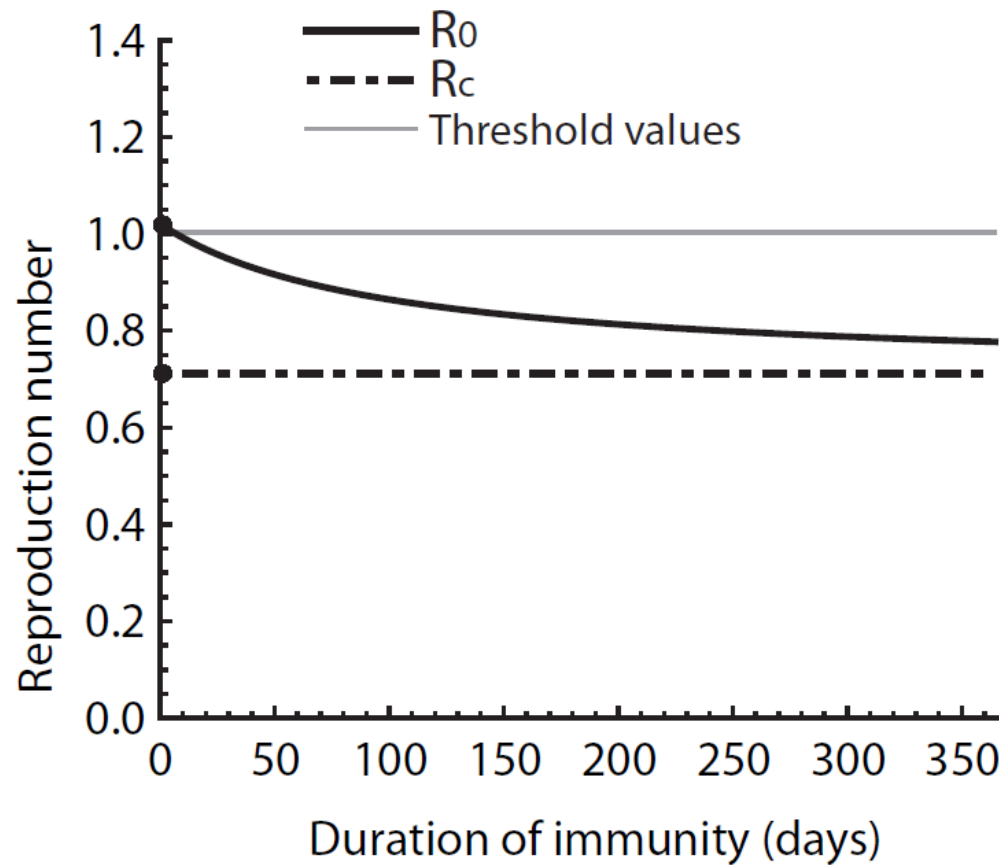


# Reproduction numbers

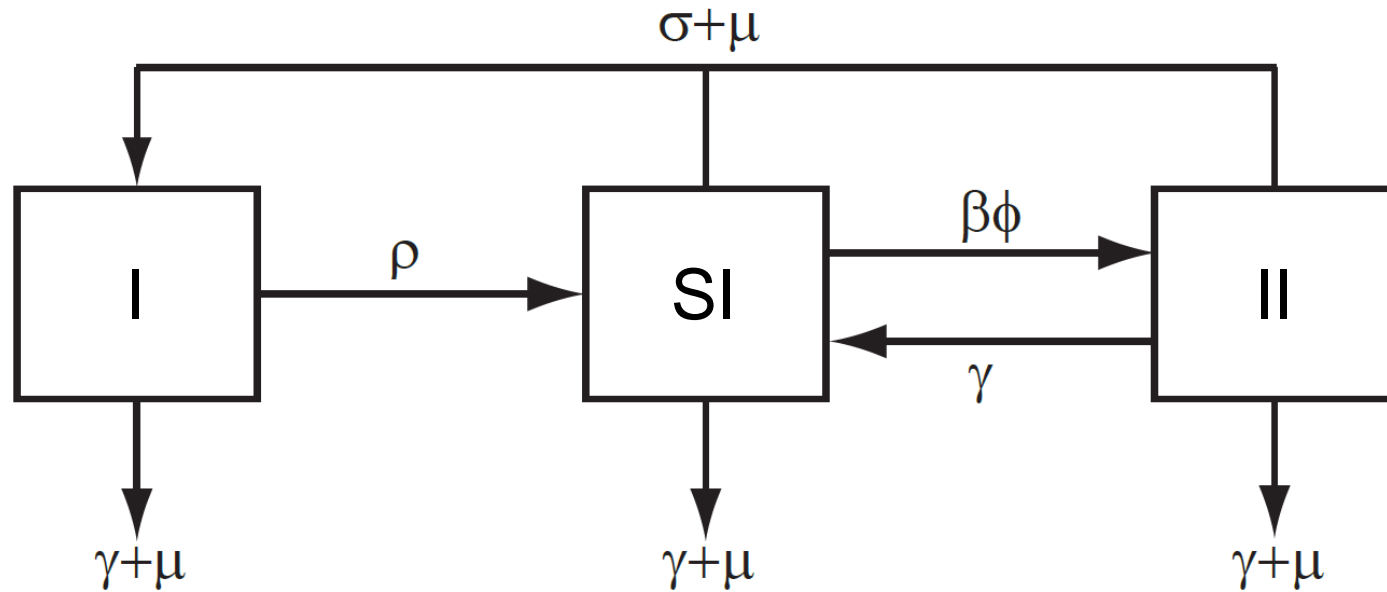
$$R_0 = \frac{\beta\phi(\gamma^2 + \gamma(\mu + \rho) + \rho(\sigma + \mu))}{(\gamma + \mu)(\sigma + 2\mu + 2\gamma + \beta\phi)(\sigma + 2\mu + \gamma + \rho)}$$

$$R_c = \frac{\beta\phi \rho(\sigma + \mu)}{(\gamma + \mu)(\sigma + 2\mu + \gamma + \beta\phi)(\sigma + 2\mu + \gamma + \rho)}$$

# Duration of immunity



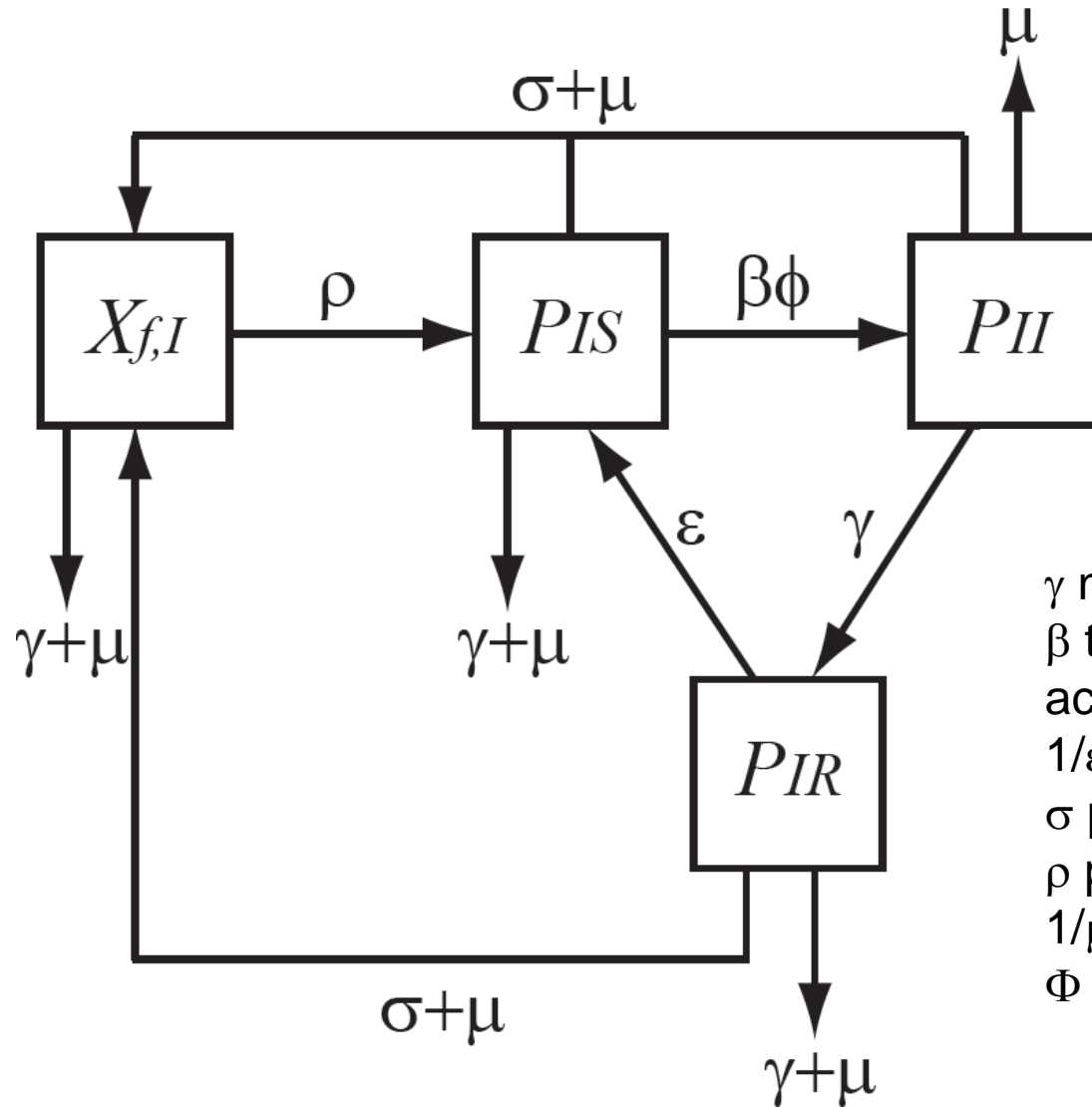
# Pair model, infected women



|            |                                      |
|------------|--------------------------------------|
| $1/\gamma$ | duration of infection                |
| $\sigma$   | pair separation rate                 |
| $\rho$     | pair formation rate                  |
| $1/\mu$    | duration of sexually active life     |
| $\beta$    | transmission probability per sex act |
| $\Phi$     | frequency of sex acts                |

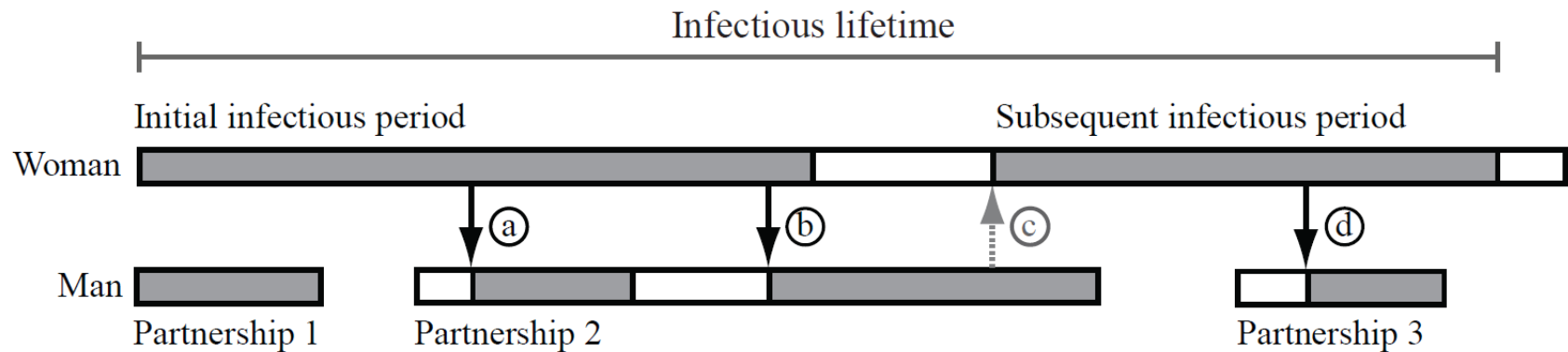
S      Susceptible  
I      Infected

# Pair model



$\gamma$  natural clearance rate  
 $\beta$  transmission probability per sex act  
 $1/\varepsilon$  duration of immunity  
 $\sigma$  pair separation rate  
 $\rho$  pair formation rate  
 $1/\mu$  duration of sexual active live  
 $\Phi$  Frequency of sex acts

# Partnership reproduction number ( $R_p$ )



# Partnership reproduction number

