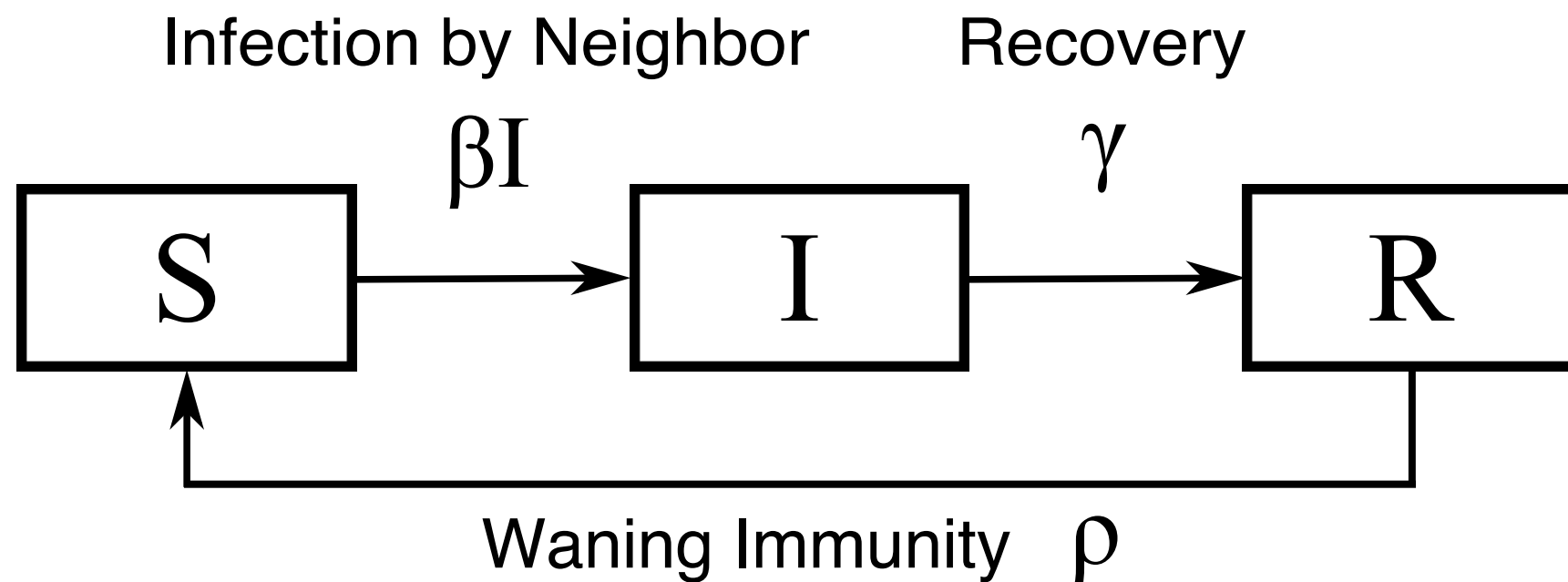


Contact Network Heterogeneity and Persistence of Endemic Disease

Daniel T. Citron
Christopher R. Myers
Department of Physics
Cornell University

Endemic Infection

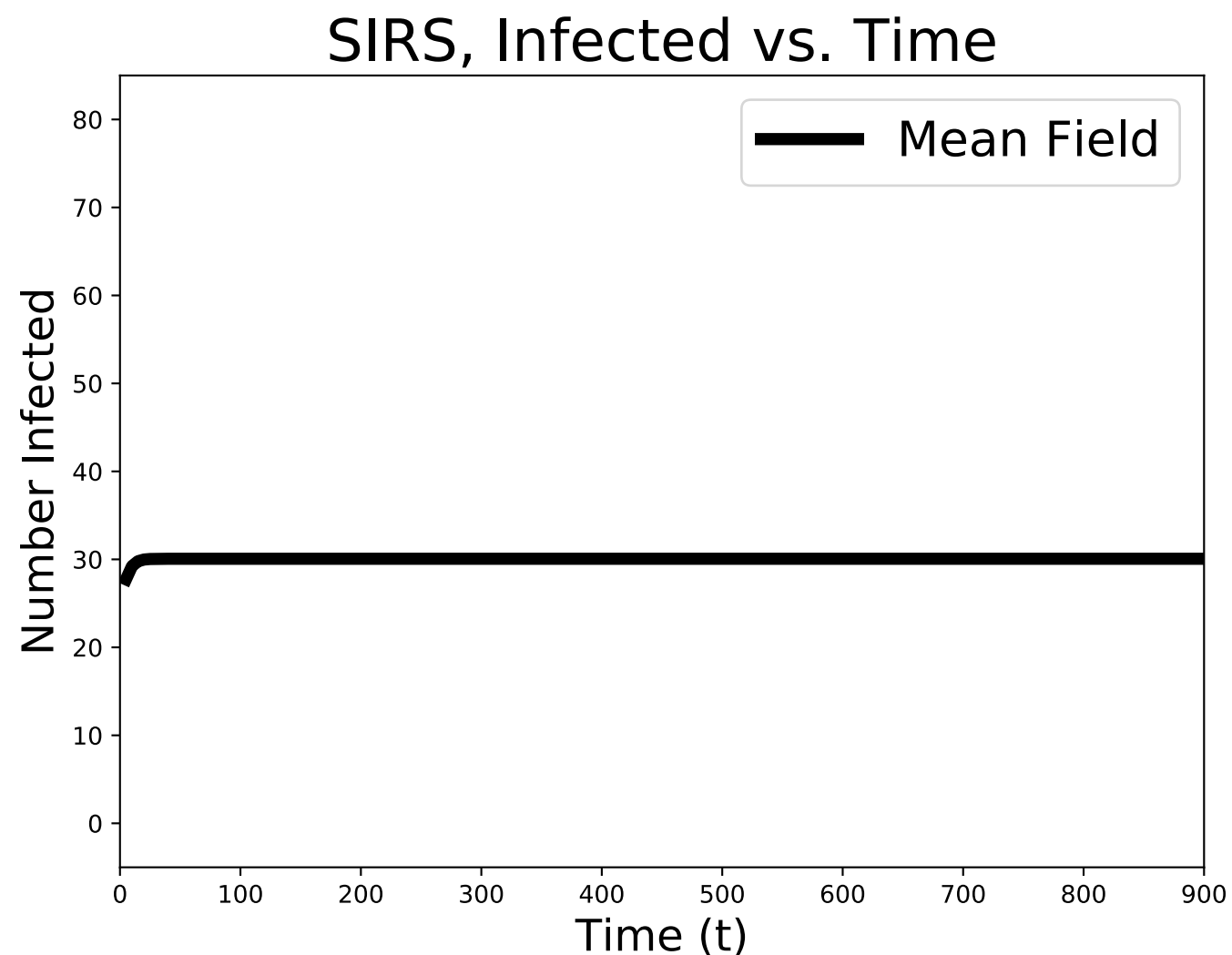
- Susceptibles are replenished
- Example: SIR with waning immunity



- Persistence of infected individuals in a population

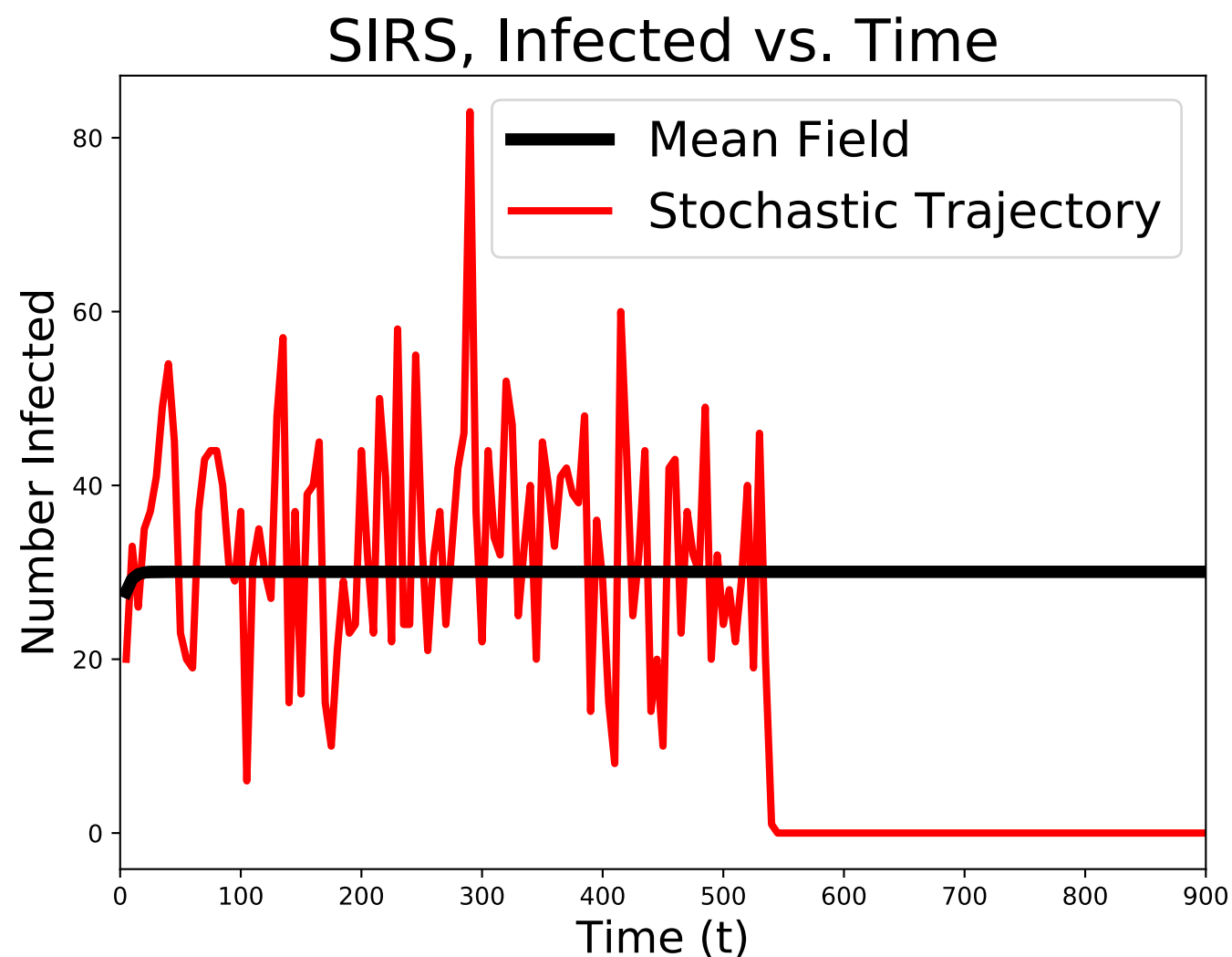
Spontaneous Extinctions

- Endemic states not stable in finite populations



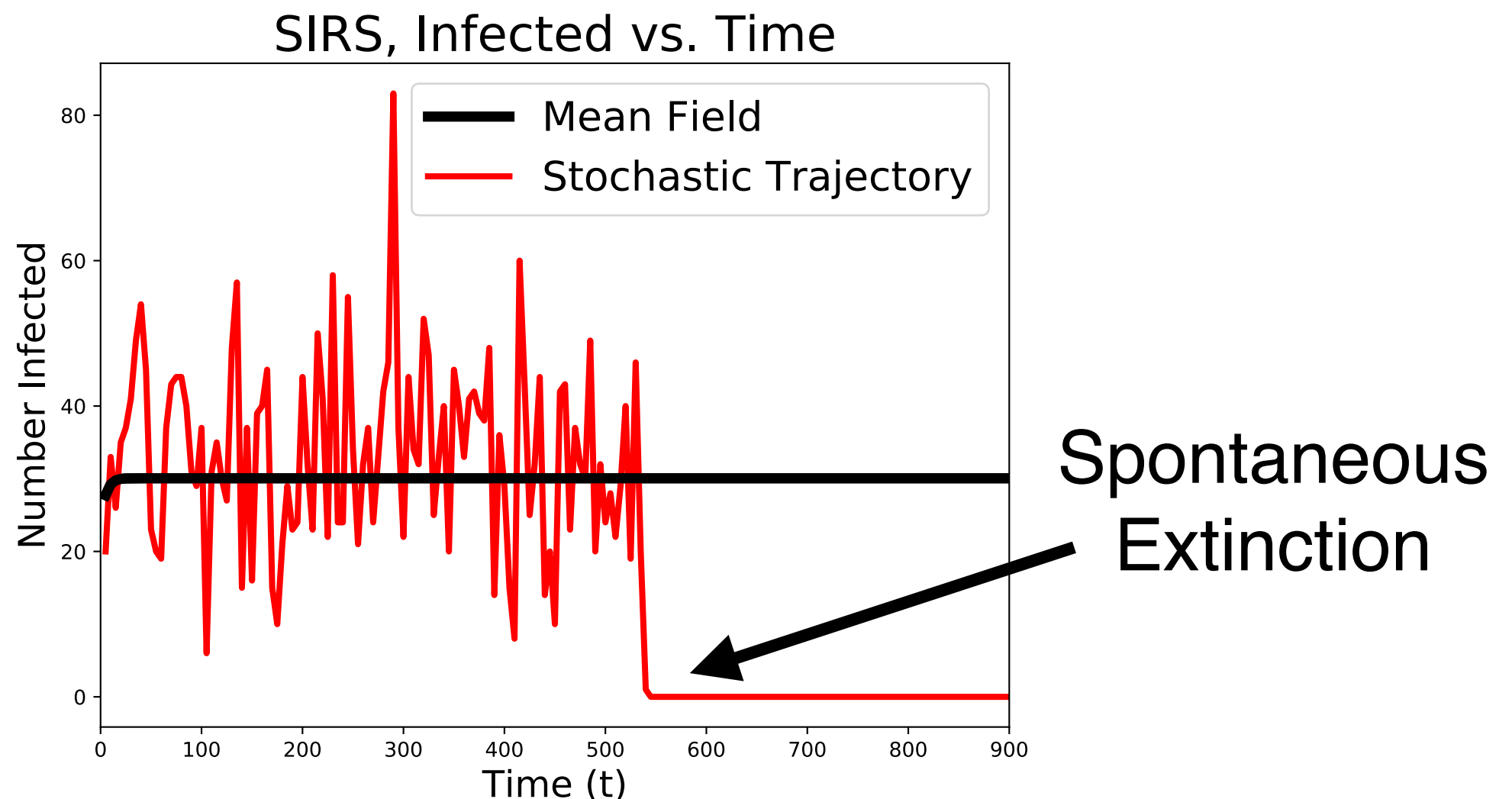
Spontaneous Extinctions

- Endemic states not stable in finite populations
- Stochastic fluctuations bring infection level to 0



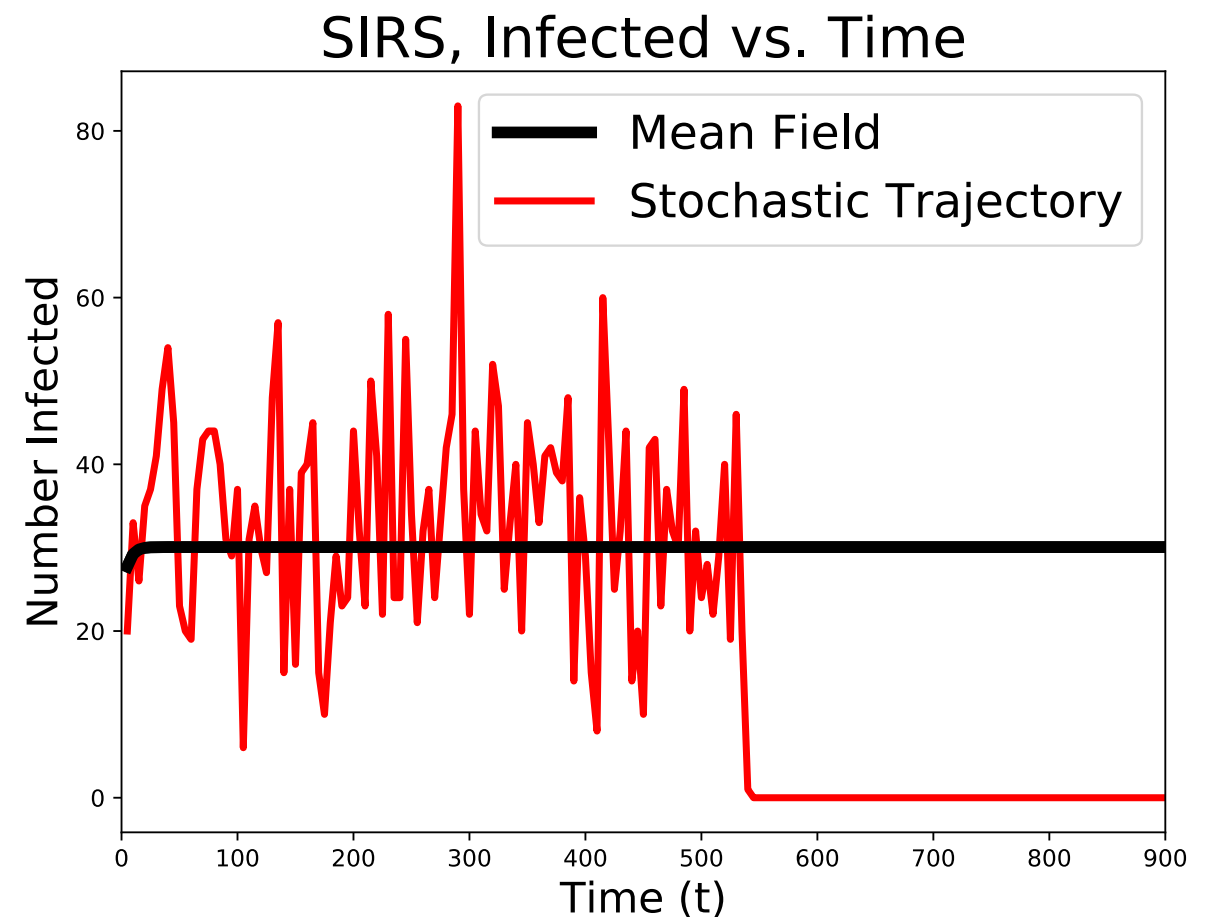
Spontaneous Extinctions

- Endemic states not stable in finite populations
- Stochastic fluctuations bring infection level to 0



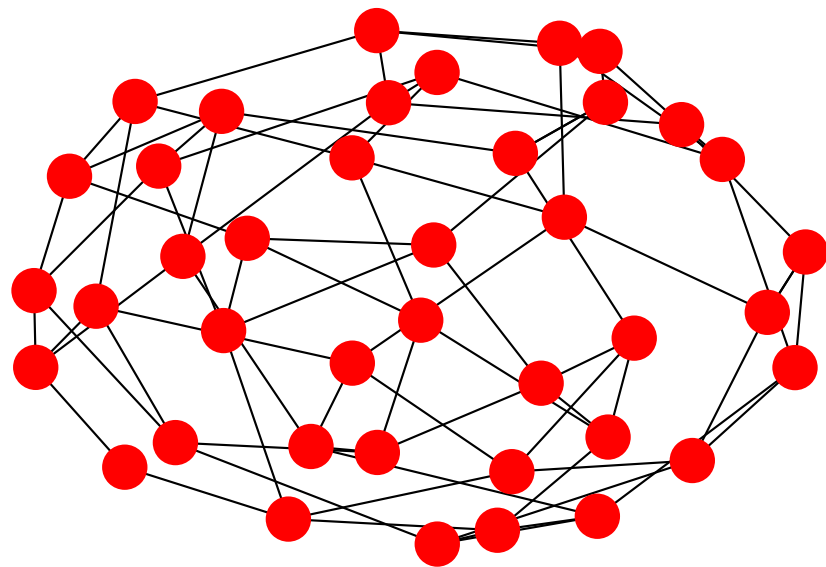
Endemic State Lifetimes

- Characteristic Lifetime
- Depends on
 - Mean Infection μ
 - Fluctuation Size σ

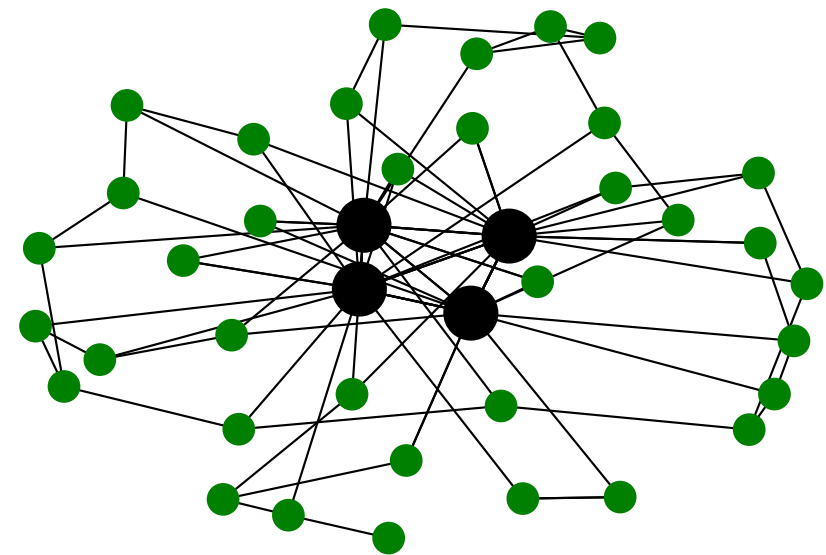


Network Effects

- Change contact network topology
- What happens to the endemic state lifetime?



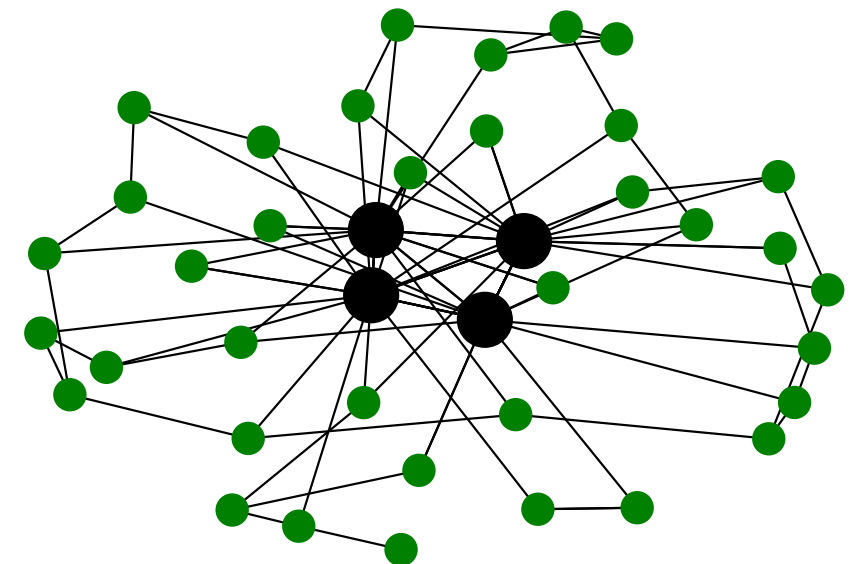
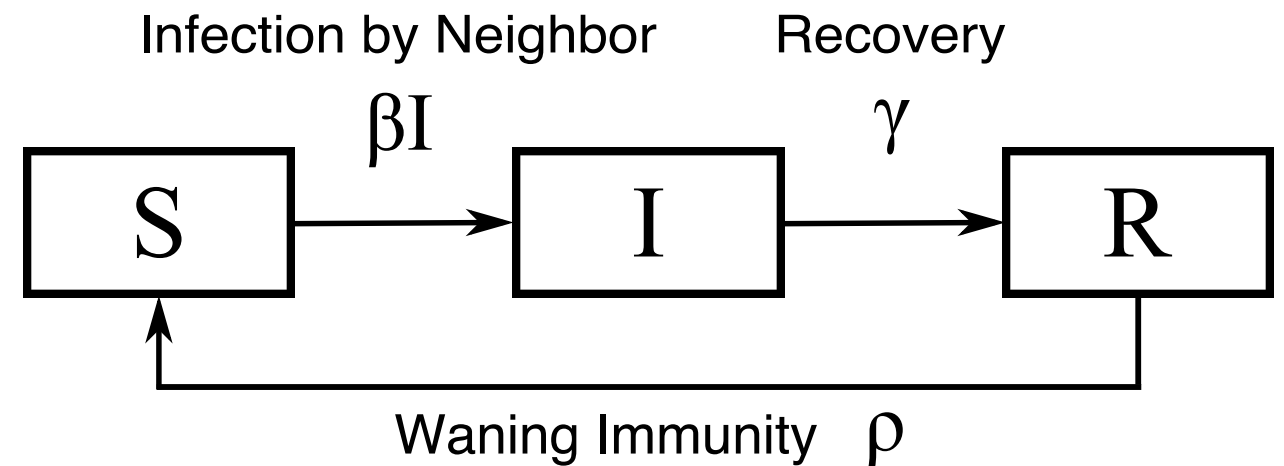
Homogeneous Graph



Heterogeneous Graph

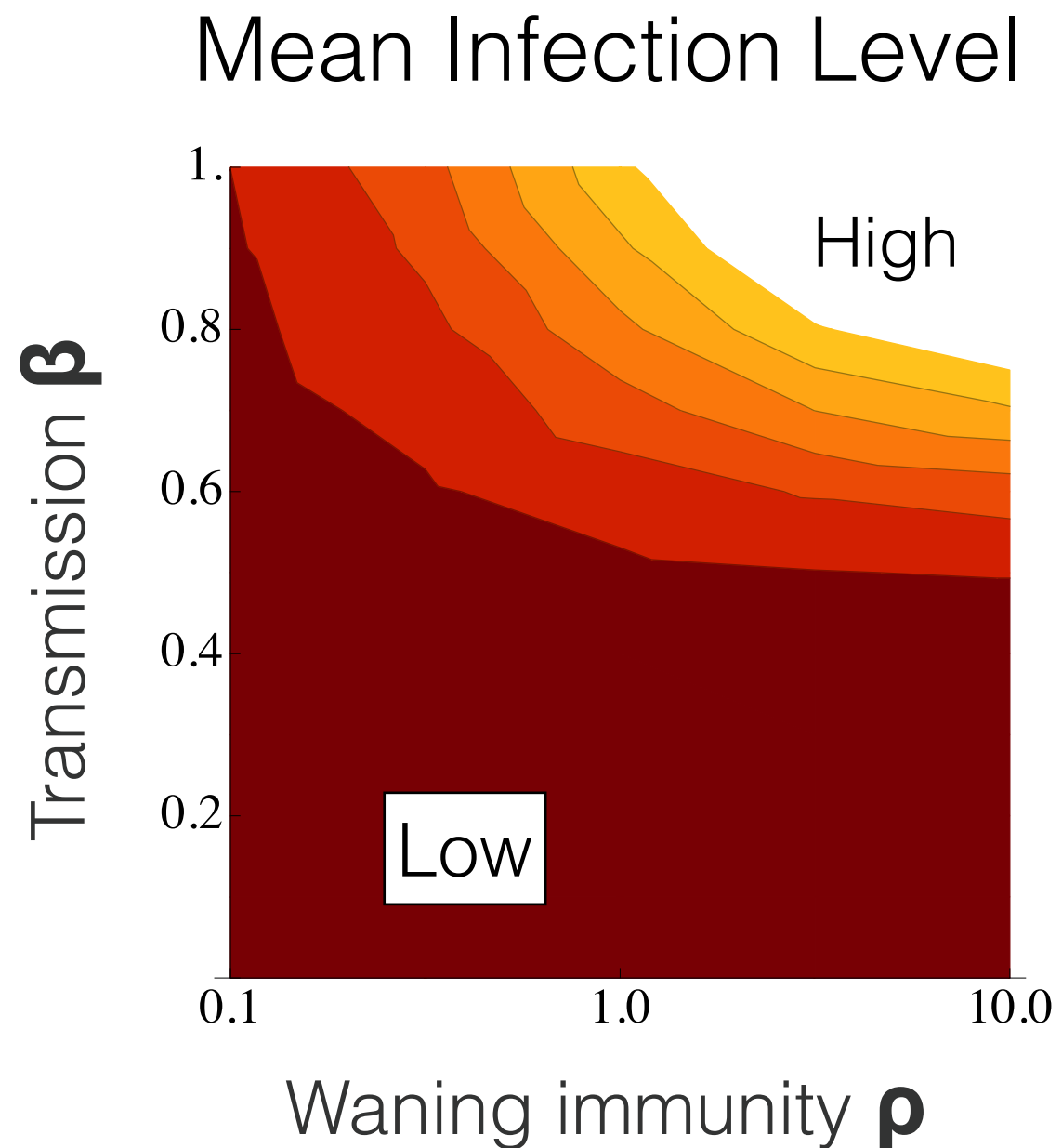
Our Simulations

- SIRS model
- Annealed networks
- Vary degree heterogeneity $\langle \sigma_k \rangle$
- Generate ensembles of trajectories
- Measure lifetime of endemic state



Results

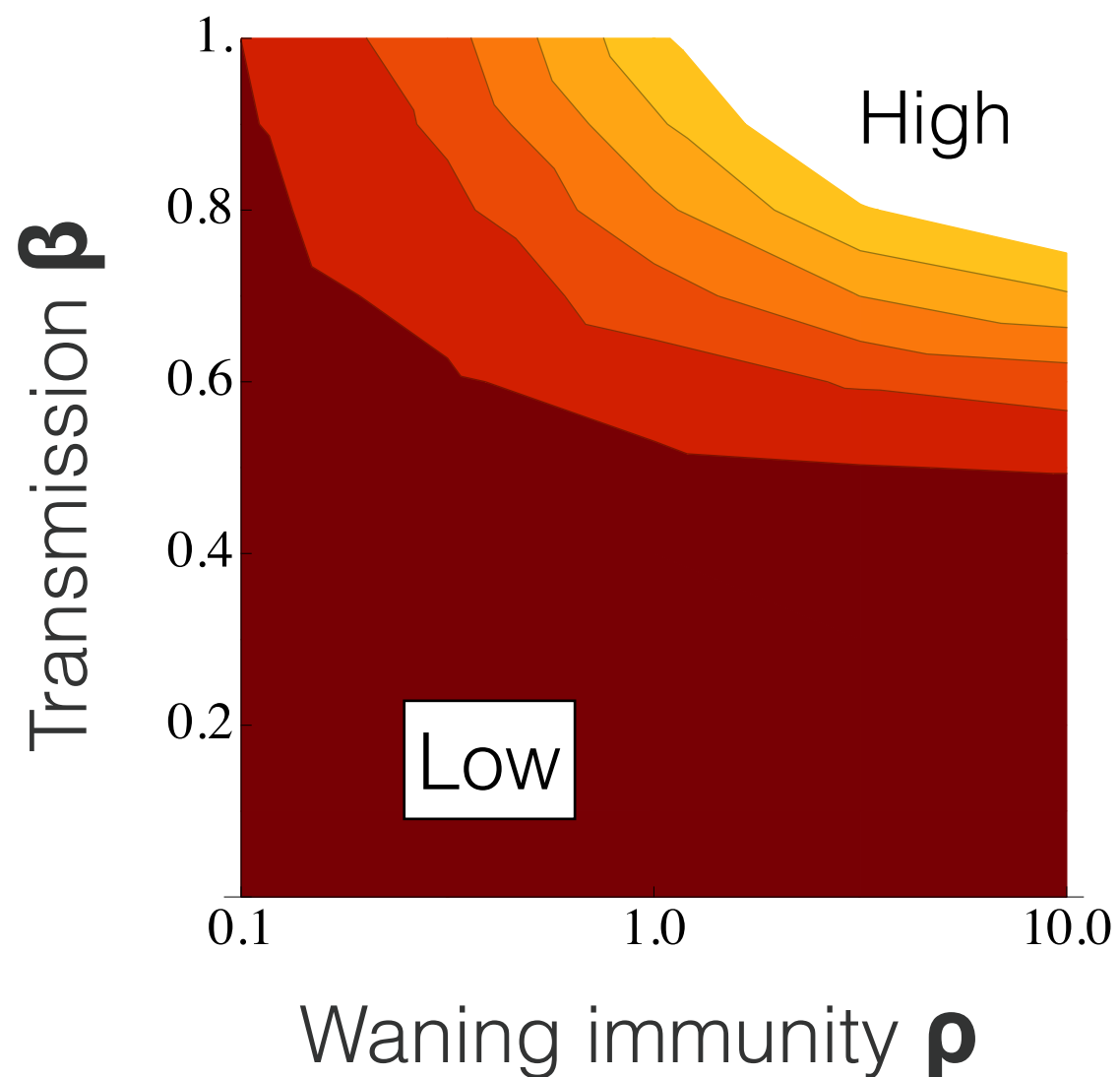
500 nodes, mean degree 10, $\langle \sigma_k \rangle = 10$



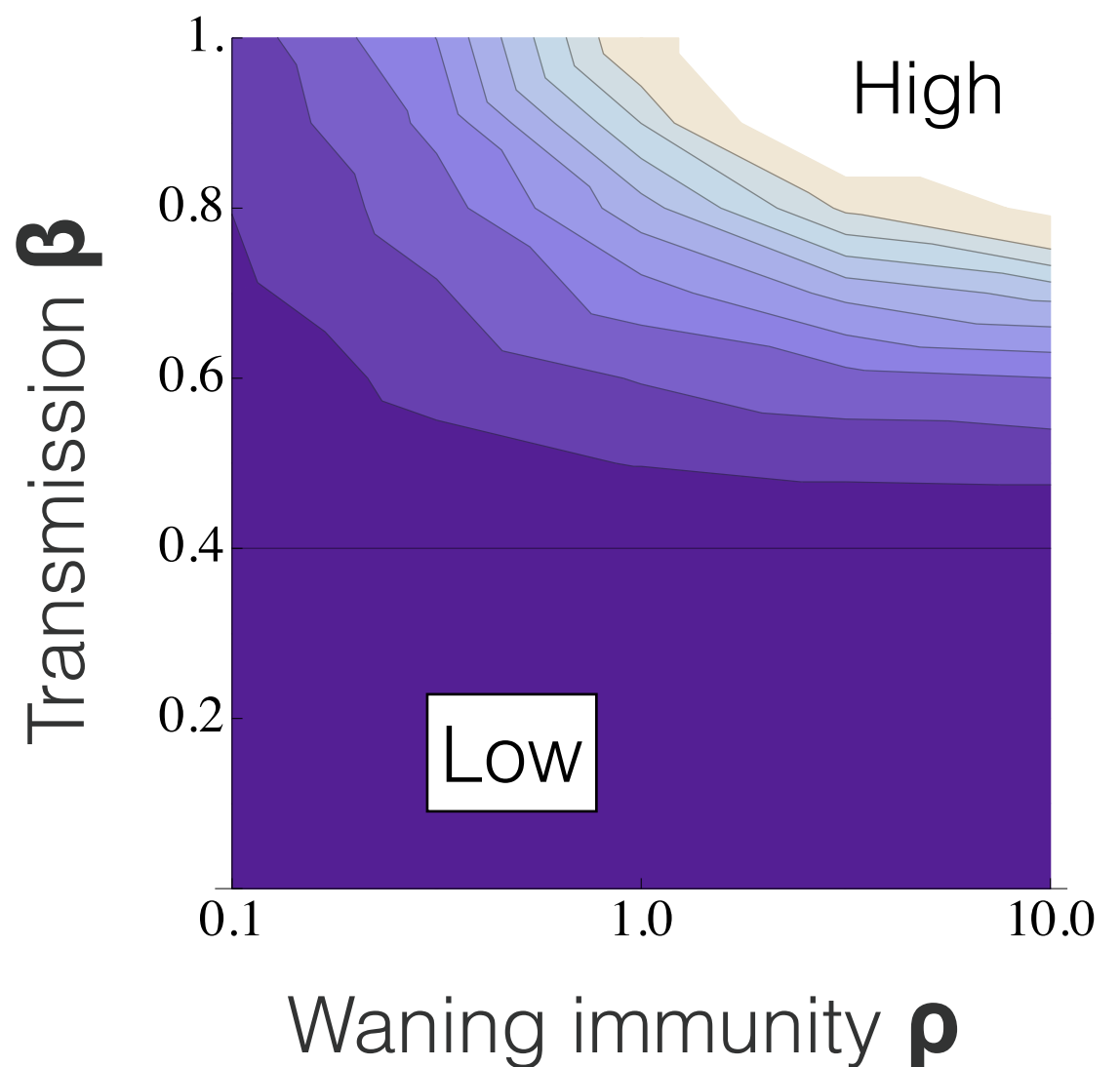
Results

500 nodes, mean degree 10, $\langle \sigma_k \rangle = 10$

Mean Infection Level



Mean Time to Extinction



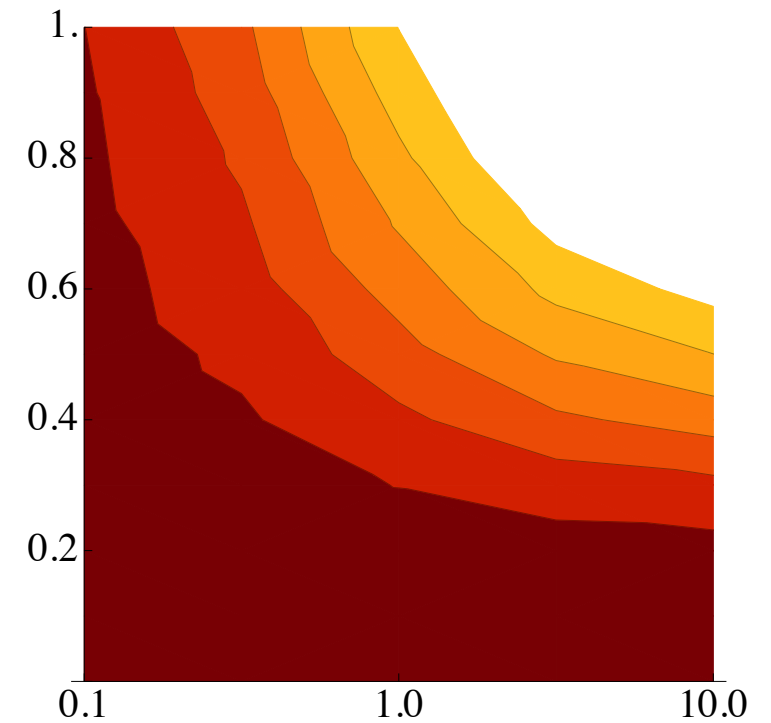
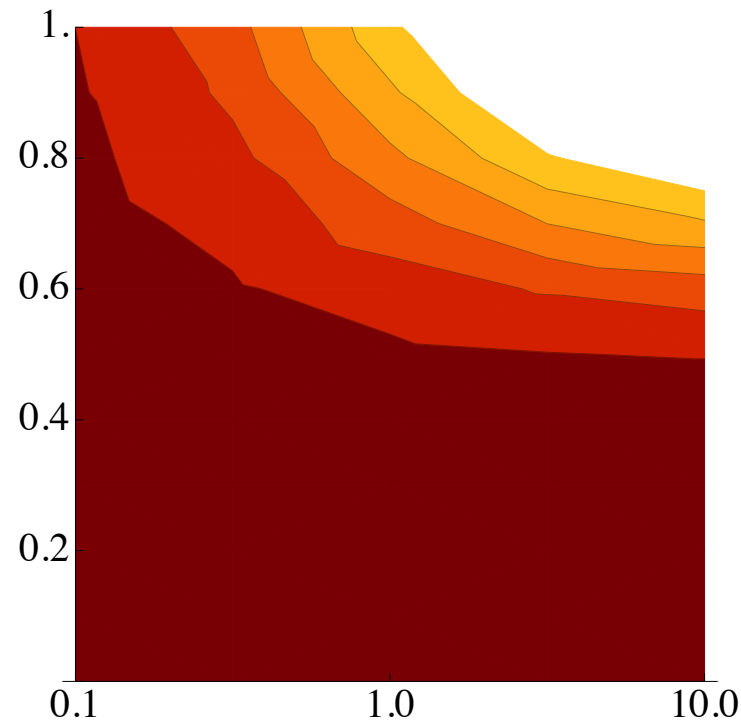
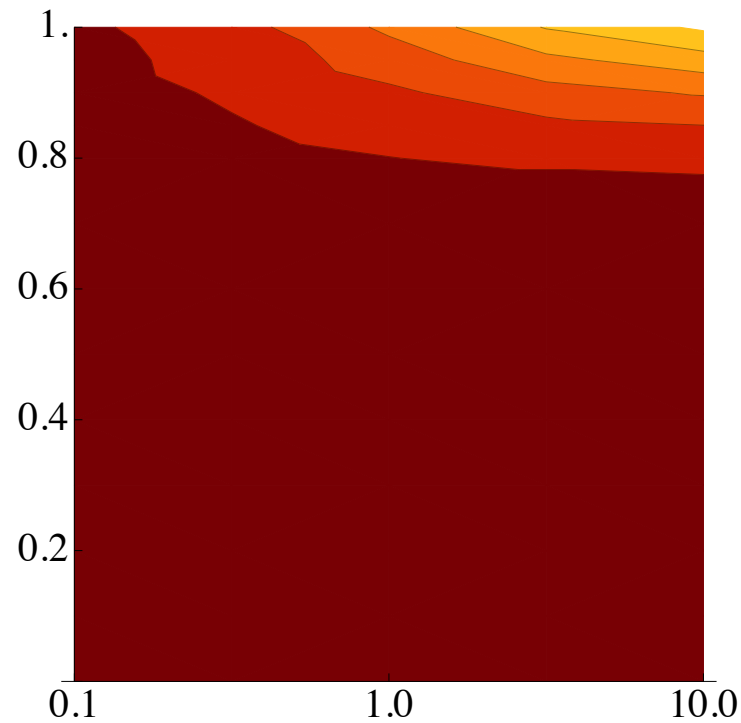
Results: Across Graphs

$\langle \sigma_k \rangle = 5$
(Homogeneous)

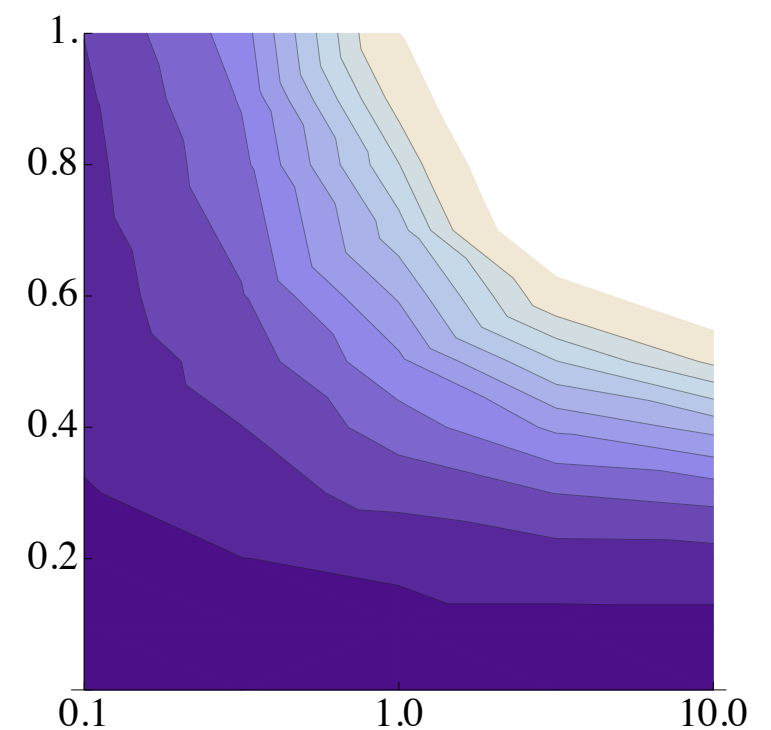
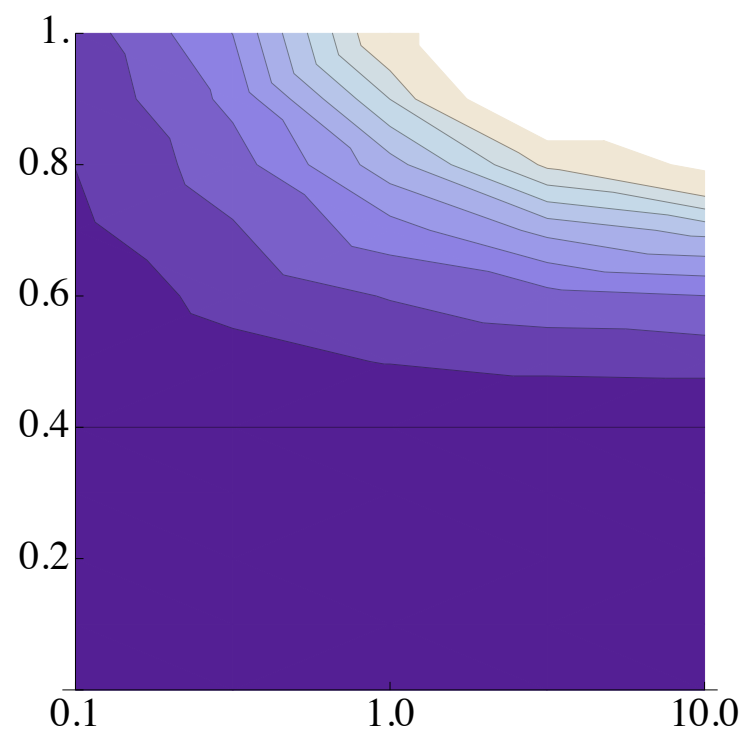
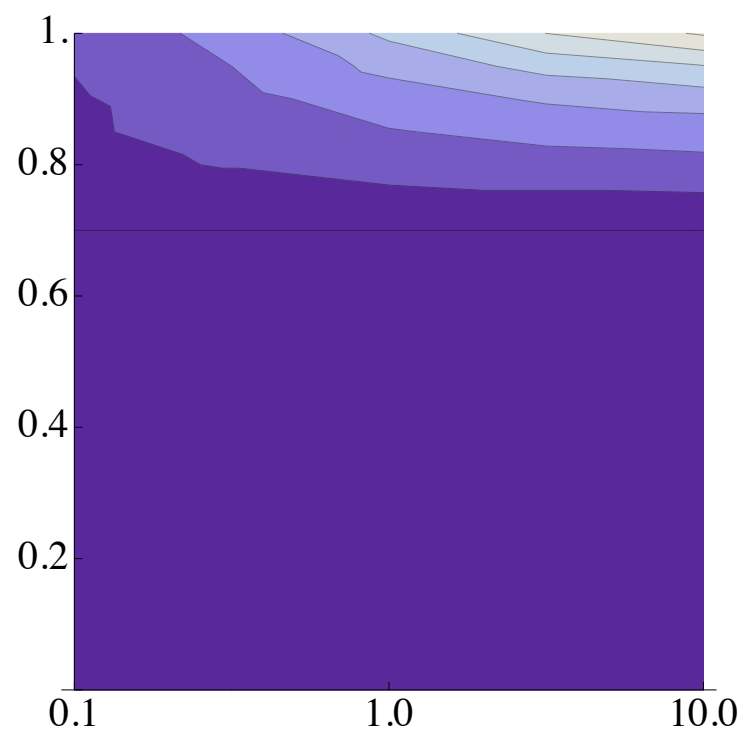
$\langle \sigma_k \rangle = 10$

$\langle \sigma_k \rangle = 20$
(Heterogeneous)

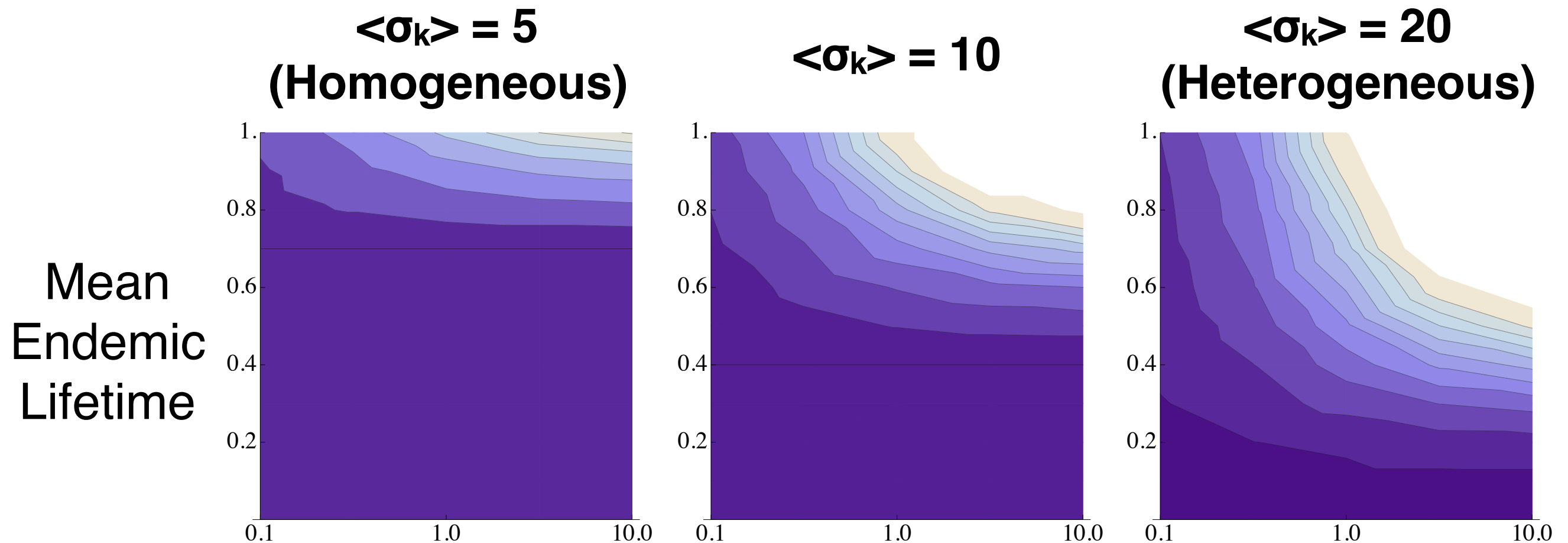
Mean
Number
Infected



Mean
Endemic
Lifetime



Results: Across Graphs



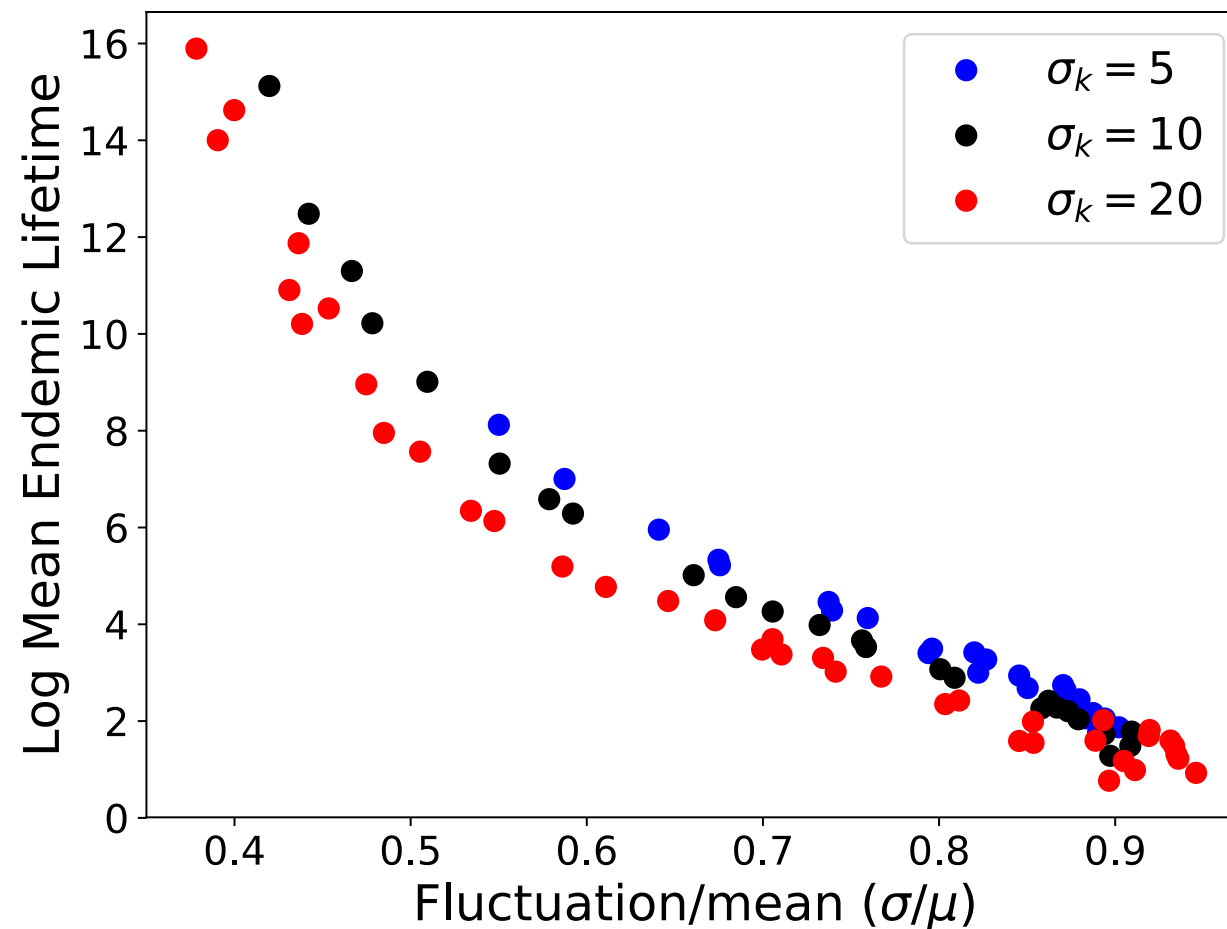
- Heterogeneous graphs have longer persistence
- Endemic lifetimes depend on topology

Predicting Lifetimes

- Can we predict endemic state lifetimes?

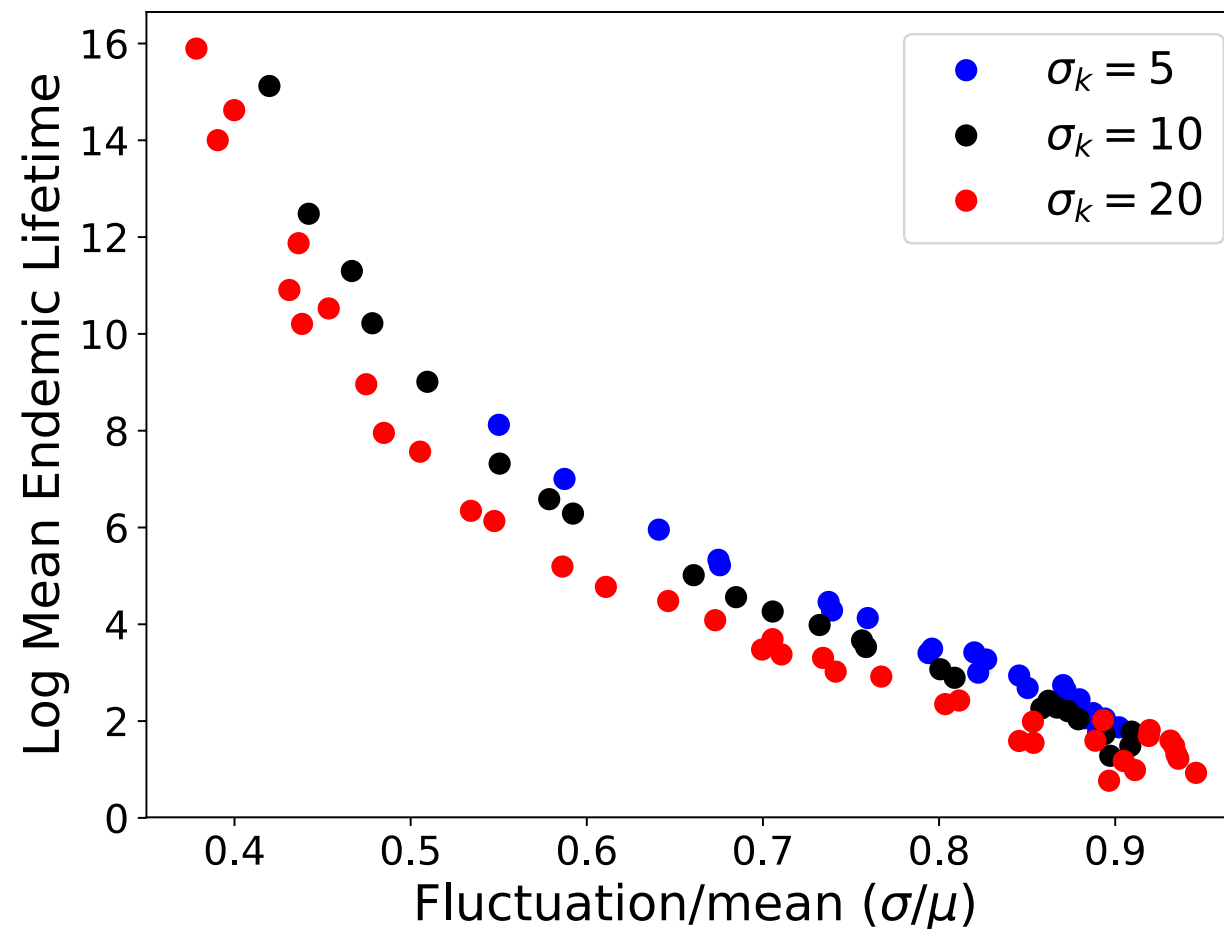
Predicting Lifetimes

- Can we predict endemic state lifetimes?



Predicting Lifetimes

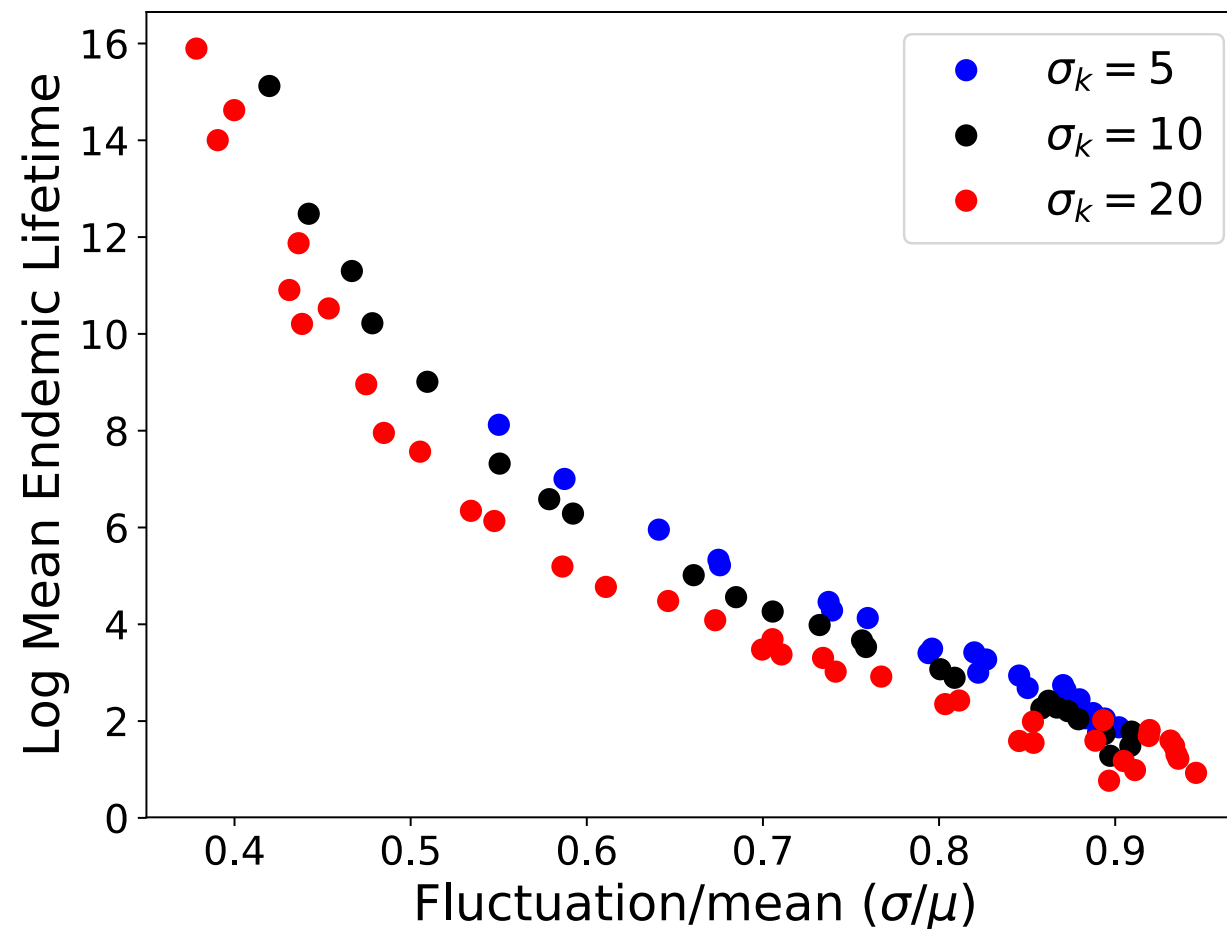
- Can we predict endemic state lifetimes?



- Fluctuation/mean ratio σ/μ is sufficient

Predicting Lifetimes

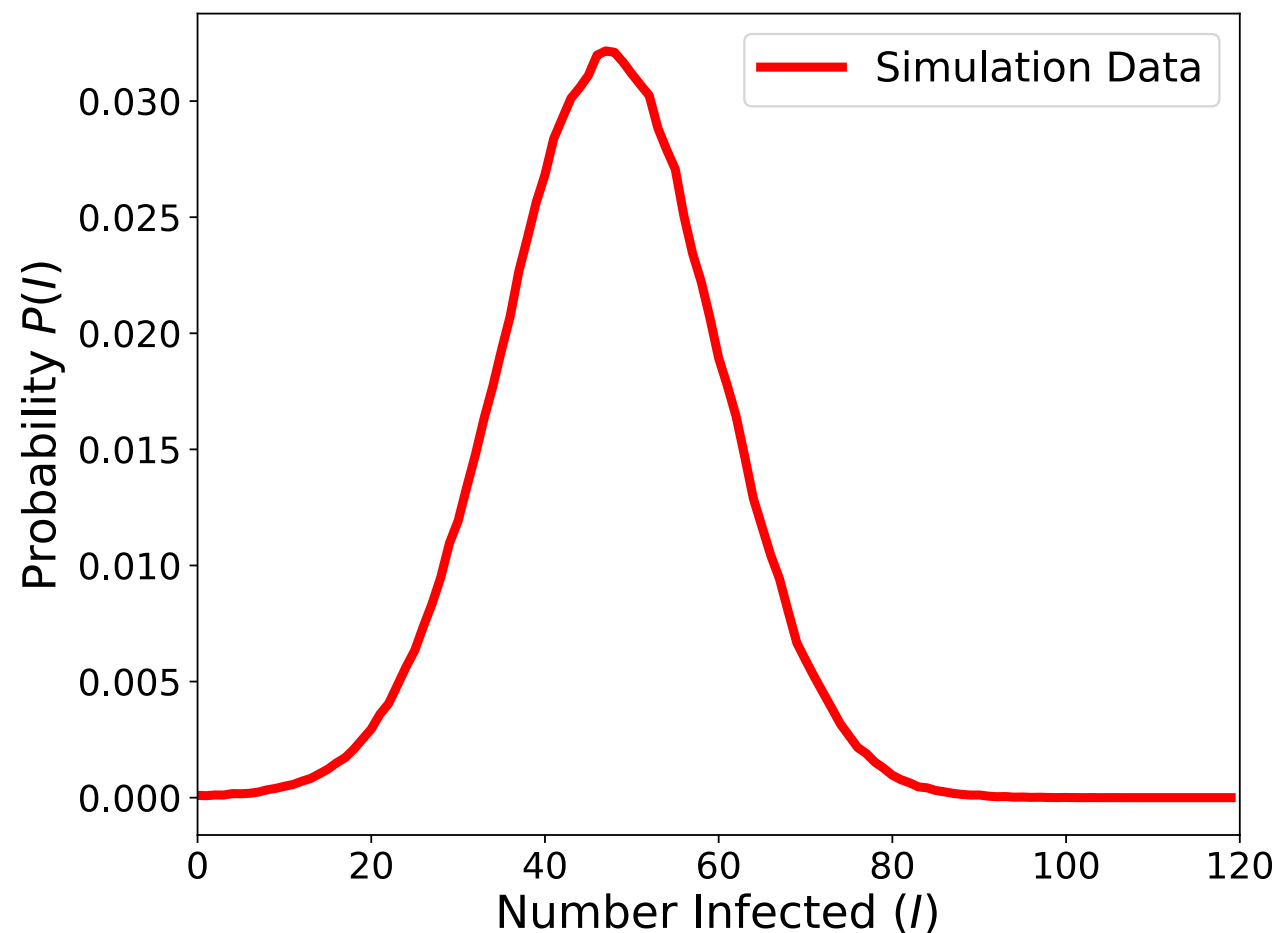
- Can we predict endemic state lifetimes?



- Fluctuation/mean ratio σ/μ is sufficient
- Corrections from changing topology

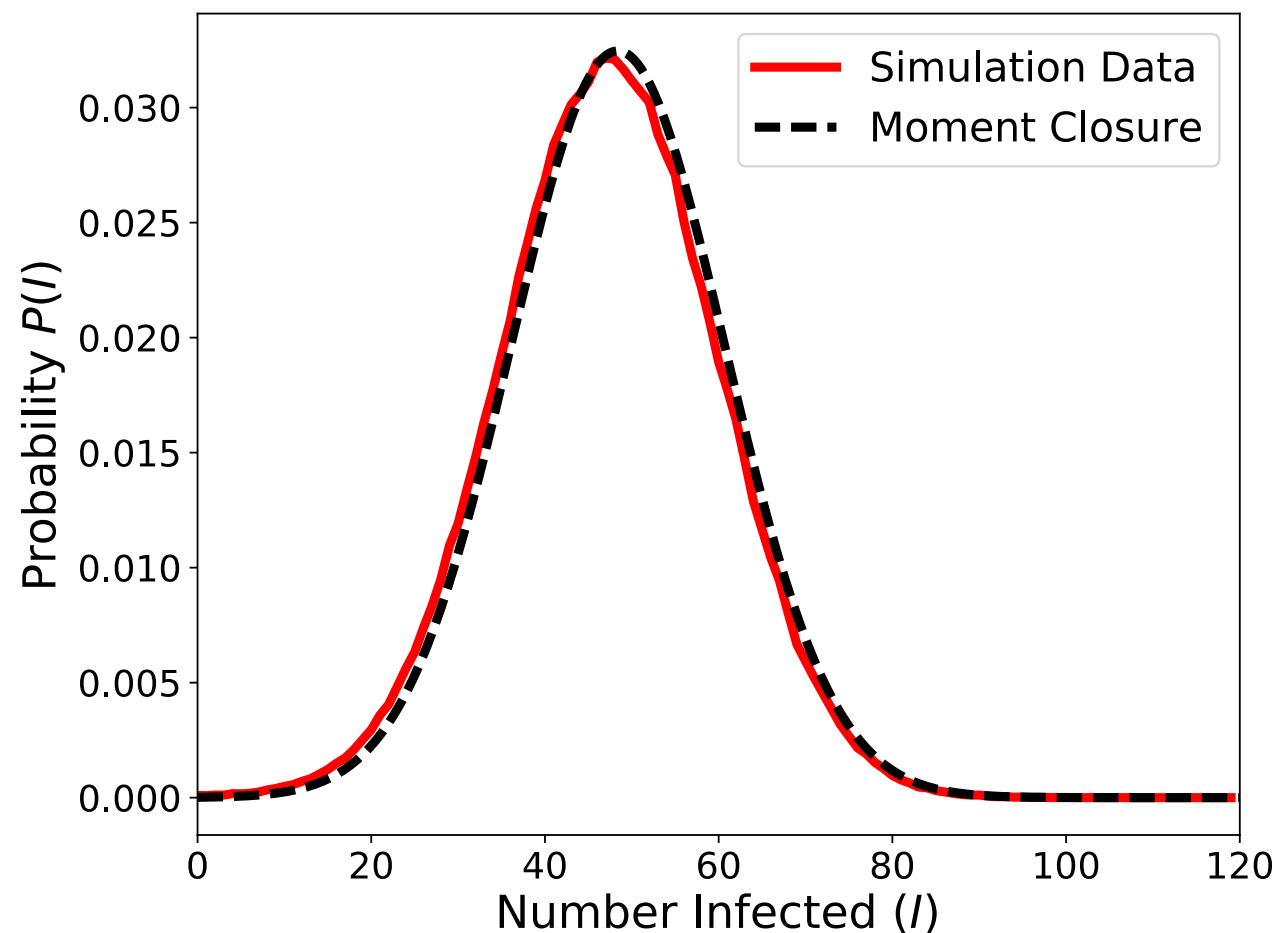
Moment Closure

- Analytical technique that predicts *both* means and fluctuation sizes



Moment Closure

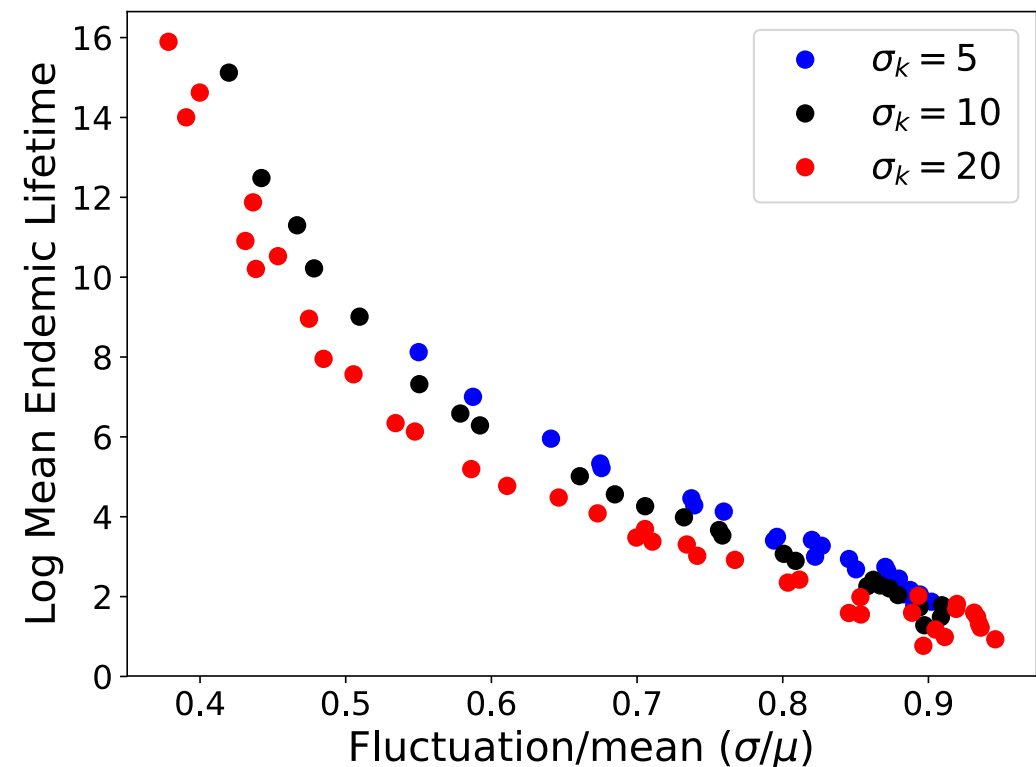
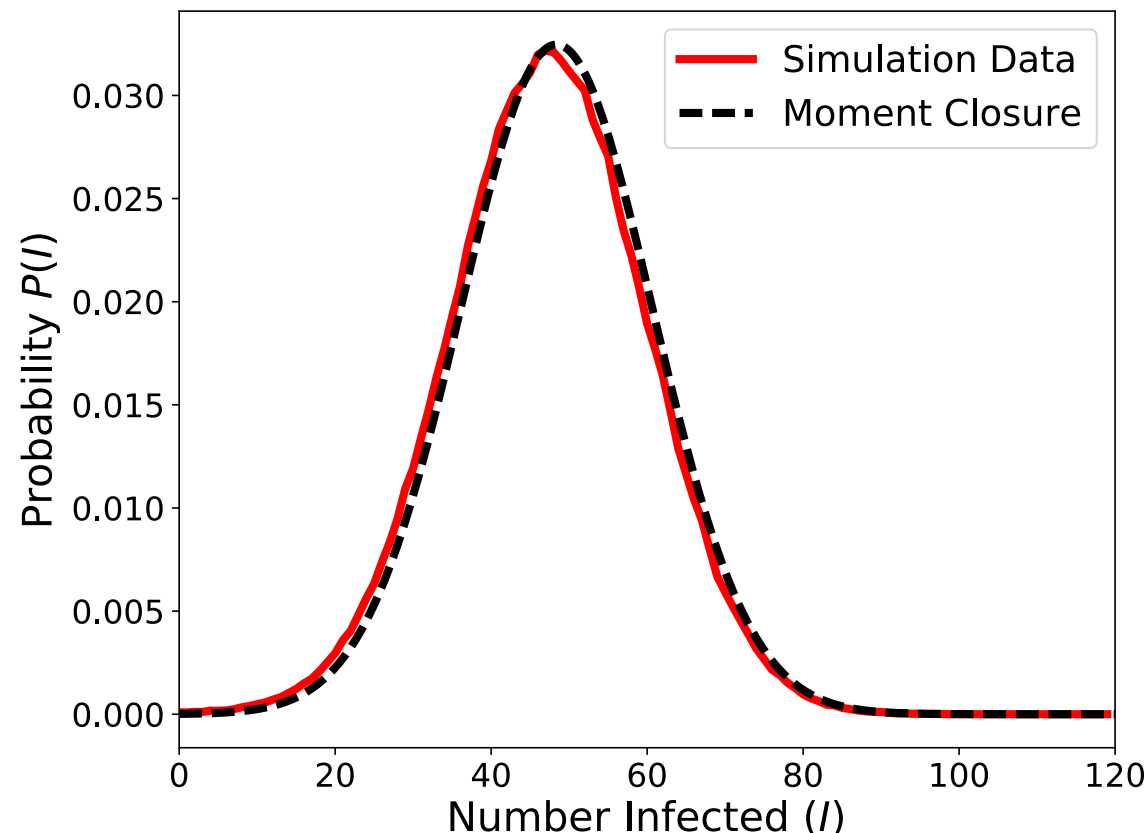
- Analytical technique that predicts *both* means and fluctuation sizes



- Good agreement for annealed networks

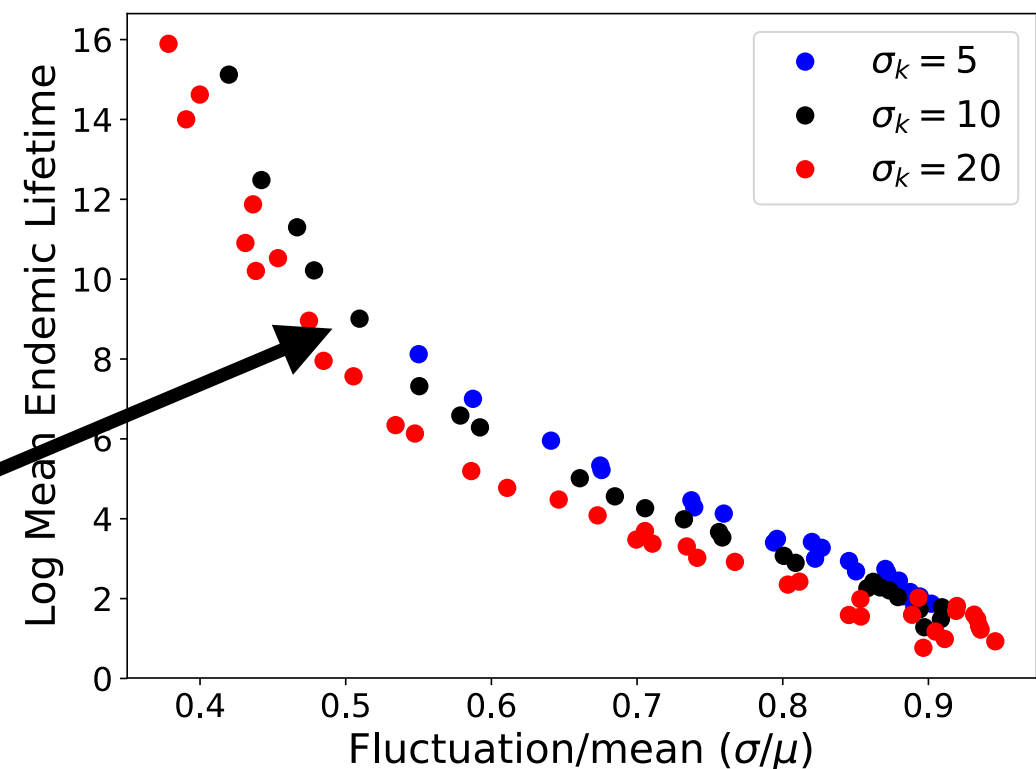
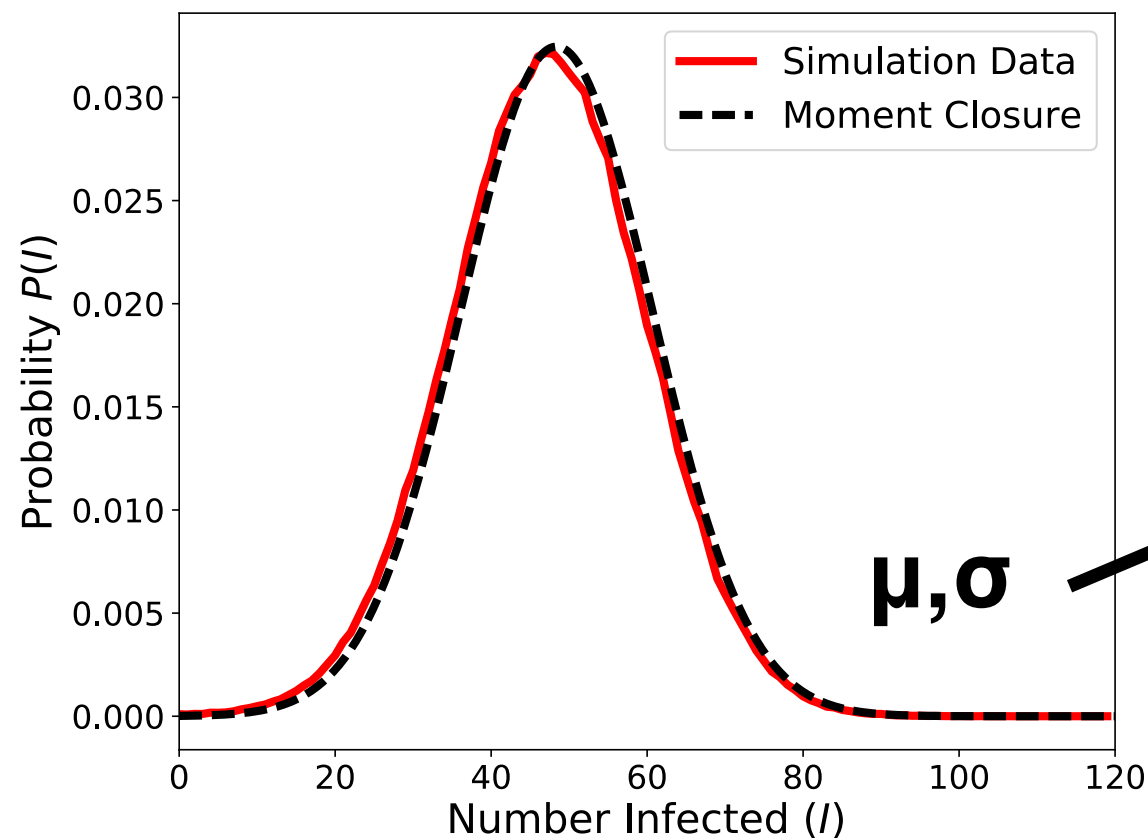
Moment Closure

- Analytical technique that predicts *both* means and fluctuation sizes
- Good agreement for annealed networks
- Hypothesis: moment closure for predicting lifetimes



Moment Closure

- Analytical technique that predicts *both* means and fluctuation sizes
- Good agreement for annealed networks
- Hypothesis: moment closure for predicting lifetimes



Summary

- Network topology can affect the persistence of endemic disease
- Higher degree heterogeneity leads to longer lifetimes
- Moment closure may be useful for predicting lifetimes

Acknowledgments

- Christopher R. Myers
- Sarabjeet Singh
- Jason Hindes
- Andrew Dolgert
- NSF Graduate Research Fellowship under Grant No. DGE-1144153

Bibliography

- M. Keeling and P. Rohani, *Modeling Infectious Diseases in Humans and Animals*
- N. T. J. Bailey, *The Elements of Stochastic Processes*
- A. Barrat et al., *Dynamical Processes on Complex Networks*
- I. Nåsell, *Mathematical Biosciences* 179, 1 (2002)
- D. Clancy and S. T. Mendy, *J of Math. Biology* 61, 527 (2009)

Constant Mean Infection Level

- Measure means, fluctuations
- Can we predict the lifetime?
- Vary graph topology

Constant Mean Infection Level

- Measure means, fluctuations
- Can we predict the lifetime?
 - 500 Nodes
 - $\langle I \rangle = 35$ Infected
- Vary graph topology
- Finite size effects
- High and low degree classes

