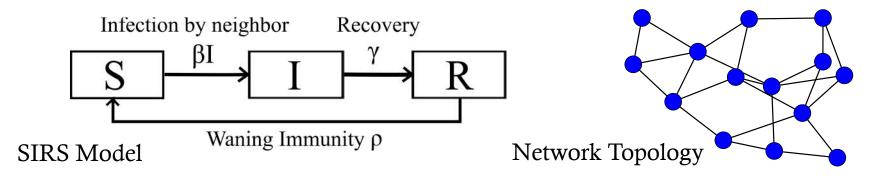
# MOMENT CLOSURE ANALYSIS OF SIRS DISEASE MODEL ON HETEROGENEOUS NETWORKS

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3/5/15

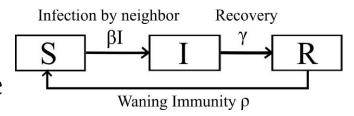
# SIR-Type Models of Disease Spread

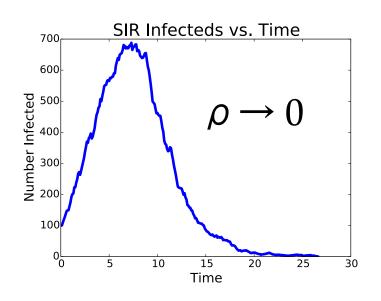
- How does disease spread through a population?
- Each individual takes on disease state
  - (S)usceptible can become infected through contact
  - (I)nfected infect susceptible nodes through contact
  - (R)ecovered no longer infected, cannot become infected again
- Transition between **extinct** and **active** phases
  - Extinct: outbreak affects finite population, dies out in finite time
  - Active: outbreak affects macroscopic fraction of population, persists

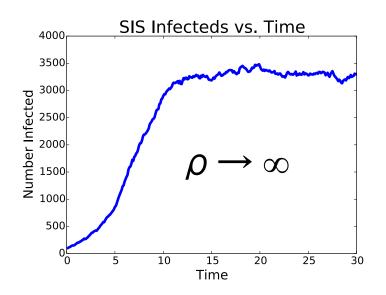


#### SIRS Model – Disease Persistence

- SIRS  $\rightarrow$  SIR as  $\rho \rightarrow 0$ , single outbreak
- SIRS  $\rightarrow$  SIS as  $\rho \rightarrow \infty$ , endemic disease



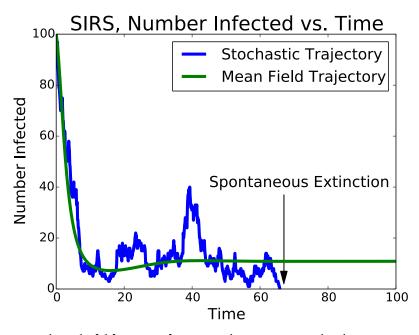




- What happens for intermediate  $\rho$ ?
- When does the SIRS model sustain endemic disease?

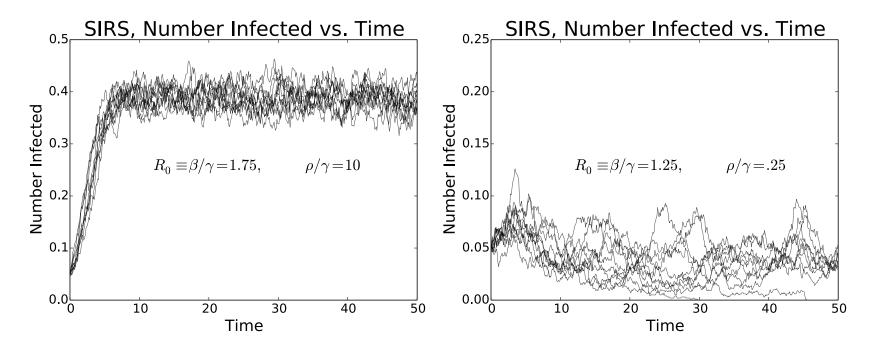
## Spontaneous Extinction from Fluctuations

- Stochastic SIRS model trajectories fluctuate
- Fluctuations not present in deterministic model
- Fluctuations lead to extinction events, end persistence



- Persists with probability given by model parameters
- To understand disease persistence, we must understand fluctuation size compared to the mean

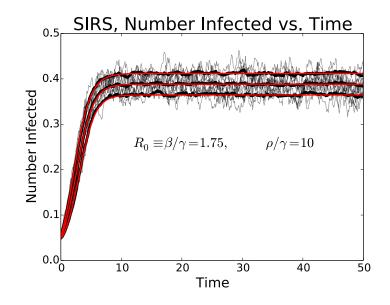
# Ensembles of Trajectories

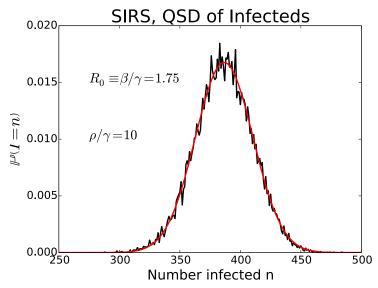


- Stochastic model trajectories drawn from an ensemble
- Want to describe the ensemble's distribution
- How does the ensemble depend on model parameters?

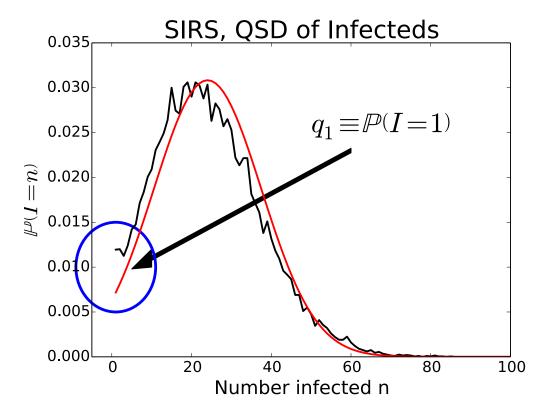
#### Moment Closure

- "Quasistatic Distribution" (QSD)
- Approximate QSD
- Assume Gaussian distribution
- Perform Moment Closure
- Obtain ODEs describing behavior of QSD mean ( $\mu_I$ ) and variance ( $\sigma_I$ )
- Produce analytical approximation to QSD as function of parameters



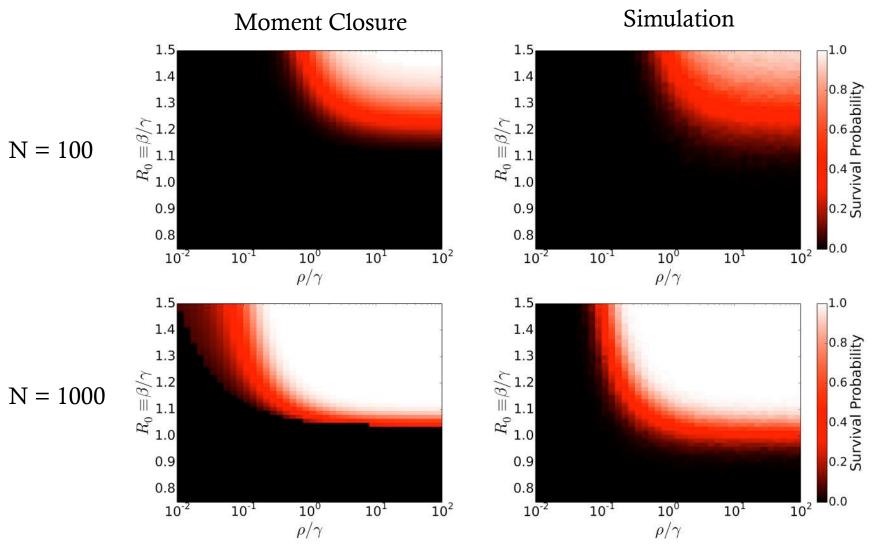


#### **Extinction Rates**



- Moment closure gives us mean and variance
- Can analytically approximate q<sub>1</sub>
- $q_1 \propto$  Rate at which trajectories become extinct
- Allows us to predict probability of persistence at time *t*

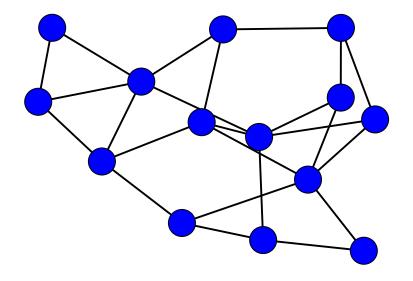
#### Persistence of Endemic Disease



• Bright regions show where disease persists at t = 50

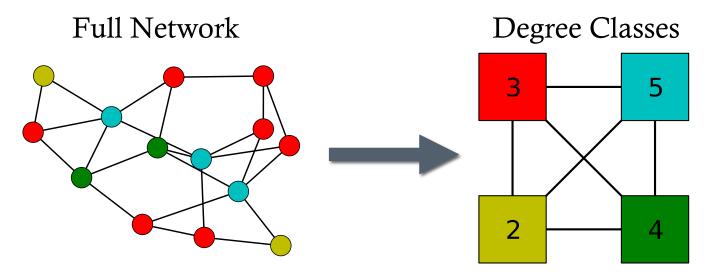
## Disease Spread on Networks

- Can we use Moment Closure to predict persistence on networks?
- Need to account for contact heterogeneity



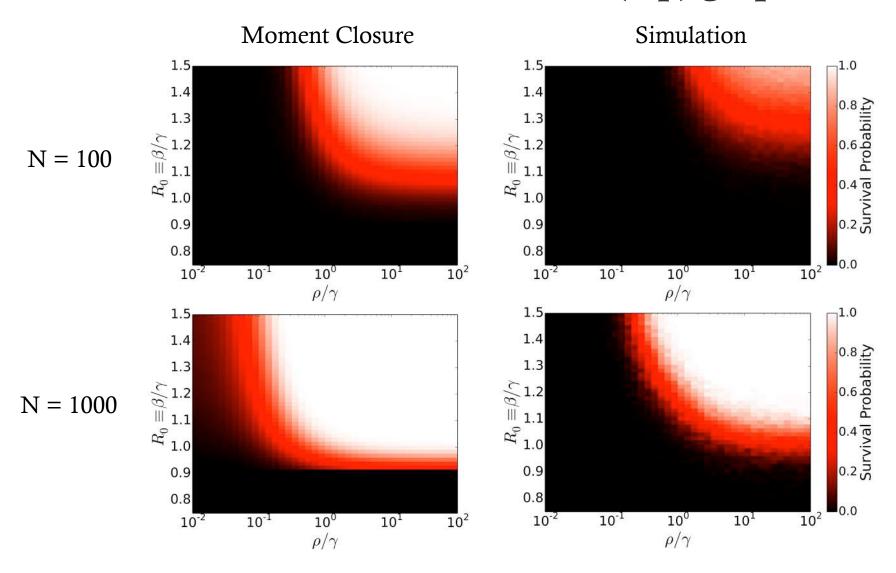
## Disease Spread on Networks

- Heterogeneous MFT
- Separate network into K coupled degree classes



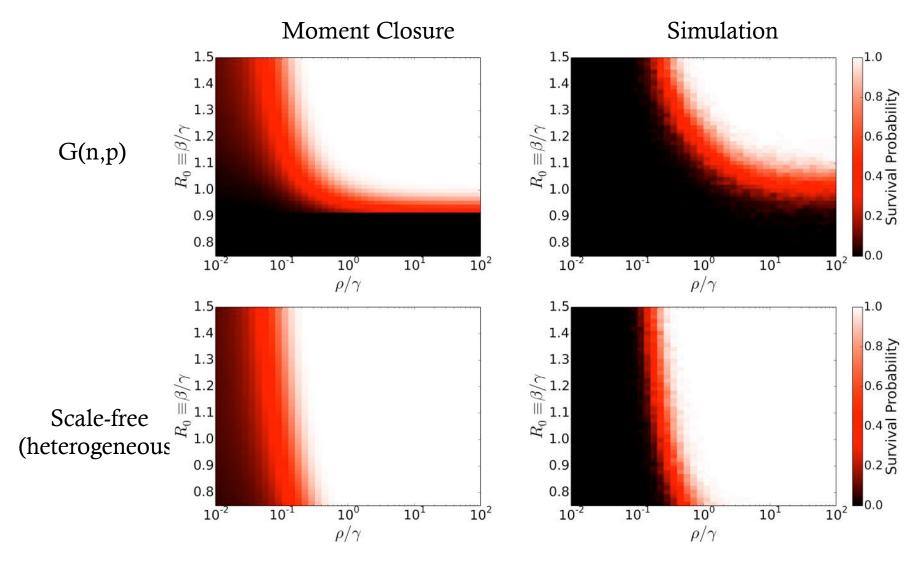
- QSD now a K-dimensional multivariate Gaussian
- Repeat Moment Closure calculations
- Obtain means and variances for all degree classes

#### Persistence of Endemic Disease, G(n,p) graphs



• Bright regions indicate persistence at t = 50

## Network Topology Affects Persistence



• Bright regions indicate persistence at t = 50

#### Conclusions

- Demonstrated utility of Moment Closure for understanding persistence of disease in stochastic SIRS model
- Extended application of Moment Closure to networks case
- Good qualitative agreement between analytic results and simulations
- Quantitative disagreement in parameter regime where assumptions fail

# Further Questions

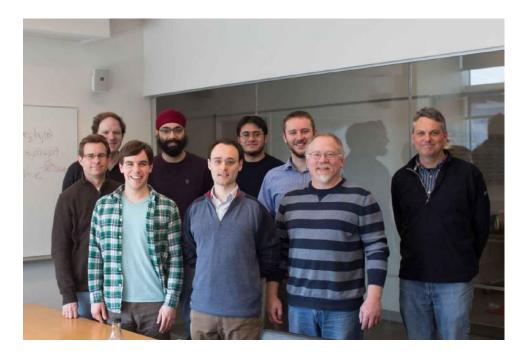
- For what types of network is our Moment Closure approximation more or less accurate?
- Are there further analytical tools to correct for when our approximations fail?

## Acknowledgements

- NSF
- DHS
- AFIDD Group
  - Andrew J. Dolgert
  - David J. Schneider
  - Sarabjeet Singh
  - Jason Hindes
  - Kevin O'Keeffe
  - Oleg B. Kogan





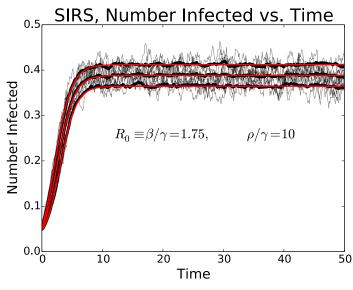


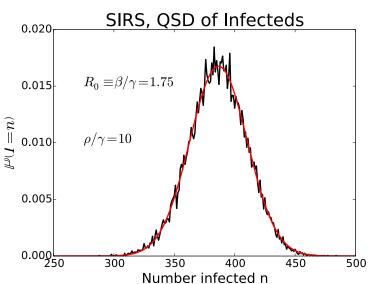
This material is based upon work supported by the National Science Foundation Graduate Research Fellowship under Grant No. DGE-1144153. Any opinion, findings, and conclusions or recommendations expressed in this material are those of the authors and do not necessarily reflect the views of the National Science Foundation. This work is also supported by the Science & Technology Directorate, Department of Homeland Security via interagency agreement no. HSHQDC-10-X-00138

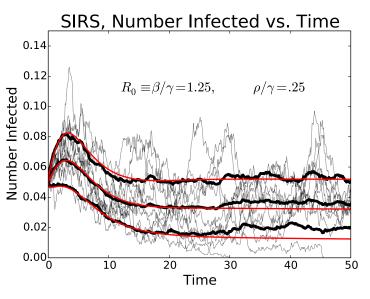
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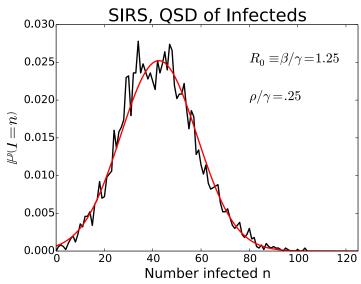
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#### Moment Closure



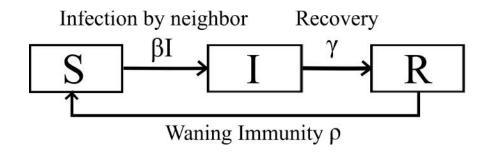






#### Stochastic SIRS Model

SIRS Model



Define transitions in stochastic model

Event	Transition	Rate
$S + I \longrightarrow I + I$	$(m,n) \longrightarrow (m-1,n+1)$	eta mn/N
$I \longrightarrow R$	$(m,n) \longrightarrow (m,n-1)$	$\gamma m$
$R \longrightarrow S$	$(m,n) \longrightarrow (m+1,n-1)$	$\rho \left(N-m-n\right)$

Kolmogorov Forward Equation

$$\frac{\partial}{\partial t} p_{m,n} = \beta / N(m+1) (n-1) p_{m+1,n-1}(t) + \gamma (n+1) p_{m,n+1}(t) + \rho (N-(m-1)-n) p_{m-1,n}(t) - (\beta m n / N + \gamma n + \rho (N-m-n)) p_{m,n}(t)$$

#### Moment Closure

- Condition on no extinction → KFE for QSD
- Extra nonlinear term appears, proportional to extinction rate
- Assume it is small (for analytical tractability)

$$\gamma q_{\cdot,1}(t) q_{m,n}(t) \longrightarrow 0$$

• Define Probability Generating Function (PGF)

$$P(x,y,t) \equiv \sum_{m,n=0}^{\infty} \underline{q}_{m,n}(t) x^{m} y^{n}$$

• Change variables in KFE to obtain PDE for PGF

$$\frac{\partial}{\partial t}P(x,y,t) = \beta(y^2 - xy)\frac{\partial^2 P}{\partial x \partial y} + \gamma(1 - y)\frac{\partial P}{\partial y} + \rho(x - 1)\left(N - x\frac{\partial}{\partial x} - y\frac{\partial}{\partial y}\right)P$$

#### Moment Closure

- Obtain PDE for *Cumulant* Generating Function (CGF)
- Assume Gaussian form of probability distribution, restricts
   CGF to quadratic form

$$K(\theta, \phi, t) = \mu_x \theta + \mu_y \phi + \sigma_{xy} \theta \phi + \frac{1}{2} \sigma_x^2 \theta^2 + \frac{1}{2} \sigma_y^2 \phi^2$$

• Obtain set of ODEs for all moments:

$$\frac{\partial \mu_{x}}{\partial t} = -\beta/N\mu_{x}\mu_{y} + \rho(N-\mu_{x}-\mu_{y}) - \beta/N\sigma_{xy} \qquad \frac{\partial}{\partial t} \sigma_{x}^{2} = \beta/N\mu_{x}\mu_{y} + \beta/N\sigma_{xy} + \rho(N-\mu_{x}-\mu_{y})$$

$$\frac{\partial \mu_{y}}{\partial t} = \beta/N\mu_{x}\mu_{y} - \gamma\mu_{y} + \beta/N\sigma_{xy} \qquad \frac{-2\beta/N(\mu_{x}\sigma_{xy} + \mu_{y}\sigma_{x}^{2}) - 2\rho(\sigma_{xy} + \sigma_{x}^{2})}{\frac{\partial}{\partial t} \sigma_{y}^{2} = \beta/N\mu_{x}\mu_{y} + \beta/N\sigma_{xy} + \gamma\mu_{y}}$$

$$\frac{\partial}{\partial t} \sigma_{y}^{2} = \beta/N\mu_{x}\mu_{y} + \beta/N\sigma_{xy} + \gamma\mu_{y}$$

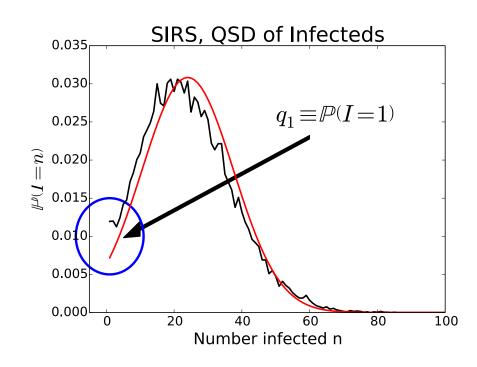
$$+\beta/N(\mu_{x}\sigma_{xy} - \mu_{y}\sigma_{xy} + \mu_{y}\sigma_{x}^{2} - \mu_{x}\sigma_{y}^{2}) - \rho\sigma_{y}^{2}$$

$$+\beta/N(\mu_{x}\sigma_{xy} - \mu_{y}\sigma_{xy} + \mu_{y}\sigma_{x}^{2} - \mu_{x}\sigma_{y}^{2}) - \rho\sigma_{y}^{2}$$

## Moment Closure Assumptions

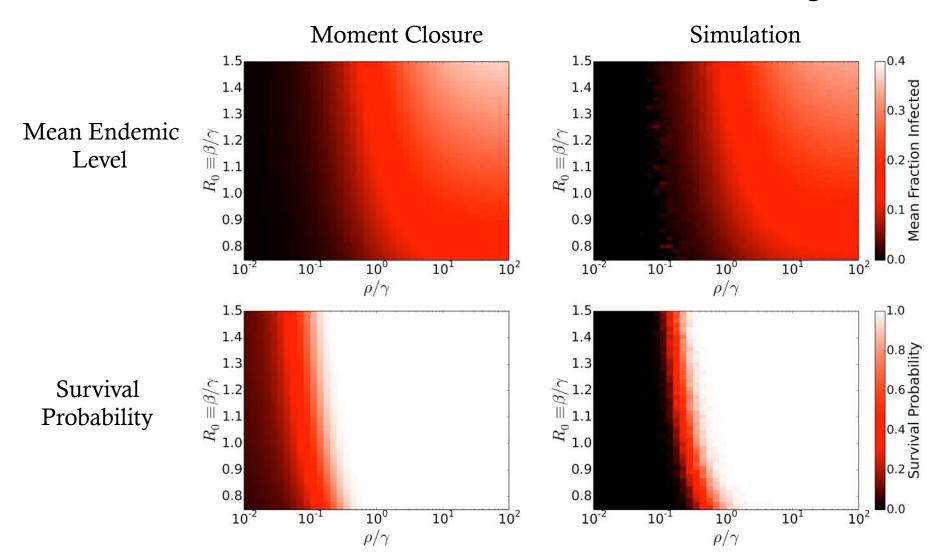
- Small rate of extinction q<sub>1</sub>
  - Fails as  $\mu_{\rm I} \downarrow$  and  $\sigma_{\rm I} \uparrow$
  - Especially tricky for low  $\mu_{\rm I}$  here we expect rapid extinction
  - Poor estimation of extinction rate when extinction rate is high
  - Is there a better analytical approximation in this regime?

- Gaussian assumption
  - Does not account for skew
  - Does not account for I > 0
  - "Renormalized Gaussian"



### Moment Closure Assumptions

• Poor estimation of extinction rate when extinction rate is high



## Other Types of Graphs

- Many different graphs with same degree distribution
- HMFT insufficient to describe
- What other graph properties affect disease persistence?

