

Epidemics in Networks

Part 3 — Qualitative observations of disease spread in networks

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Recall our key questions

For SIR:

- ▶ \mathcal{P} , the probability of an epidemic.
- ▶ \mathcal{A} , the “attack rate”: the fraction infected if an epidemic happens (better named the attack ratio).
- ▶ \mathcal{R}_0 , the average number of infections caused by those infected early in the epidemic.
- ▶ $I(t)$, the time course of the epidemic.

For SIS:

- ▶ \mathcal{P}
- ▶ $I(\infty)$, the equilibrium level of infection
- ▶ \mathcal{R}_0
- ▶ $I(t)$

Introduction

Sample stochastic simulations

Impact of network properties

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Assumptions

We start with some simple assumptions:

- ▶ SIS or SIR disease on a fixed static network.
- ▶ Susceptible nodes , infected nodes , and recovered nodes .

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- ▶ SIS or SIR disease on a fixed static network.
- ▶ Susceptible nodes , infected nodes , and recovered nodes .
- ▶ Disease transmits along an edge with rate β (many authors use τ)
- ▶ Infected individuals recover with rate γ

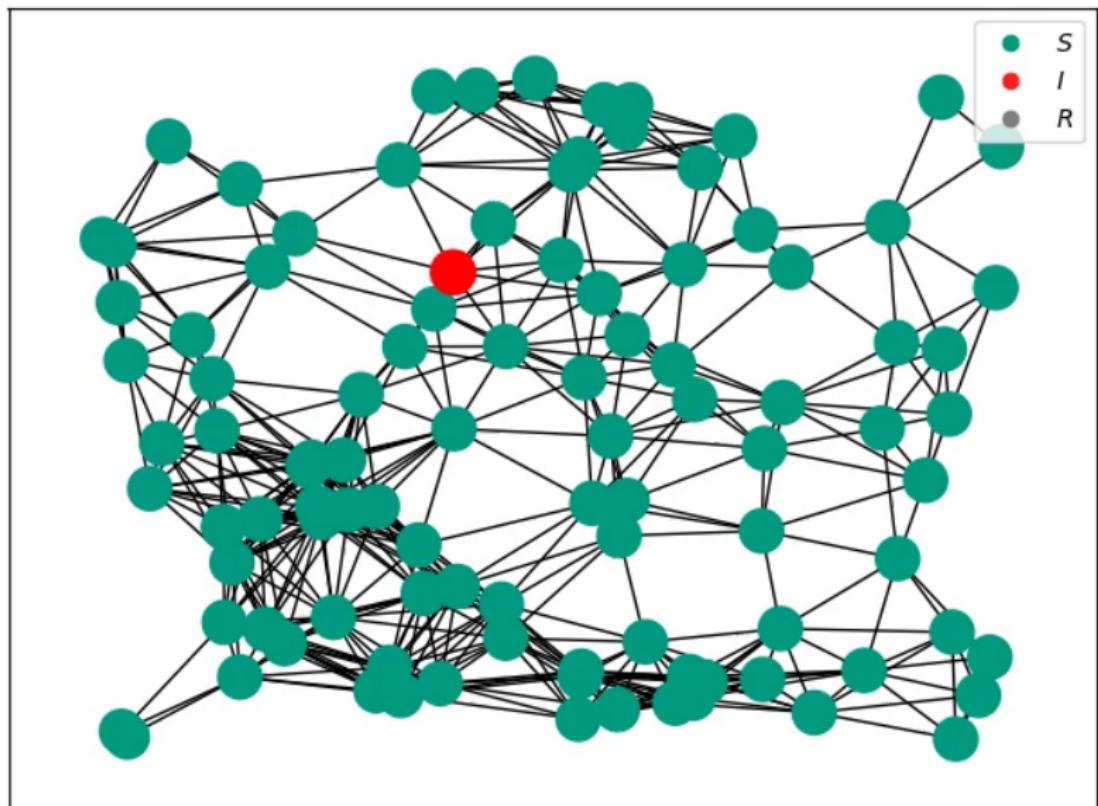
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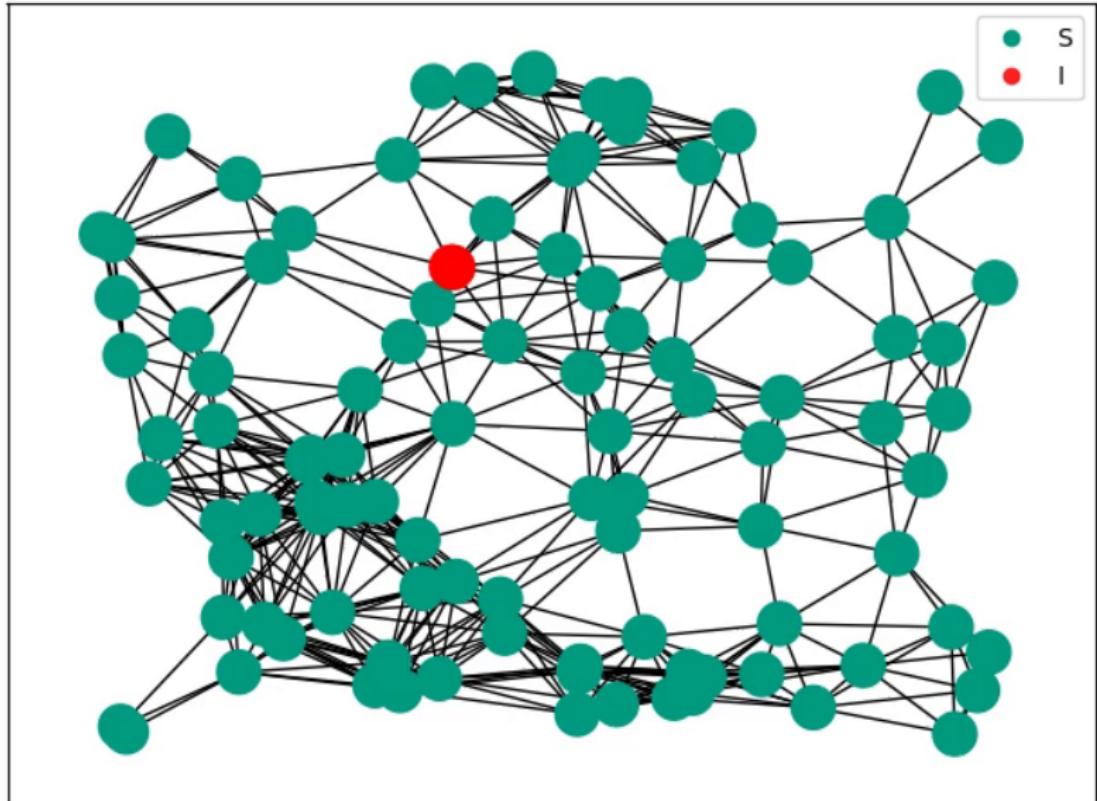
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Sample SIR epidemic



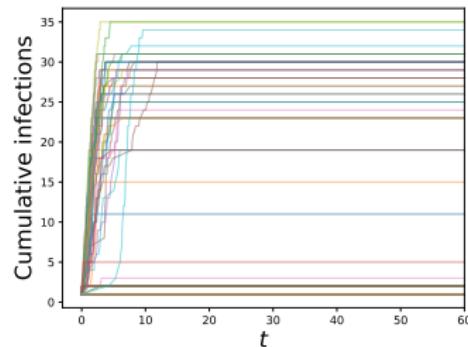
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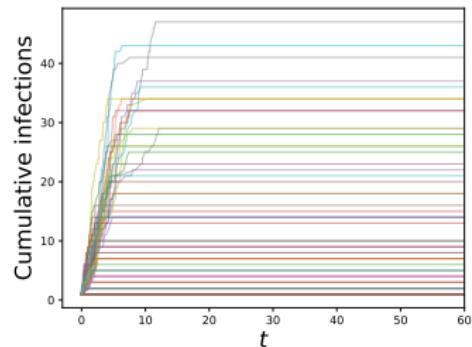
Stochastic simulation — SIR on network case

SIR disease spread with $\langle K \rangle = 5$, $\beta = 0.4$, and $\gamma = 1$.

$$P(5) = 1,$$



$$P(1) = P(9) = 0.5$$



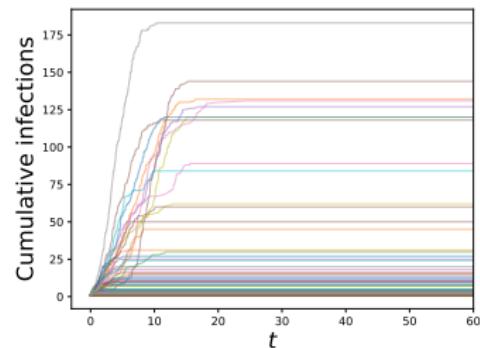
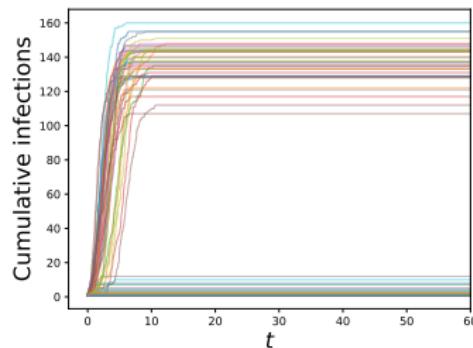
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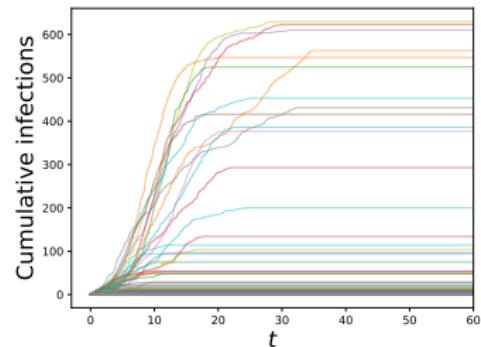
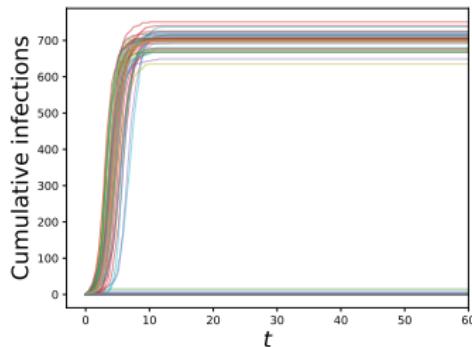
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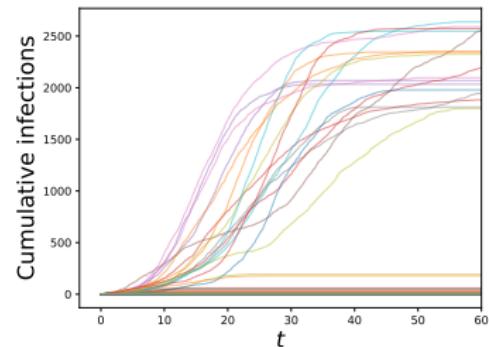
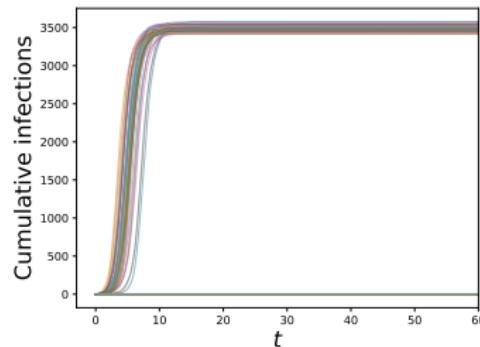
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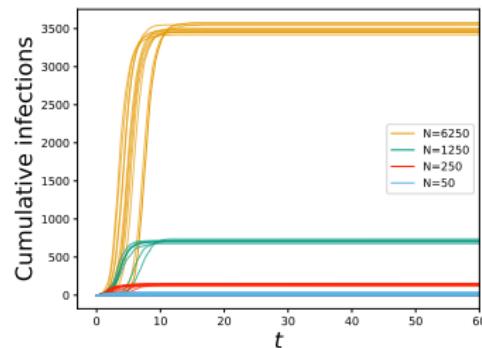


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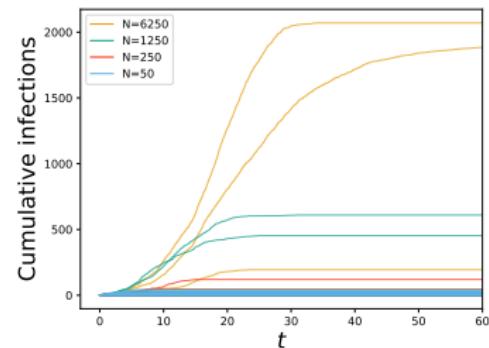
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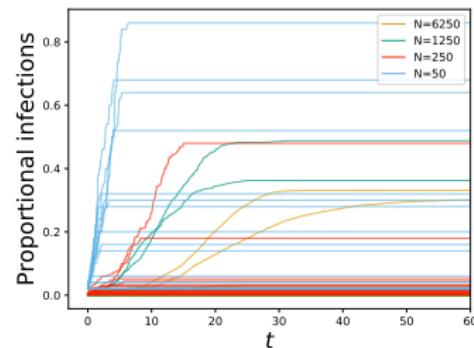
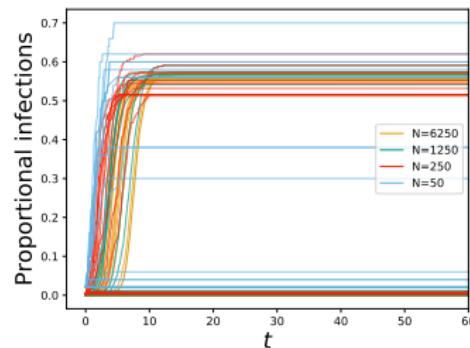


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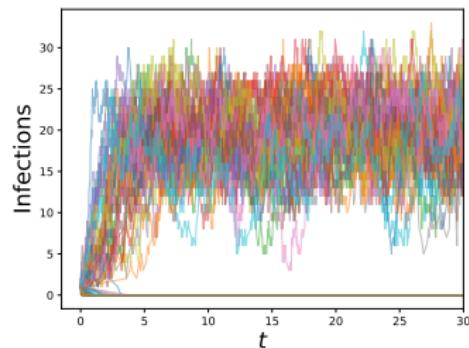
SIR observations

- ▶ In large networks outbreaks are either small (non-epidemic) or large (epidemic).
- ▶ Small outbreaks don't care about network size (once network is sufficiently large).
- ▶ Epidemic sizes are proportional to network size.
- ▶ The degree distribution affects the final size and the early growth.

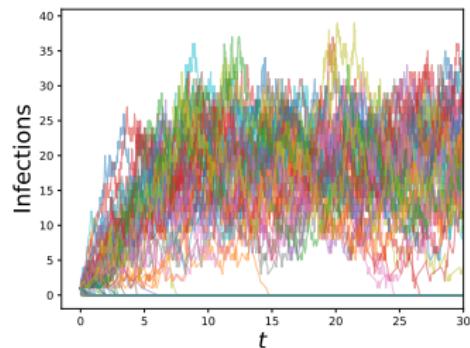
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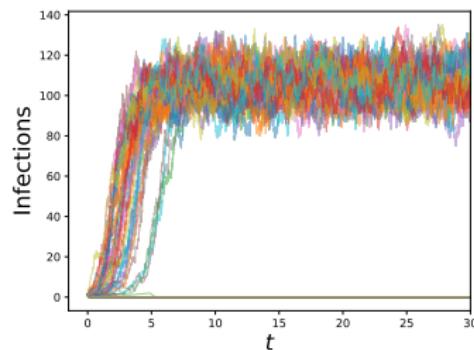


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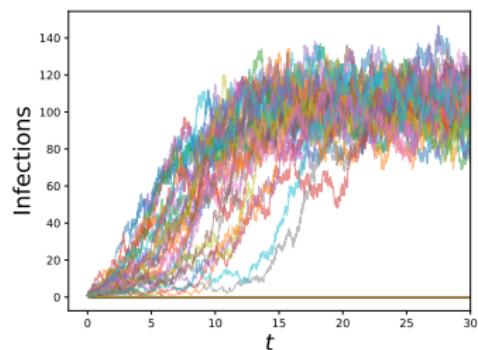
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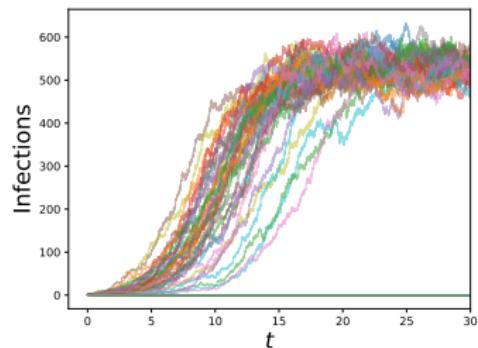
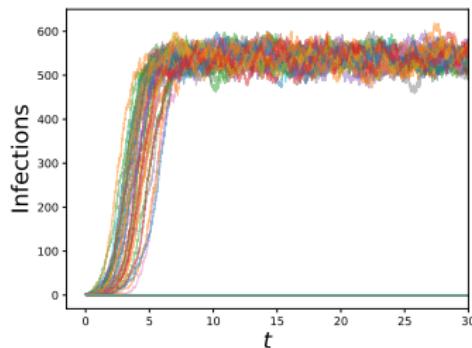
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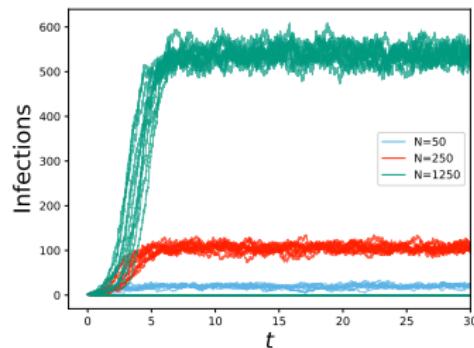


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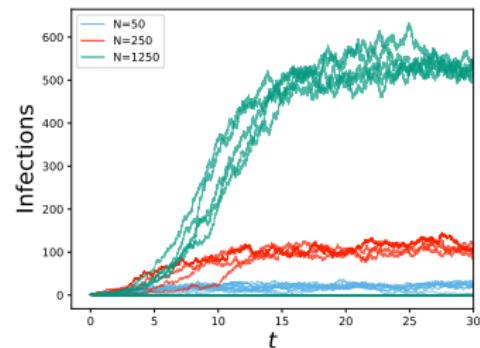
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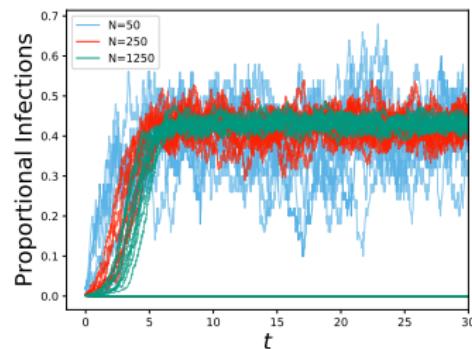
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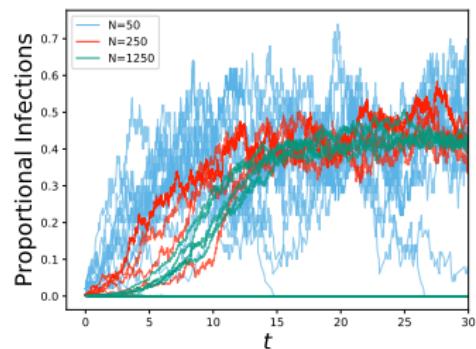
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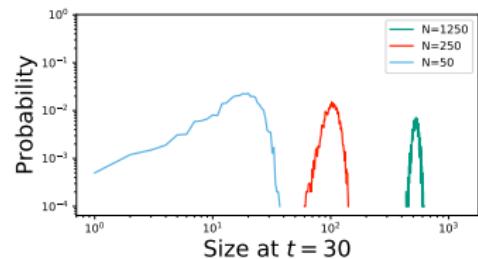
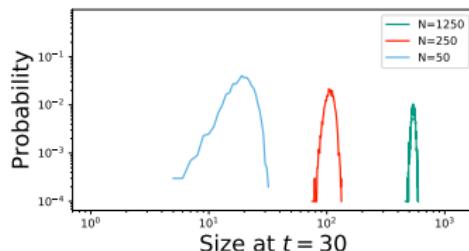
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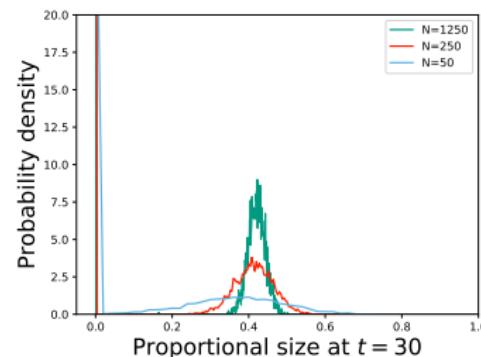
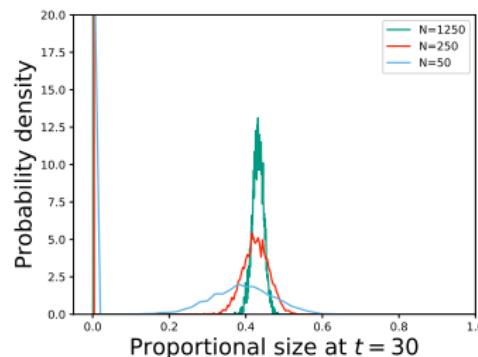
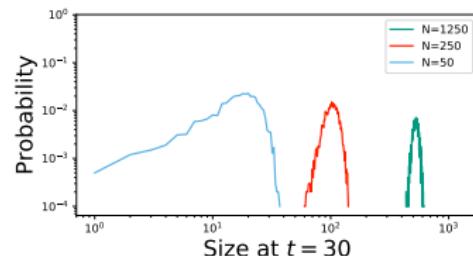
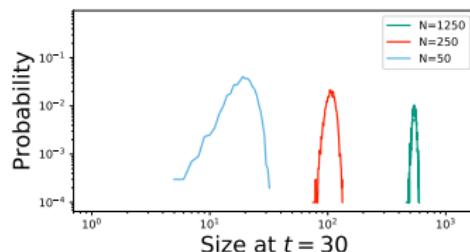


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SIS observations

- ▶ In large networks outbreaks either go extinct quickly (non-epidemic) or reach an endemic equilibrium (epidemic).
- ▶ Small outbreaks don't care about network size.
- ▶ Epidemic equilibrium sizes are proportional to network size.
- ▶ Coefficient of variation decreases for large networks. [typical deviation from mean is small compared to mean.]

Introduction

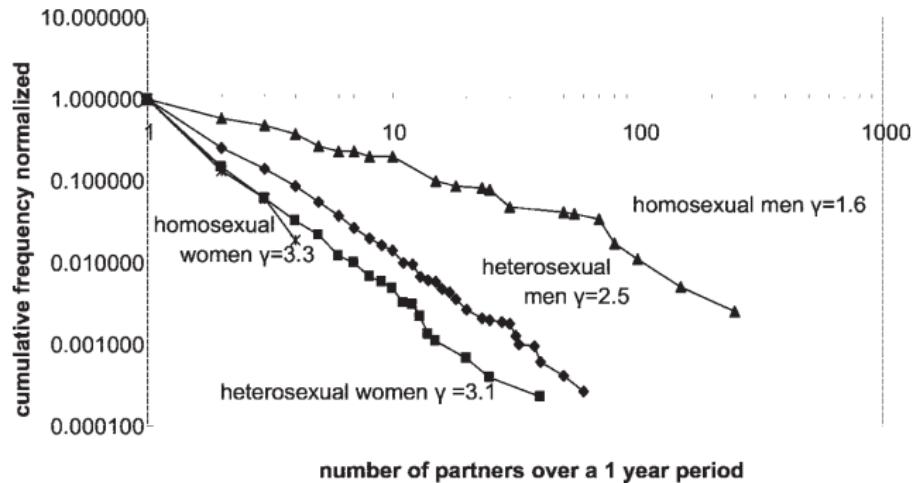
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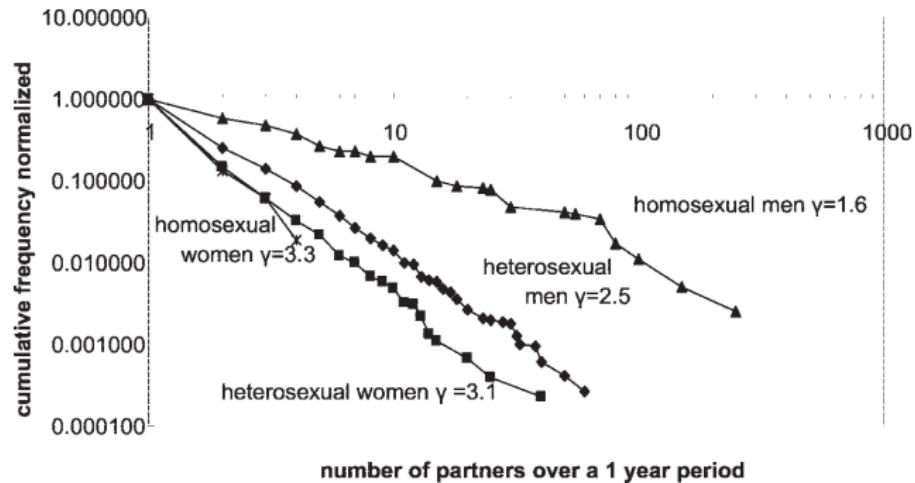
Degree distribution

From [1]:



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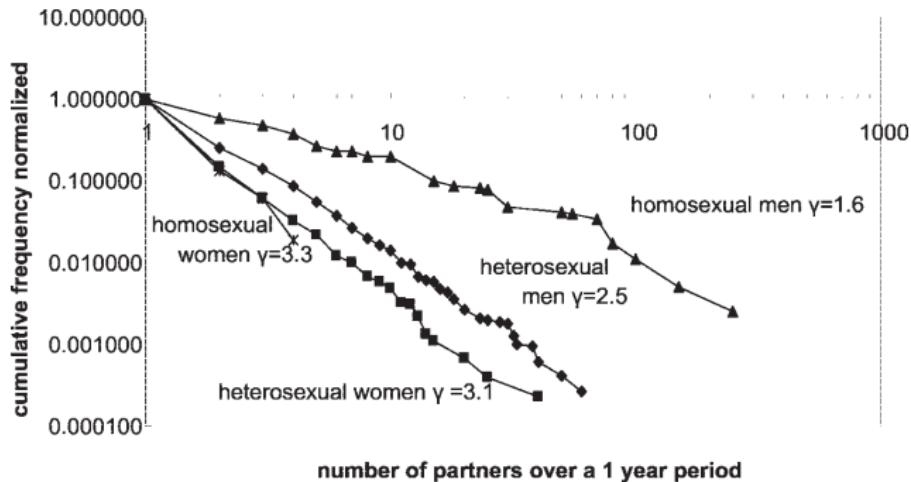
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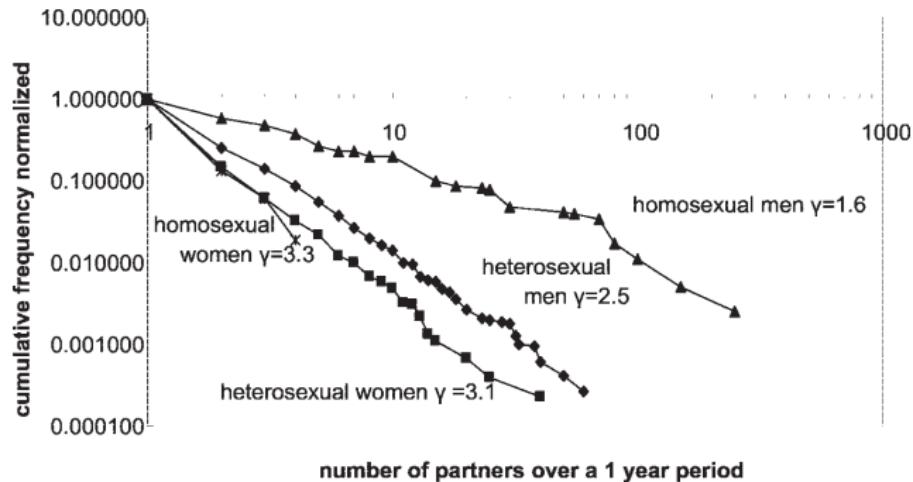


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Holding $\langle K \rangle$ fixed, increasing heterogeneity increases \mathcal{R}_0 .

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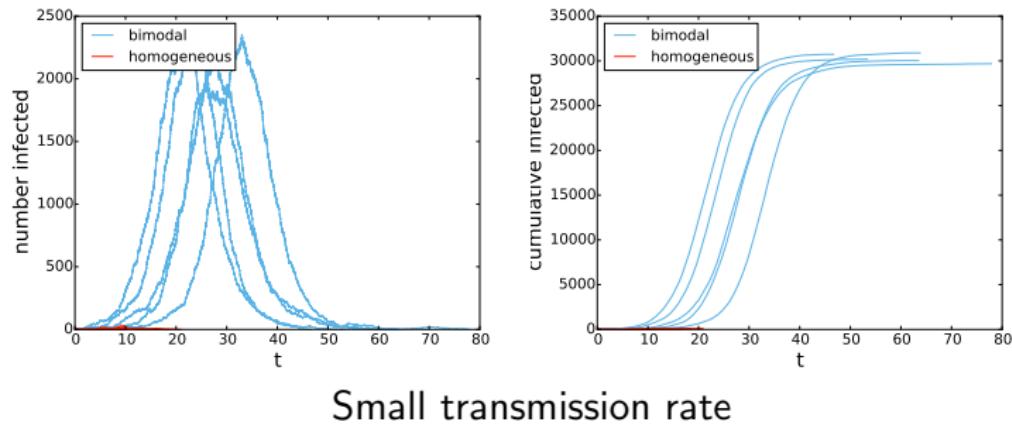


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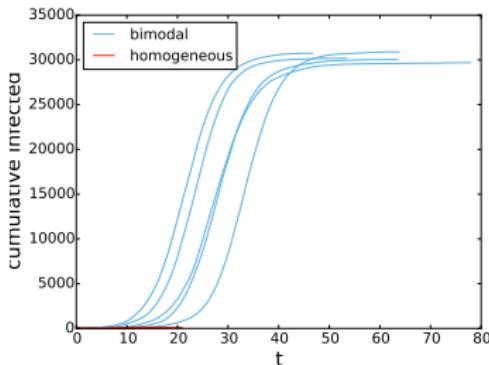
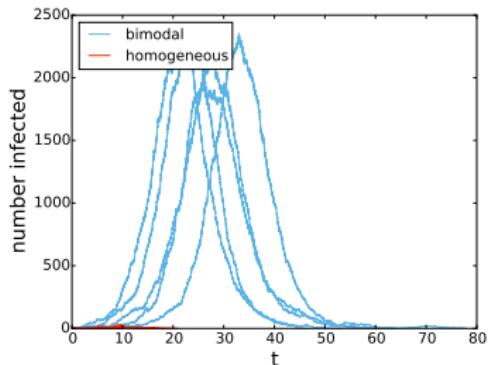
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Why does this increase \mathcal{R}_0 ?

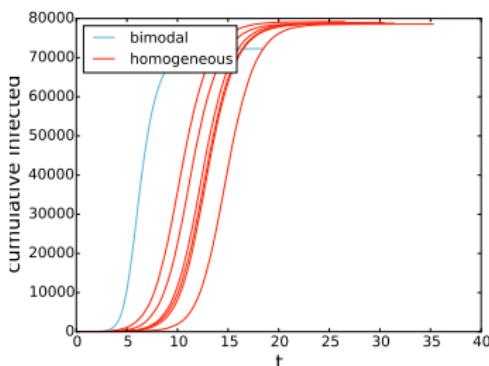
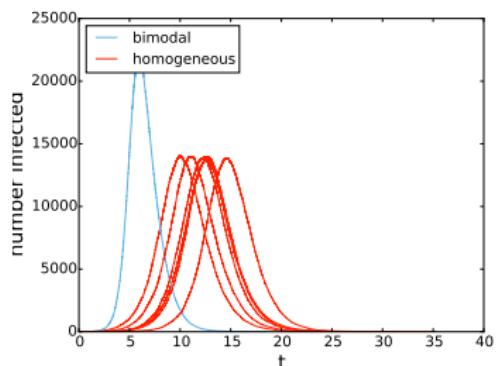
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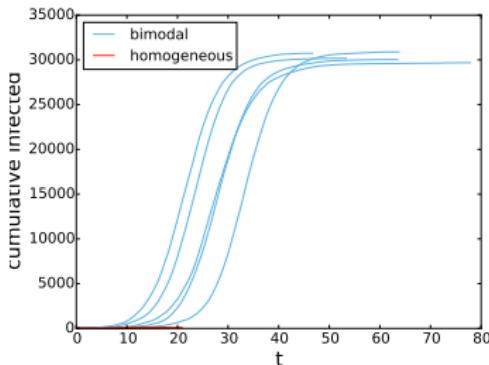
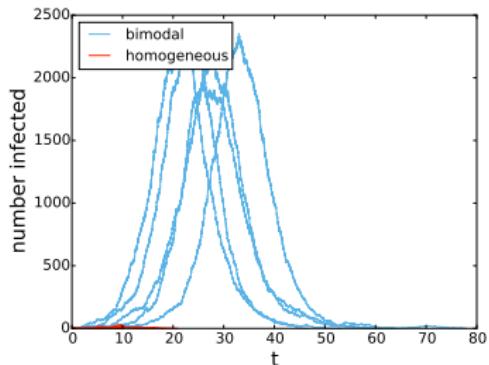


Small transmission rate

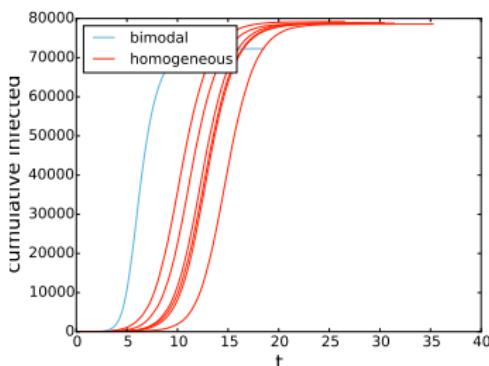
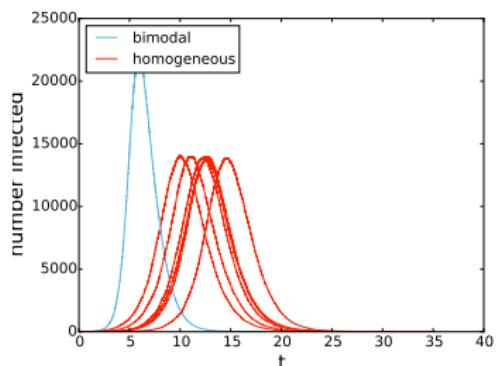


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Understanding impact on final size

- ▶ Why does degree heterogeneity increase final epidemic size at smaller transmission rate?

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- ▶ Why does degree heterogeneity decrease final epidemic size at higher transmission rate?

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If high degree individuals preferentially contact high degree individuals, impact on \mathcal{R}_0 : Increases it.

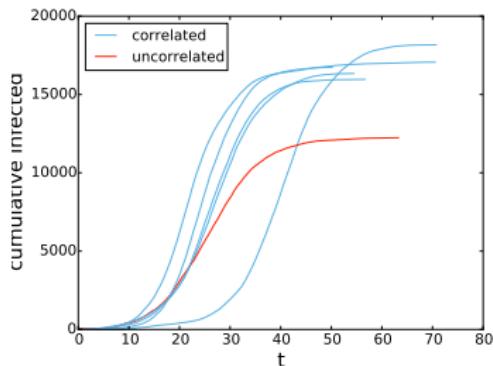
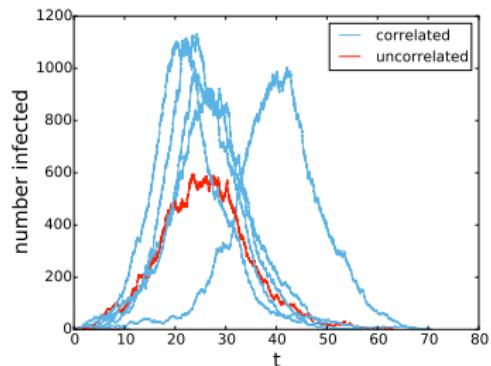
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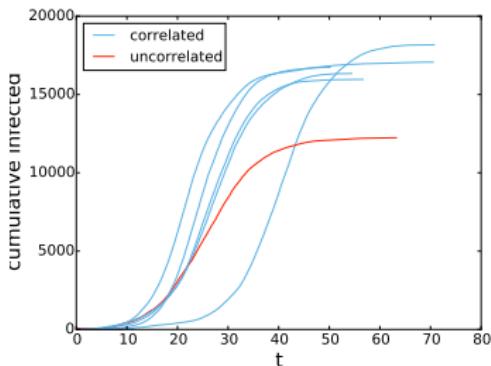
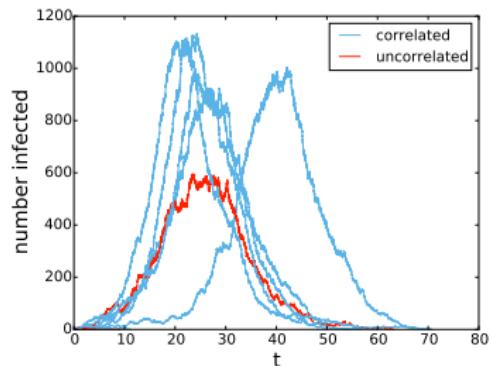
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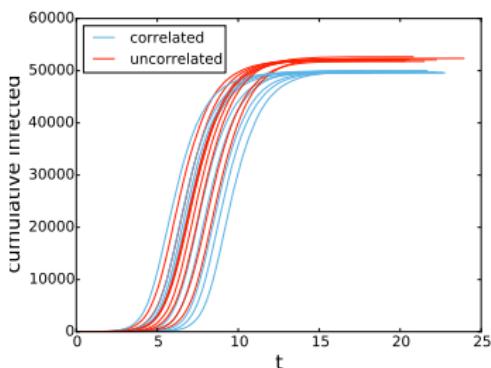
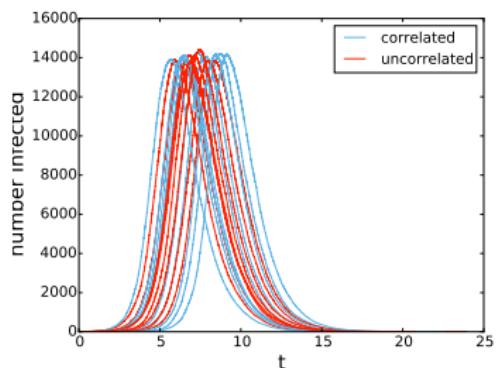


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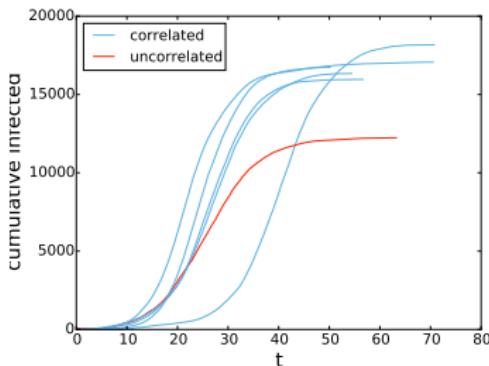
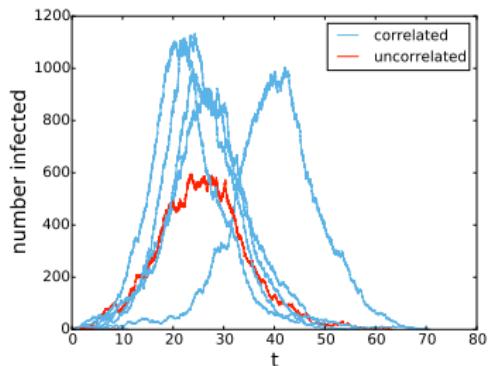


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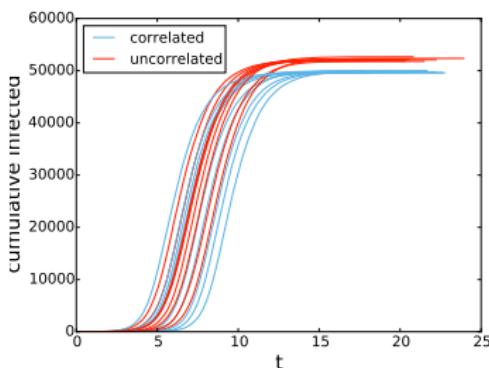
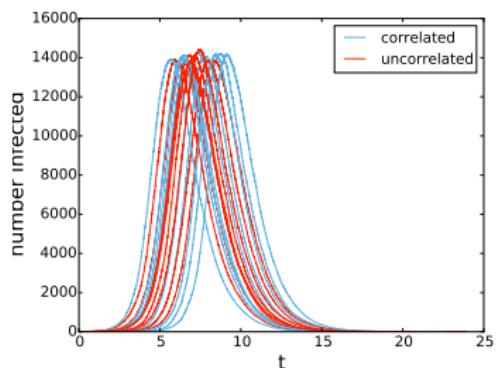


Large transmission rate

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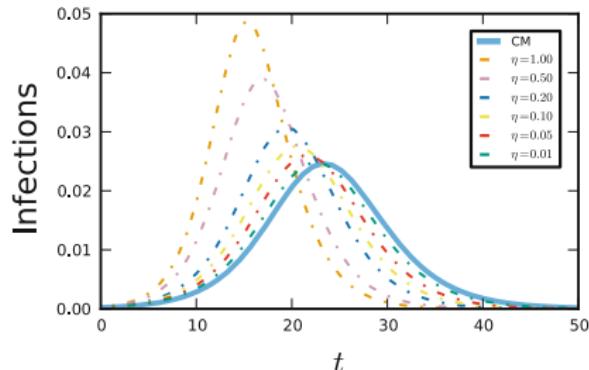
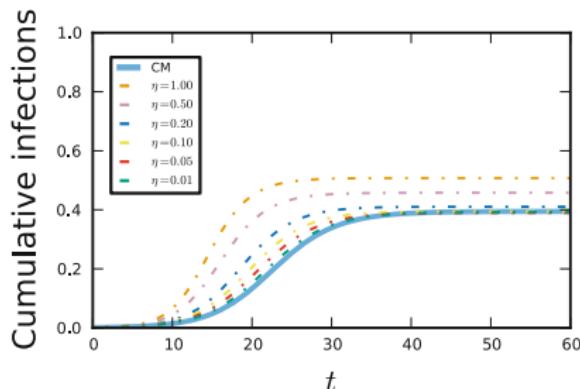
Impact on \mathcal{R}_0 :

For SIR, long partnership duration decreases \mathcal{R}_0 because repeated transmissions are wasted.

For SIS, it is complex — repeated transmissions are wasted, but long-lasting partnerships help ensure that newly-recovered high degree nodes are quickly reinfected [2].

Partnership duration

Sample SIR epidemics from [3]



(η is inverse partnership duration, “CM” is static Configuration Model)

Clustering

If partnerships are clustered, even early on individuals who become infected are likely to have partners who are infected by others.

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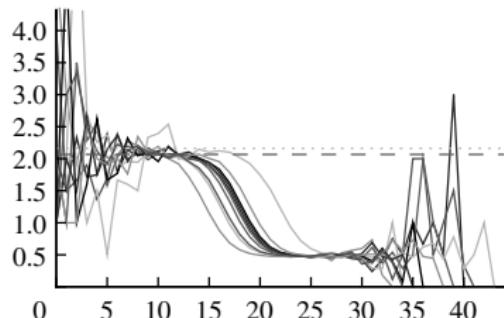
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Ratio of successive generation sizes from [4]

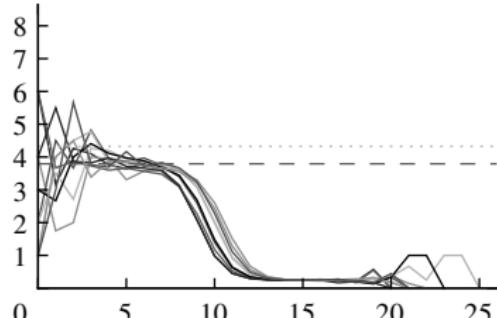
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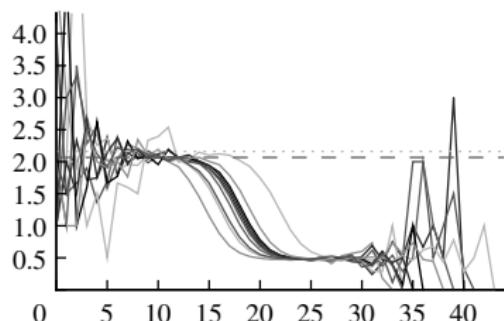
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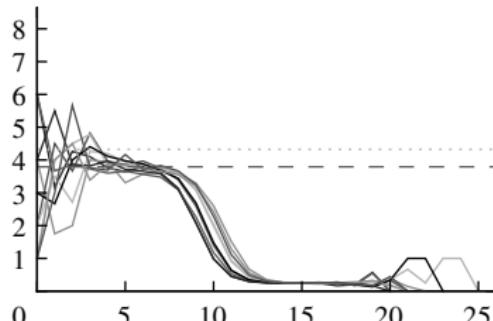
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Dotted line is prediction ignoring clustering. Dashed line is correction accounting for triangles and squares.

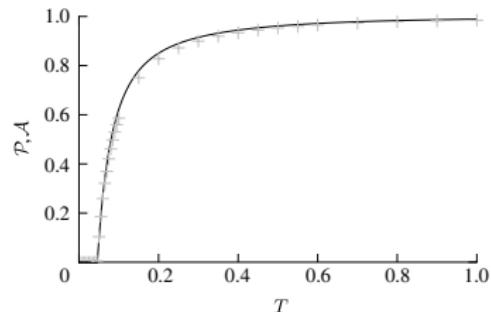
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Comparison of unclustered prediction (line) with stochastic simulation (symbols)



(horizontal axis is transmission probability, vertical is fraction infected.)

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References

References |

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