

Spreading Processes

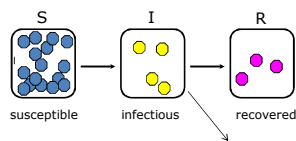
Modeling Infectious Disease Dynamics with Networks



What is epidemiology?

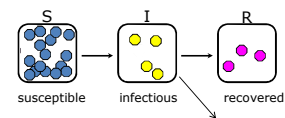
- **Terms**
 - Susceptible
 - Infected
 - Epidemic
- **Questions asked:**
 - will an epidemic occur?
 - what is the typical size of an outbreak?
 - what determines the probability of an epidemic?
 - How do we control the spread?

Compartmental models



$$\begin{aligned}\frac{dS}{dt} &= -\beta IS \\ \frac{dI}{dt} &= \beta IS - \gamma I - \alpha I \\ \frac{dR}{dt} &= \gamma I\end{aligned}$$

Compartmental models

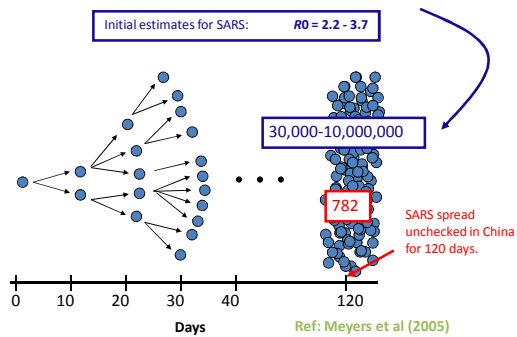


$$R_0 = \frac{\text{Infection rate}}{\text{Mortality} + \text{Recovery rate}}$$

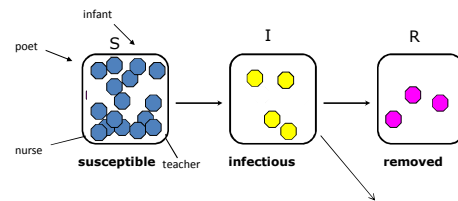
Reproductive Ratio/Number: average number of secondary cases per infected individual

SARS and its Reproductive Ratio (R_0)

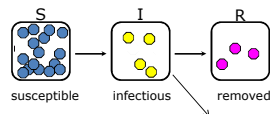
R_0 = reproductive ratio → number of secondary infections caused by a single infected person



What's missing in this model?



Review: Compartmental Models



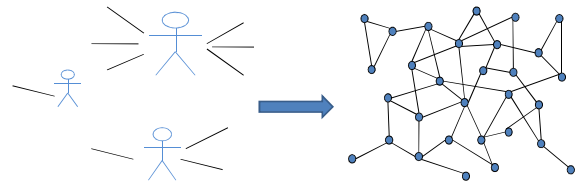
Advantages:

- Simple
- Extendable
- Amenable to mathematical analysis

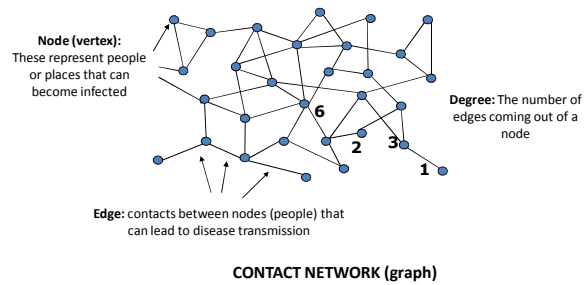
Disadvantage:

- Assume everyone in the population is equally vulnerable to infection and to spreading infection.

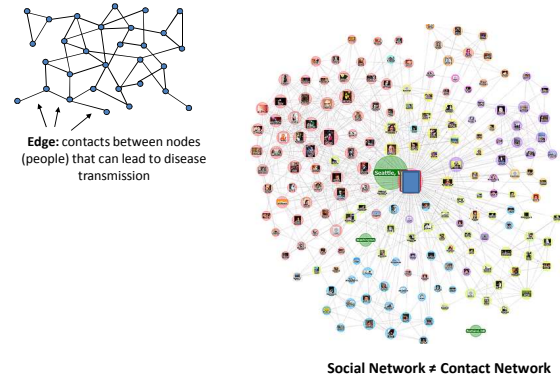
What can we do to fix this?



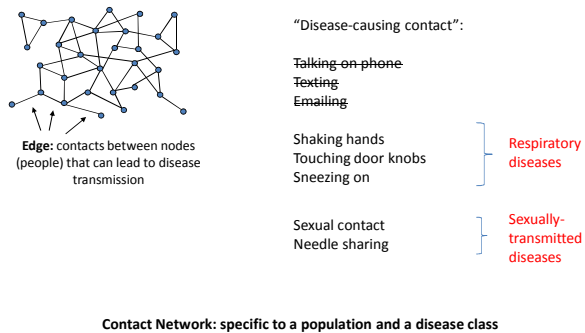
Terminology



What makes up a contact network?



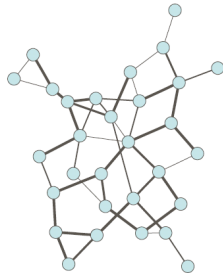
What makes up a contact network?



Capturing Contacts in Networks

Weighted Contacts

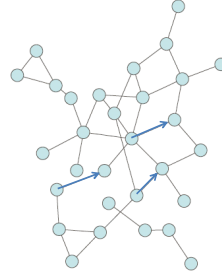
A way to represent the strength of a contact



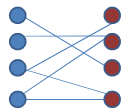
E.g.
Duration of contact,
Number of passengers

Directed Contacts

A way to represent asymmetry in contact

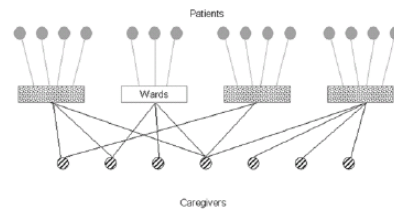


Bipartite Networks



Transmission of STD in
heterosexual network

Co-location Network



Ref: Meyers et al (2003)
Emerging Infectious Diseases 9, 204-210

Office Hours

Shweta Bansal

Office: Mueller Lab 510 C

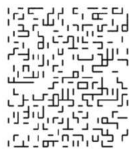
Email: sbansal@psu.edu

Office Hours: All week 1-2pm

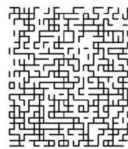
Predicting Epidemics

Bond percolation

- Start with a lattice (or network)
- Draw (or mark) the edges with a certain probability p
- The remaining edges are open (unmarked)
- At a critical probability p_c a spanning cluster appears



$p=0.315$

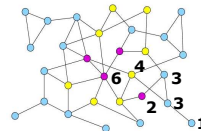


$p=0.525$

Ref: H. L. Frisch and J. M. Hammersley, J. SIAM 11, 894 (1963)

Predicting Epidemics

Degree Distribution



Transmissibility



Probability of transmission:

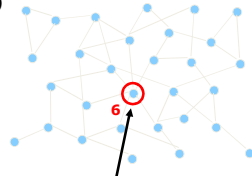
- rate of contact
- infectiousness
- susceptibility

Predicting epidemics: Percolation

Probability generating function (PGF) for the degree distribution

$$G_0(x) = \sum_{k=0}^{\infty} p_k x^k$$

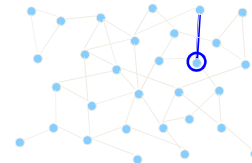
Degree	Frequency
1	3/30 = 0.1
2	7/30 = 0.23
3	11/30 = .37
4	6/30 = 0.2
5	2/30 = 0.07
6	1/30 = 0.03



G_0 tells you about the degrees of randomly chosen vertices

Ref: H. Wilf, *Generatingfunctionology* (1994)

Predicting epidemics: Percolation



G_1 tells you about the degrees of vertices arrived at along randomly chosen edges

$$G_1(x) = \frac{1}{\langle k \rangle} G'_0(x)$$

PGF for the excess degree distribution

Ref: Newman (2002) *Phys. Rev. E* 66, 016128 (2002)

Predicting epidemics: Percolation

Including disease transmission

The pgf for the number of secondary infections leading from an infected vertex

$$G_0(x; T) = \sum_{m=0}^{\infty} \sum_{k=0}^{\infty} p_k \binom{k}{m} T^m (1-T)^{k-m} x^m = G_0(1 + (x-1)T)$$

the probability that a randomly selected vertex has degree k

the probability m of the k edges transmit infection

$$G_1(x; T) = G_1(1 + (x-1)T)$$

Predicting epidemics: Percolation

$H_0(x; T)$ pgf for infected cluster size distribution

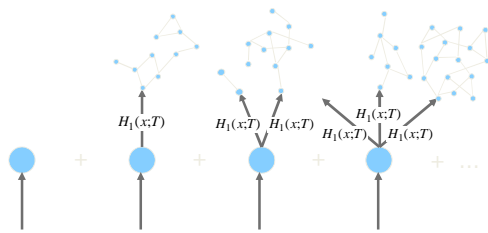
$H_1(x; T)$ pgf for infected cluster size distribution at end of randomly chosen edge

Ref: Newman (2002) *Phys. Rev. E* 66, 016128 (2002)

Predicting epidemics: Percolation

The pgf for the cluster size distribution at the end of a randomly chosen edge:

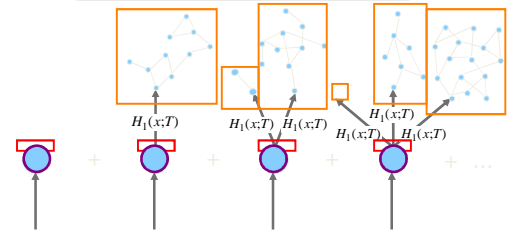
$$H_1(x;T)$$



Predicting epidemics: Percolation

The pgf for the cluster size distribution at the end of a randomly chosen edge:

$$H_1(x;T) = xG_1(H_1(x;T), T)$$



Predicting Epidemics: the Math

$H_0(x;T)$ pgf for infected cluster size distribution

$H_0'(1;T) = \langle s \rangle$ average infected cluster size

$$\langle s \rangle = 1 + \frac{G_0'(1;T)}{1 - G_1'(1;T)} = 1 + \frac{T G_0'(1)}{1 - T G_1'(1)}$$

$$T_c = 1/G_1'(1) \quad \text{Epidemic threshold}$$

Ref: Newman (2002) Phys. Rev. E 66, 016128 (2002)

Predicting Epidemics: Epidemic Threshold

$$T < T_c$$

only small disease outbreaks occur whose avg. size will be $\langle s \rangle$

$$T > T_c$$

larger epidemics can occur, and $\langle s \rangle$ does not apply

Predicting Epidemics: Epidemic Size

$T > T_c$ Epidemic possible

What will be the probability of an epidemic?

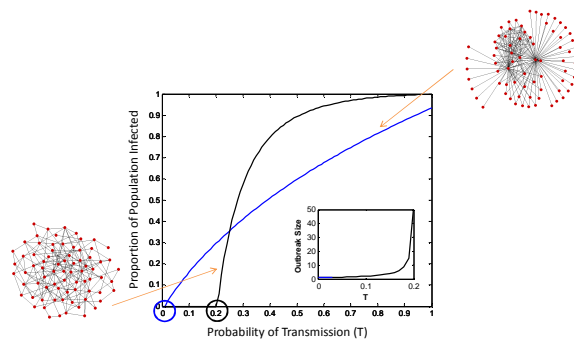
$P = 1 - \text{Prob}(\text{only small outbreaks})$

$$= 1 - H_0(1; T)$$

$$S = 1 - H_0(1; T)$$

Size of epidemic

Predicting Epidemics: Results



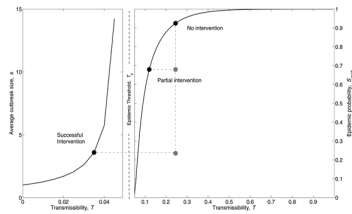
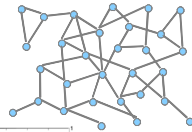
Dynamical Models on Networks

- Pair approximation methods
Ref: Keeling (1999) Proc. Roy. Soc. Lond. B 266 859-869
- Heterogeneous-mixing methods
Ref: Pastor-Satorras & Vespignani (2002) Phys. Rev. E 65, 035108
- Dynamical PGF methods
Ref: Volz (2008) Journal of Mathematical Biology, 56 3

Controlling Epidemics

Assessing Control Strategies (I)

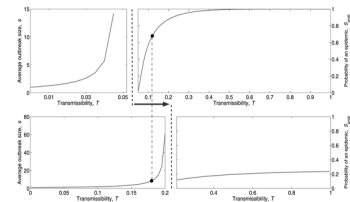
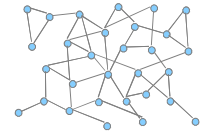
- Transmission-reducing interventions (reducing T_{ij} on some or all edges)
 - Face masks, gloves
 - Washing hands



Ref: Pourbohloul, Meyers, et al. (2005) EID Vol 11

Assessing Control Strategies (II)

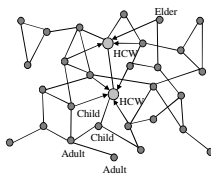
- Contact-reducing interventions (reducing number of edges)
 - Quarantining a patient
 - Closing schools
 - Social distancing



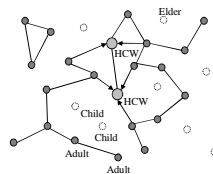
Ref: Pourbohloul, Meyers, et al. (2005) EID Vol 11

Assessing Control Strategies (III)

- Immunization



Total Population



Vaccinated Population