# Contagion

Complex Networks, Course 303A, Spring, 2009

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Basic Contagion Models

Social Contagion Models

Network version All-to-all networks

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### **Outline**

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# Some large questions concerning network contagion:

- For a given spreading mechanism on a given network, what's the probability that there will be global spreading?
- If spreading does take off, how far will it go?
- 3. How do the details of the network affect the outcome?
- 4. How do the details of the spreading mechanism affect the outcome?
- 5. What if the seed is one or many nodes?
- Next up: We'll look at some fundamental kinds of spreading on generalized random networks.

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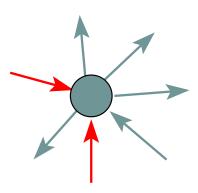
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## Spreading mechanisms



uninfected infected

- General spreading mechanism: State of node *i* depends on history of i and i's neighbors' states.
- Doses of entity may be
- May have multiple,

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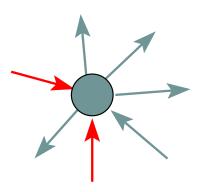
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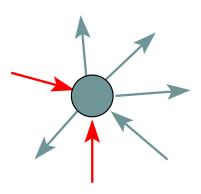
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## Spreading mechanisms



- uninfected
- infected

- General spreading mechanism: State of node *i* depends on history of i and i's neighbors' states.
- Doses of entity may be stochastic and history-dependent.
- May have multiple, interacting entities spreading at once.

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- For random networks, we know local structure is pure branching.
- Successful spreading is ∴ contingent on single edges infecting nodes.

► Focus on binary case with edges and nodes either infected or not

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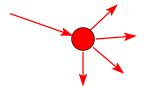
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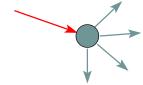




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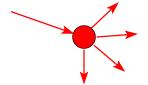
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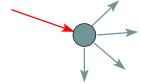
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We need to find:

r = the average # of infected edges that one random infected edge brings about.

 $\triangleright$  Define  $\beta_k$  as the probability that a node of degree k

$$r = \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle} \cdot \underbrace{\beta_k} \cdot \underbrace{(k-1)}_{\mbox{Prob. of infection}}$$

$$+\sum_{k=0}^{\infty} \frac{\widehat{kP_k}}{\langle k \rangle}$$

$$\underbrace{(1-\beta_k)}_{\text{Prob. of}}$$

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- We need to find:
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 $r = \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle} \cdot \frac{\beta_k}{\text{Prob. of}} \cdot \frac{(k-1)}{\text{infection}}$ prob. of infection infected edges

# outgoing infected edges

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$$r = \sum_{k=0}^{\infty} \underbrace{\frac{kP_k}{\langle k \rangle}}_{\begin{subarray}{c} \text{prob. of} \\ \text{connecting to} \\ \text{a degree } k \ \text{node} \end{subarray}} \quad \underbrace{\frac{\beta_k}{\langle k \rangle}}_{\begin{subarray}{c} \text{Prob. of} \\ \text{infection} \end{subarray}} \quad \# \ \text{outgoing} \\ \# \ \text{outgoi$$

$$+\sum_{k=0}^{\infty} \frac{\widehat{kP_k}}{\langle k \rangle} \cdot \underbrace{\begin{pmatrix} 1-\beta_k \end{pmatrix}}_{\text{Prob. of no infection}} \cdot \underbrace{\begin{pmatrix} 0 \\ \text{# outgoing infected edges} \end{pmatrix}}_{\text{infected edges}}$$

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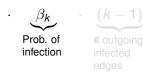
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  $\underbrace{\frac{kP_k}{\langle k \rangle}}_{\text{prob. of connecting to a degree } k \text{ node}}$   $\cdot$   $\underbrace{}_{\text{Prointegration}}_{\text{prop. of the properties}}$ 

$$\frac{\frac{k}{\langle k \rangle}}{\frac{\langle k \rangle}{\langle k \rangle}} \cdot \underbrace{\frac{\beta_k}{\text{Prob. of}}}_{\text{infection}} \cdot \underbrace{\frac{(k-1)}{\text{# outgoing}}}_{\text{infected}}$$

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Our contagion condition is then:

$$r = \sum_{k=0}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle} \beta_k > 1.$$

▶ Case 1: If  $\beta_k = 1$  then

$$r = \frac{\langle k(k-1)\rangle}{\langle k\rangle} > 1.$$

Good: This is just our giant component condition again.

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▶ Case 2: If  $\beta_k = \beta < 1$  then

$$r = \beta \frac{\langle k(k-1) \rangle}{\langle k \rangle} > 1$$

- ▶ A fraction  $(1-\beta)$  of edges do not transmit infection.
- ▶ Analogous phase transition to giant component case but critical value of  $\langle k \rangle$  is increased.
- Aka bond percolation.
- ▶ Resulting degree distribution  $P'_k$ :

$$P'_{k} = \beta^{k} \sum_{i=k}^{\infty} {i \choose k} (1-\beta)^{i-k} P_{i}.$$

▶ We can show  $F_{P'}(x) = F_P(\beta x + 1 - \beta)$ .

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- ▶ Cases 3, 4, 5, ...: Now allow  $\beta_k$  to depend on k
- Asymmetry: Transmission along an edge depends on node's degree at other end.
- ▶ Possibility:  $\beta_k$  increases with k... unlikely.
- ▶ Possibility:  $\beta_k$  is not monotonic in k... unlikely.
- ▶ Possibility:  $\beta_k$  decreases with k... hmmm
- ► The story:

  More well connected people are harder to influence.

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- ▶ Possibility:  $\beta_k$  is not monotonic in k... unlikely.
- ▶ Possibility:  $\beta_k$  decreases with k... hmmm.
- $ightharpoonup eta_k \searrow$  is a plausible representation of a simple kind of social contagion.
- ► The story:

  More well connected people are harder to influence.

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- ▶ Cases 3, 4, 5, ...: Now allow  $\beta_k$  to depend on k
- Asymmetry: Transmission along an edge depends on node's degree at other end.
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 $r = \sum_{k=1}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle} \beta_k = \sum_{k=1}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle k}$  $= \sum_{k=1}^{\infty} \frac{(k-1)P_k}{\langle k \rangle} = \frac{\langle k \rangle - 1}{\langle k \rangle} = 1 - \frac{1}{\langle k \rangle}$ 

- Since r is always less than 1, no spreading can occur for this mechanism.
- ▶ Decay of  $\beta_k$  is too fast.
- Result is independent of degree distribution.

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• Example:  $\beta_k = 1/k$ .

$$r = \sum_{k=1}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle} \beta_k = \sum_{k=1}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle k}$$
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- **Example:**  $\beta_k = H(\frac{1}{k} \phi)$  where  $0 < \phi \le 1$  is a threshold and H is the Heaviside function.
- Infection only occurs for nodes with low degree.
- Call these nodes vulnerables: they flip when only one of their friends flips

$$r = \sum_{k=1}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle} \beta_k = \sum_{k=1}^{\infty} \frac{(k-1)kP_k}{\langle k \rangle} H(\frac{1}{k} - \phi)$$

$$=\sum_{k=1}^{\lfloor \frac{1}{\theta} \rfloor} \frac{(k-1)kP_k}{\langle k \rangle} \quad \text{where } \lfloor \cdot \rfloor \text{ means floor.}$$

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▶ The contagion condition:

$$r = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} \frac{(k-1)kP_k}{\langle k \rangle} > 1.$$

- As  $\phi \to 1$ , all nodes become resilient and  $r \to 0$ .
- As  $\phi \to 0$ , all nodes become vulnerable and the contagion condition matches up with the giant component condition.
- ▶ Key: If we fix  $\phi$  and then vary  $\langle k \rangle$ , we may see two phase transitions.
- Added to our standard giant component transition, we will see a cut off in spreading as nodes become more connected.

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### Some important models (recap from CSYS 300)

- ▶ Tipping models—Schelling (1971) [8, 9, 10]
  - Simulation on checker boards.
  - Idea of thresholds
- ► Threshold models—Granovetter (1978) [7]
- ► Herding models—Bikhchandani et al. (1992) [1, 2]
  - Social learning theory, Informational cascades,...

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### Original work:

"A simple model of global cascades on random networks" D. J. Watts. Proc. Natl. Acad. Sci., 2002 [12]

- ▶ Mean field Granovetter model → network model
- Individuals now have a limited view of the world

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- Interactions between individuals now represented by a network
- Network is sparse
- Individual i has k<sub>i</sub> contacts
- ▶ Influence on each link is reciprocal and of unit weight
- **Each** individual *i* has a fixed threshold  $\phi_i$
- Individuals repeatedly poll contacts on network
- Synchronous, discrete time updating
- ▶ Individual *i* becomes active when fraction of active contacts  $a_i \ge \phi_i k_i$
- Activation is permanent (SI)

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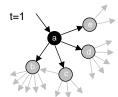
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▶ All nodes have threshold  $\phi = 0.2$ .

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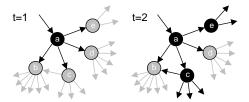
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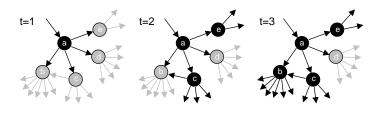


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#### Vulnerables:

- Recall definition: individuals who can be activated by just one contact being active are vulnerables.
- ▶ The vulnerability condition for node i:  $1/k_i \ge \phi_i$ .
- ▶ Means # contacts  $k_i \leq \lfloor 1/\phi_i \rfloor$ .
- ► Key: For global cascades on random networks, must have a *global component of vulnerables* [12]
- For a uniform threshold  $\phi$ , our contagion condition tells us when such a component exists:

$$r = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} \frac{(k-1)kP_k}{\langle k \rangle} > 1.$$

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- ► Key: For global cascades on random networks, must have a *global component of vulnerables* [12]
- For a uniform threshold  $\phi$ , our contagion condition tells us when such a component exists:

$$r = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} \frac{(k-1)kP_k}{\langle k \rangle} > 1.$$

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#### Vulnerables:

- Recall definition: individuals who can be activated by just one contact being active are vulnerables.
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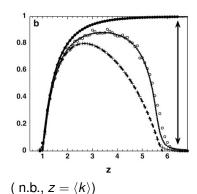
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- Top curve: final fraction infected if successful.
- Middle curve: chance of starting a global spreading event (cascade).
- Bottom curve: fractional size of vulnerable subcomponent. [12]
- Cascades occur only if size of vulnerable subcomponent > 0.
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- 'Ignorance' facilitates spreading.

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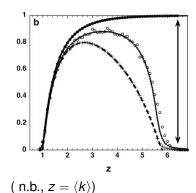
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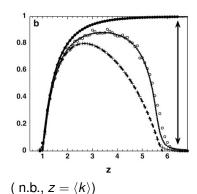
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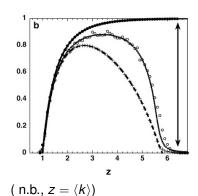
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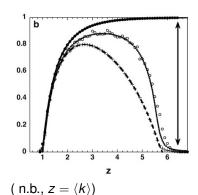
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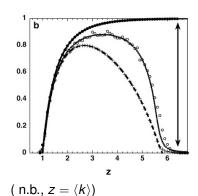
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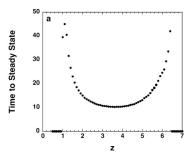
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- Time taken for cascade to spread through network. [12]
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( n.b., 
$$z = \langle k \rangle$$
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- ► Largest vulnerable component = critical mass.
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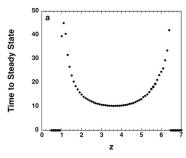
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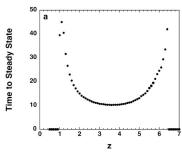
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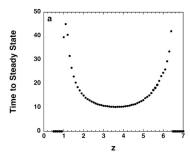
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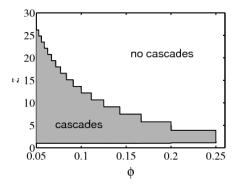
Non-spontaneous collective sheep phenomena (⊞)

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### Cascade window for random networks



(n.b., 
$$z = \langle k \rangle$$
)

Outline of cascade window for random networks.

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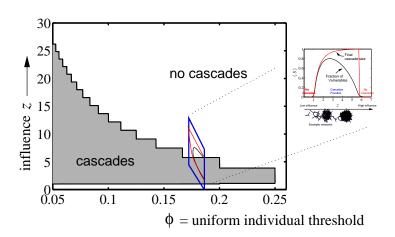
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# Cascade window for random networks



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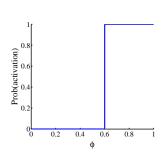
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# Granovetter's Threshold model—recap



# Assumes deterministic response functions

- $\phi_*$  = threshold of an individual.
- $f(\phi_*)$  = distribution of thresholds in a population
- ►  $F(\phi_*)$  = cumulative distribution =  $\int_{\phi_*}^{\phi_*} f(\phi_*') d\phi_*'$
- $\phi_t$  = fraction of people 'rioting' at time step t.

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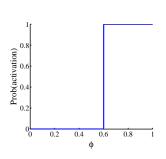
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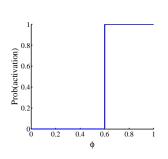
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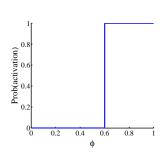
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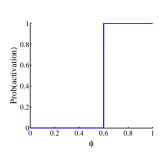
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At time t+1, fraction rioting = fraction with  $\phi_* \leq \phi_t$ .

$$\phi_{t+1} = \int_0^{\phi_t} f(\phi_*) d\phi_* = F(\phi_*)|_0^{\phi_t} = F(\phi_t)$$

 $\rightarrow$  lterative maps of the unit interval [0, 1].

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# Models

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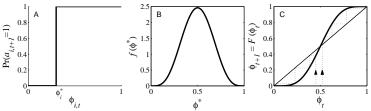
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Action based on perceived behavior of others.



- Two states: S and I
- Recover now possible (SIS)
- $\phi$  = fraction of contacts 'on' (e.g., rioting)
- Discrete time, synchronous update (strong assumption!)
- ▶ This is a Critical mass model

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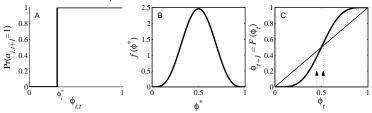
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Basic Contagion Models

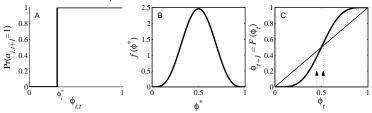
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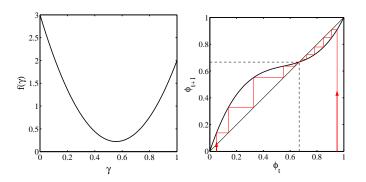
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Example of single stable state model

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# Implications for collective action theory:

- 1. Collective uniformity *⇒* individual uniformity
- Small individual changes ⇒ large global changes

#### Next

- Connect mean-field model to network model.
- Single seed for network model: 1/N → 0.
- Comparison between network and mean-field model sensible for vanishing seed size for the latter.

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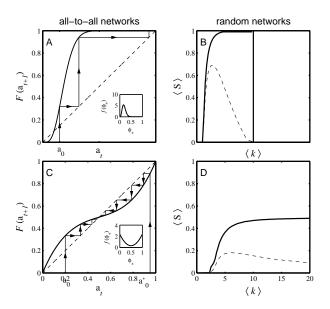
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## All-to-all versus random networks



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# Three key pieces to describe analytically:





# Threshold contagion on random networks

# Three key pieces to describe analytically:

- 1. The fractional size of the largest subcomponent of vulnerable nodes. Syuln.

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# Three key pieces to describe analytically:

- 1. The fractional size of the largest subcomponent of vulnerable nodes,  $S_{\text{vuln}}$ .
- 2. The chance of starting a global spreading event,  $P_{\text{trig}} = S_{\text{trig}}$ .
- 3. The expected final size of any successful spread, *S*.

   n.b., the distribution of *S* is almost always bimodal.

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- First goal: Find the largest component of vulnerable nodes.
- Recall that for finding the giant component's size, we had to solve:

$$F_{\pi}(x) = xF_{P}(F_{\rho}(x))$$
 and  $F_{\rho}(x) = xF_{R}(F_{\rho}(x))$ 

- We'll find a similar result for the subset of nodes that are vulnerable.
- ► This is a node-based percolation problem.
- For a general monotonic threshold distribution  $f(\phi)$ , a degree k node is vulnerable with probability

$$\beta_k = \int_0^{1/k} f(\phi) \mathrm{d}\phi.$$

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Everything now revolves around the modified generating function:

$$F_P^{(\text{vuln})}(x) = \sum_{k=0}^{\infty} \beta_k P_k x^k.$$

Generating function for friends-of-friends distribution is related in same way as before:

$$F_R^{\text{(vuln)}}(x) = \frac{\frac{d}{dx} F_P^{\text{(vuln)}}(x)}{\frac{d}{dx} F_P^{\text{(vuln)}}(x)|_{x=1}}$$

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Functional relations for component size g.f.'s are almost the same...

$$F_{\pi}^{(\text{vuln})}(x) = \underbrace{1 - F_{P}^{(\text{vuln})}(1)}_{\text{central node is not vulnerable}} + x F_{P}^{(\text{vuln})} \left( F_{\rho}^{(\text{vuln})}(x) \right)$$

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$$F_{\pi}^{(\text{vuln})}(x) = \underbrace{1 - F_{P}^{(\text{vuln})}(1)}_{\text{central node is not vulnerable}} + x F_{P}^{(\text{vuln})} \left( F_{\rho}^{(\text{vuln})}(x) \right)$$

$$F_{\rho}^{(\text{vuln})}(x) = \underbrace{1 - F_{R}^{(\text{vuln})}(1)}_{\text{first node is not}} + x F_{R}^{(\text{vuln})} \left(F_{\rho}^{(\text{vuln})}(x)\right)$$

► Can now solve as before to find  $S_{\text{vuln}} = 1 - F_{\pi}^{(\text{vuln})}(1)$ .

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#### Threshold contagion on random networks

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#### Second goal: Find probability of triggering largest vulnerable component.

- Assumption is first node is randomly chosen.
- Same set up as for vulnerable component except now we don't care if the initial node is vulnerable or not:

$$F_{\pi}^{(\text{trig})}(x) = x F_{P} \left( F_{\rho}^{(\text{vuln})}(x) \right)$$
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### Threshold contagion on random networks

- ► Third goal: Find expected fractional size of spread.
- ▶ Not obvious even for uniform threshold problem.
- ▶ Difficulty is in figuring out if and when nodes that need ≥ 2 hits switch on.
- Problem solved for infinite seed case by Gleeson and Cahalane:
  - "Seed size strongly affects cascades on random networks," Phys. Rev. E, 2007. [6]
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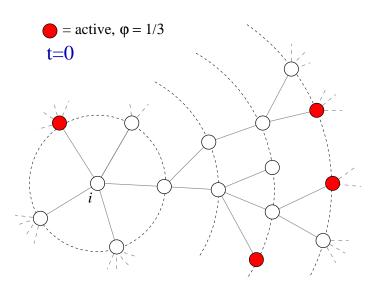
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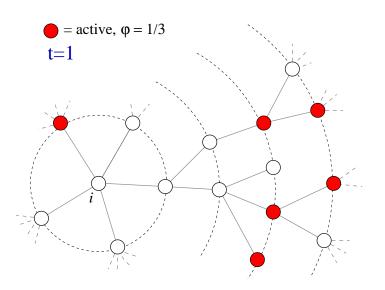
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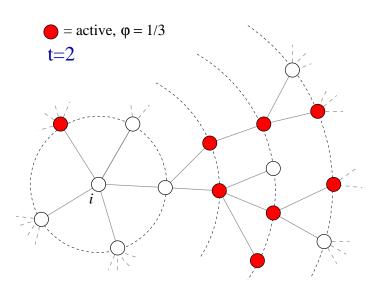
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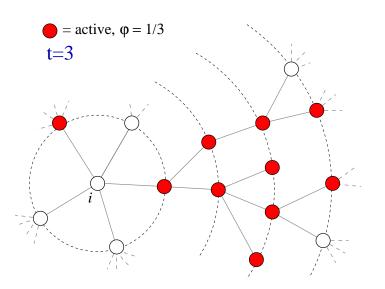
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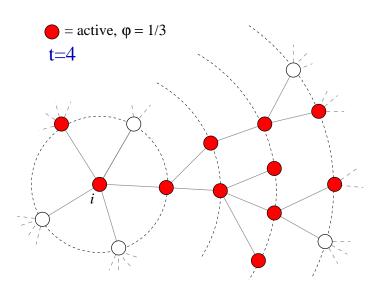
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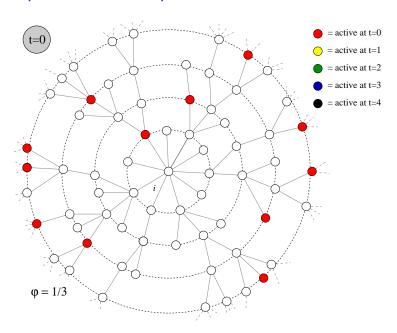
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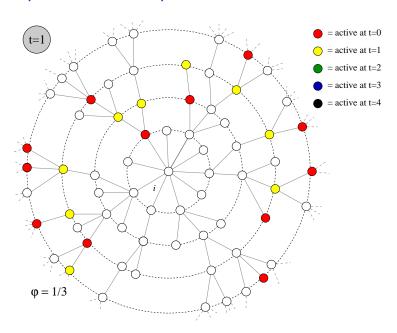
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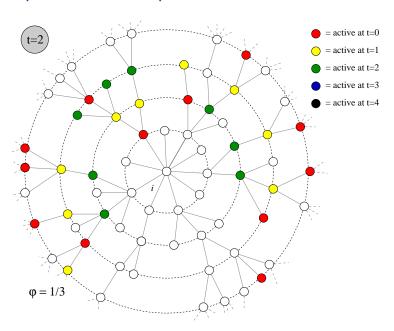
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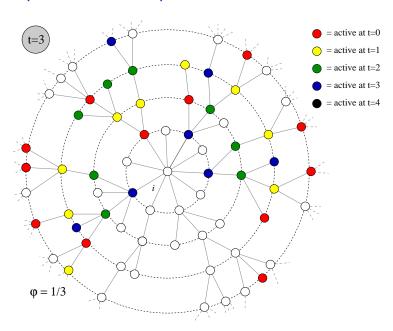
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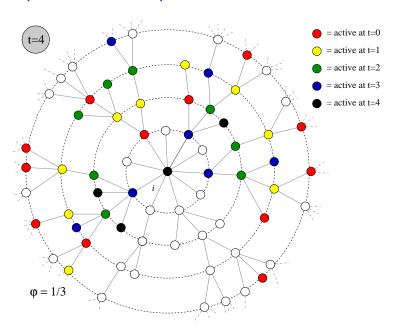
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#### Notes:

- Calculations are possible nodes do not become inactive.
- Not just for threshold model—works for a wide range of contagion processes.
- We can analytically determine the entire time evolution, not just the final size.
- ▶ We can in fact determine Pr(node of degree k switches on at time t).
- Asynchronous updating can be handled too.

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#### Pleasantness:

- Taking off from a single seed story is about expansion away from a node.
- Extent of spreading story is about contraction at a node

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- Notation: **Pr**(node *i* becomes active at time t) =  $\phi_{i,t}$ .
- Notation:  $\beta_{kj} = \mathbf{Pr}$  (a degree k node becomes active if j neighbors are active).
- ▶ Our starting point:  $\phi_{i,0} = \phi_0$ .
- $\binom{k_i}{j} \phi_0^j (1 \phi_0)^{k_i j} = \mathbf{Pr}$  (*j* of node *i*'s  $k_i$  neighbors were seeded at time t = 0).
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- For general t, we need to know the probability an edge coming into node i at time t is active.
- Notation: call this probability  $\theta_t$ .
- ▶ We already know  $\theta_0 = \phi_0$ .
- ▶ Story analogous to t = 1 case:

$$\phi_{i,t+1} = \phi_0 + (1 - \phi_0) \sum_{j=0}^{k_i} {k_j \choose j} \theta_t^j (1 - \theta_t)^{k_i - j} \beta_{k_i j}.$$

▶ Average over all nodes to obtain expression for  $\phi_{t+1}$ :

$$\phi_{t+1} = \phi_0 + (1 - \phi_0) \sum_{k=0}^{\infty} P_k \sum_{j=0}^{k} \binom{k}{j} \theta_t^j (1 - \theta_t)^{k-j} \beta_{kj}.$$

▶ So we need to compute  $\theta_t$ ...

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- ▶ Notation: call this probability  $\theta_t$ .
- ▶ We already know  $\theta_0 = \phi_0$ .
- ▶ Story analogous to t = 1 case:

$$\phi_{i,t+1} = \phi_0 + (1 - \phi_0) \sum_{j=0}^{k_i} {k_j \choose j} \theta_t^j (1 - \theta_t)^{k_i - j} \beta_{k_i j}.$$

▶ Average over all nodes to obtain expression for  $\phi_{t+1}$ :

$$\phi_{t+1} = \phi_0 + (1 - \phi_0) \sum_{k=0}^{\infty} P_k \sum_{j=0}^{k} \binom{k}{j} \theta_t^j (1 - \theta_t)^{k-j} \beta_{kj}.$$

▶ So we need to compute  $\theta_t$ ...

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▶ So we need to compute  $\theta_t$ ...

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▶ So we need to compute  $\theta_t$ ... massive excitement...

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### First connect $\theta_0$ to $\theta_1$ :

▶  $\theta_1 = \phi_0 +$ 

$$(1 - \phi_0) \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \sum_{j=0}^{k-1} {k-1 \choose j} \theta_0^{j} (1 - \theta_0)^{k-1-j} \beta_{kj}$$

- $ightharpoonup rac{kP_k}{\langle k \rangle} = R_k = \mathbf{Pr} \text{ (edge connects to a degree } k \text{ node)}.$
- ▶  $\sum_{j=0}^{k-1}$  piece gives **Pr**(degree node k activates) of its neighbors k-1 incoming neighbors are active.
- $\phi_0$  and  $(1 \phi_0)$  terms account for state of node at time t = 0.
- ▶ See this all generalizes to give  $\theta_{t+1}$  in terms of  $\theta_t$ ...

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### First connect $\theta_0$ to $\theta_1$ :

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### Two pieces:

1. 
$$\theta_{t+1} = \phi_0 +$$

$$(1 - \phi_0) \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \sum_{j=0}^{k-1} {k-1 \choose j} \theta_t^{j} (1 - \theta_t)^{k-1-j} \beta_{kj}$$

with  $\theta_0 = \phi_0$ .

2. 
$$\phi_{t+1} = \phi_0 +$$

$$(1-\phi_0)\sum_{k=0}^{\infty}P_k\sum_{j=0}^{k}\binom{k}{j}\theta_t^j(1-\theta_t)^{k-j}\beta_{kj}.$$

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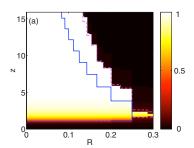
> Social Contagion Models

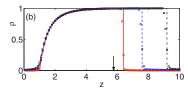
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From Gleeson and Cahalane [6]

- Pure random networks with simple threshold responses
- ► R = uniform threshold (our  $\phi_*$ ); z = average degree;  $\rho = \phi$ ;  $q = \theta$ ;  $N = 10^5$ .
- $\phi_0 = 10^{-3}, 0.5 \times 10^{-2},$  and  $10^{-2}.$
- Cascade window is for  $\phi = 10^{-2}$  case.
- Sensible expansion of cascade window as φ<sub>0</sub> increases.

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- ▶ Retrieve cascade condition for spreading from a single seed in limit  $\phi_0 \rightarrow 0$ .
- ▶ Depends on map  $\theta_{t+1} = G(\theta_t; \phi_0)$ .
- First: if self-starters are present, some activation is assured:

$$G(0; \phi_0) = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \beta_{k0} > 0.$$

meaning  $\beta_{k0} > 0$  for at least one value of  $k \ge 1$ .

If  $\theta = 0$  is a fixed point of G (i.e.,  $G(0; \phi_0) = 0$ ) then spreading occurs if

$$G'(0;\phi_0) = \frac{1}{\langle k \rangle} \sum_{k=0}^{\infty} (k-1)k P_k \beta_{k1} > 1.$$

Insert question from assignment 5 (⊞)

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#### In words:

- ▶ If  $G(0; \phi_0) > 0$ , spreading must occur because some nodes turn on for free.
- ▶ If *G* has an unstable fixed point at  $\theta = 0$ , then cascades are also always possible.

### Non-vanishing seed case:

- ightharpoonup Cascade condition is more complicated for  $\phi_0>0$ .
- If G has a stable fixed point at  $\theta=0$ , and an unstable fixed point for some  $0<\theta_*<1$ , then for  $\theta_0>\theta_*$ , spreading takes off.
- ▶ Tricky point: G depends on  $\phi_0$ , so as we change  $\phi_0$ , we also change G.

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- ▶ Cascade condition is more complicated for  $\phi_0 > 0$ .
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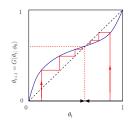
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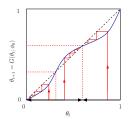
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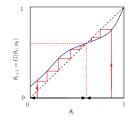
Network version All-to-all networks Theory

References









- ▶ Given  $\theta_0$ (=  $\phi_0$ ),  $\theta_\infty$  will be the nearest stable fixed point, either above or below.
- n.b., adjacent fixed points must have opposite stability types.
- ▶ Important: Actual form of *G* depends on  $\phi_0$ .
- So choice of  $\phi_0$  dictates both G and starting point—can't start anywhere for a given G.

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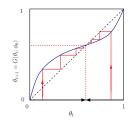
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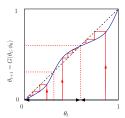
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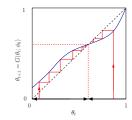
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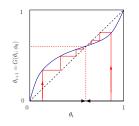
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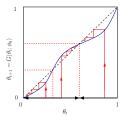
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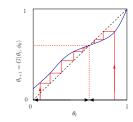
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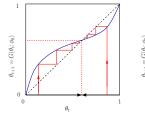
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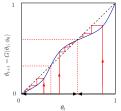
References

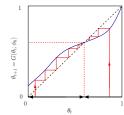
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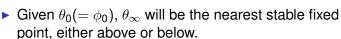
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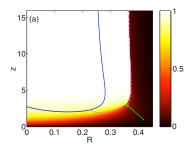
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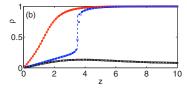
Social Contagion Models Network version

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From Gleeson and Cahalane [6]

Now allow thresholds to be distributed according to a Gaussian with mean R.

- $R = 0.2, 0.362, and 0.38; \sigma = 0.2.$
- $\phi_0 = 0$  but some nodes have thresholds  $\leq 0$  so effectively  $\phi_0 > 0$ .
- Now see a (nasty) discontinuous phase transition for low \(\lambda\rangle\).

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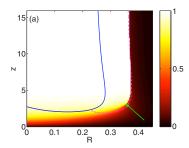
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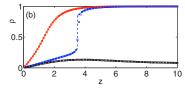
Network version All-to-all network Theory

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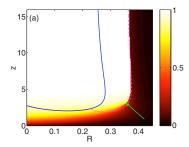
Social Contagion Models

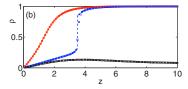
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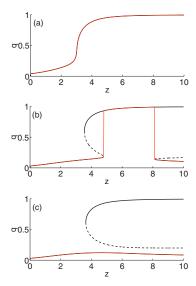
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From Gleeson and Cahalane [6]

▶ Plots of stability points for  $\theta_{t+1} = G(\theta_t; \phi_0)$ .

 n.b.: 0 is not a fixed point here: θ<sub>0</sub> = 0 always takes off.

- ► Top to bottom: R = 0.35, 0.371, and 0.375.
- n.b.: higher values of θ<sub>0</sub> for (b) and (c) lead to higher fixed points of G.
- Saddle node bifurcations appear and merge (b and c).

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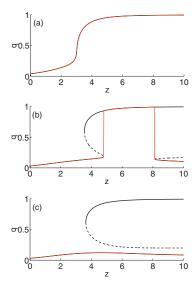
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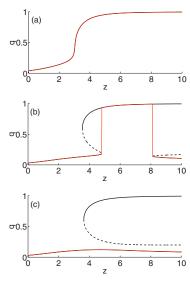
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# Comparison between theory and simulations



From Gleeson and Cahalane [6]

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 Saddle node bifurcations appear and merge (b and c). Basic Contagion Models

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## Bridging to single seed case:

- Consider largest vulnerable component as initial set of seeds.
- Not quite right as spreading must move through
- But we can usefully think of the vulnerable
- ▶ Rebuild  $\phi_t$  and  $\theta_t$  expressions...

**Basic Contagion** Models

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## Bridging to single seed case:

- Consider largest vulnerable component as initial set of seeds.
- Not quite right as spreading must move through vulnerables.
- But we can usefully think of the vulnerable
- ▶ Rebuild  $\phi_t$  and  $\theta_t$  expressions...

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### Two pieces modified for single seed:

1. 
$$\theta_{t+1} = \theta_{\text{vuln}} +$$

$$(1 - \theta_{\text{vuln}}) \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \sum_{j=0}^{k-1} {k-1 \choose j} \theta_t^{j} (1 - \theta_t)^{k-1-j} \beta_{kj}$$

with  $\theta_0 = \theta_{\text{vuln}} = \mathbf{Pr}$  an edge leads to the giant vulnerable component (if it exists).

2. 
$$\phi_{t+1} = S_{\text{vuln}} +$$

$$(1 - S_{\text{vuln}}) \sum_{k=0}^{\infty} P_k \sum_{j=0}^{k} {k \choose j} \theta_t^j (1 - \theta_t)^{k-j} \beta_{kj}.$$

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### Synchronous update

▶ Done: Evolution of  $\phi_t$  and  $\theta_t$  given exactly by the maps we have derived.

### Asynchronous updates

- ▶ Update nodes with probability  $\alpha$
- ightharpoonup As  $\alpha \to 0$ , updates become effectively independent.
- ▶ Now can talk about  $\phi(t)$  and  $\theta(t)$ .
- More on this later...

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