## Contagion

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Principles of Complex Systems, Vols. 1, 2, & 3D CSYS/MATH 6701, 6713, & a pretend number, 2023–2024 | @pocsvox

Prof. Peter Sheridan Dodds | @peterdodds

Computational Story Lab | Vermont Complex Systems Center Santa Fe Institute | University of Vermont

























The PoCSverse Contagion 1 of 88

Basic Contagion Models

Global spreading

Social Contagion Models

Network version All-to-all networks

Theory

Spreading possibility Spreading probability Physical explanation



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### Outline

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Social Contagion Models Network version All-to-all networks

### Theory

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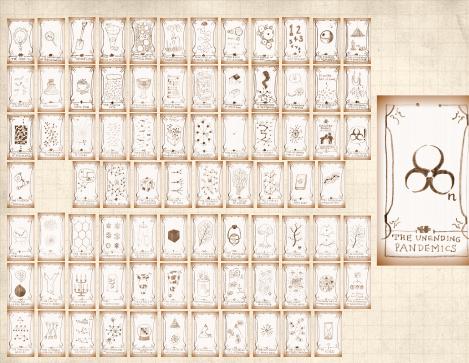
Social Contagion Models

Network version All-to-all networks

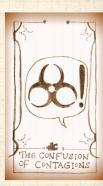
Theory

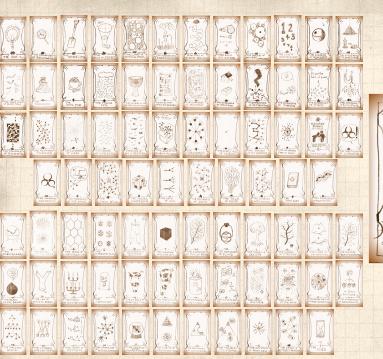
Spreading possibility Spreading probability Physical explanation Final size

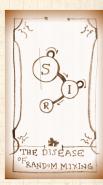


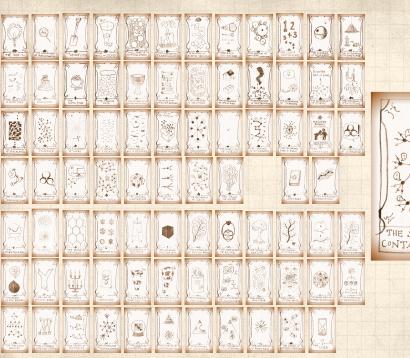


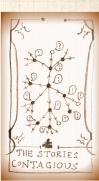


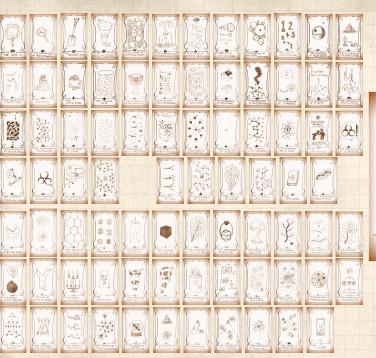


















Some large questions concerning network contagion:

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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? The PoCSverse Contagion 11 of 88

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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?
- 2. If spreading does take off, how far will it go?

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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?
- 2. If spreading does take off, how far will it go?
- 3. How do the details of the network affect the outcome?

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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?
- 2. If spreading does take off, how far will it go?
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- 4. How do the details of the spreading mechanism affect the outcome?

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- For a given spreading mechanism on a given network, what's the probability that there will be global spreading?
- 2. 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?

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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?
- 2. 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. The PoCSverse Contagion 11 of 88

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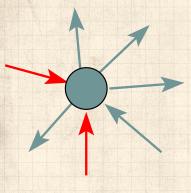
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## Spreading mechanisms



General spreading mechanism:

State of node i depends on history of i and i's neighbors' states.

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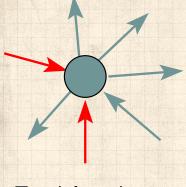
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uninfected infected



## Spreading mechanisms



General spreading mechanism:

State of node *i* depends on history of *i* and *i*'s neighbors' states.

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Doses of entity may be stochastic and history-dependent.

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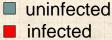
Global spreading condition

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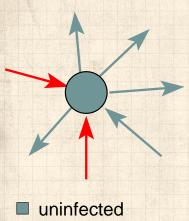
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## Spreading mechanisms



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.

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

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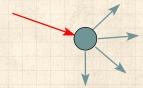
For random networks, we know local structure is pure branching.

Successful spreading is a contingent on single edges infecting nodes.

Success



Failure:



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For random networks, we know local structure is pure branching.

Successful spreading is a contingent on single edges infecting nodes.

Success Failure:



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

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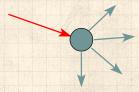


For random networks, we know local structure is pure branching.

Successful spreading is a contingent on single edges infecting nodes.

Success Failure:





- Focus on binary case with edges and nodes either infected or not.
- First big question: for a given network and contagion process, can global spreading from a single seed occur?

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We need to find: [5]

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

Call R the gain ratio.

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$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle}$$
 prob. of connecting to a degree  $k$  node

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$$\underbrace{(k-1)}_{\text{\# outgoing infected edges}}$$

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Prob. of

infection

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$$\mathbf{R} = \sum_{k=0}^{\infty} \underbrace{\frac{kP_k}{\langle k \rangle}}_{\begin{subarray}{c} \text{prob. of} \\ \text{connecting to} \\ \text{a degree $k$ node} \end{subarray}}_{\begin{subarray}{c} \text{prob. of} \\ \text{on a degree $k$ node} \end{subarray}}$$

$$\underbrace{B_{k1}}_{\text{Prob. of infection}}$$

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edges

# outgoing

infected

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Prob. of

infection



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$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{\frac{kP_k}{\langle k \rangle}}{\text{prob. of }}$$
 prob. of connecting to a degree  $k$  node

$$\underbrace{(k-1)}_{\text{\# outgoing infected edges}} \bullet \underbrace{B_{k1}}_{\text{Prob. of infection}}$$

$$+\sum_{k=0}^{\infty} \frac{\widehat{kP_k}}{\langle k \rangle}$$
 •  $\underbrace{0}_{\mbox{\# outgoing infected edges}}$  •  $\underbrace{(1-B_{k1})}_{\mbox{Prob. of no infection}}$ 

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Our global spreading condition is then:

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Our global spreading condition is then:

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} > 1.$$



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Our global spreading condition is then:

**Solution** Case 1: If  $B_{k_1} = 1$ 

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Our global spreading condition is then:

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} > 1.$$

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) = \frac{\langle k(k-1) \rangle}{\langle k \rangle} > 1.$$

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Our global spreading condition is then:

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$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) = \frac{\langle k(k-1) \rangle}{\langle k \rangle} > 1.$$

Good: This is just our giant component condition again.

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**Solution** Case 2: If 
$$B_{k1} = \beta < 1$$

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$$\red{ }$$
 Case 2: If  $B_{k1}=\beta<1$  then

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet \beta > 1.$$

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 $\clubsuit$  Case 2: If  $B_{k,1} = \beta < 1$  then

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet \beta > 1.$$



 $\clubsuit$  A fraction (1- $\beta$ ) of edges do not transmit infection.

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$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet \beta > 1.$$



 $\mathbb{A}$  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.

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- & 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 ☑.

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$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet \beta > 1.$$

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- Analogous phase transition to giant component case but critical value of  $\langle k \rangle$  is increased.
- Aka bond percolation .
- $\mathbb{R}$  Resulting degree distribution  $\tilde{P}_{k}$ :

$$\tilde{P}_k = \beta^k \sum_{i=k}^{\infty} \binom{i}{k} (1-\beta)^{i-k} P_i.$$

Insert assignment question

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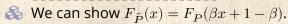


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Insert assignment question



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 $\ \$  Cases 3, 4, 5, ...: Now allow  $B_{k1}$  to depend on k

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A Cases 3, 4, 5, ...: Now allow  $B_{k1}$  to depend on k



Asymmetry: Transmission along an edge depends on node's degree at other end.

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- $\ensuremath{ \begin{subarray}{c} \& \ensuremath{ \begin{subarray}{c} Cases 3, 4, 5, ...: \ensuremath{ \begin{subarray}{c} Now allow $B_{k1}$ to depend on $k$ \end{subarray}}$
- Asymmetry: Transmission along an edge depends on node's degree at other end.
- $\ensuremath{\mathfrak{S}}$  Possibility:  $B_{k1}$  increases with k...

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Kererences



- $\clubsuit$  Cases 3, 4, 5, ...: Now allow  $B_{k1}$  to depend on k
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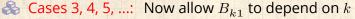
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Asymmetry: Transmission along an edge depends on node's degree at other end.

 $\clubsuit$  Possibility:  $B_{k1}$  increases with k... unlikely.

 $\mathbb{A}$  Possibility:  $B_{k1}$  is not monotonic in k...

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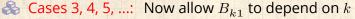
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Asymmetry: Transmission along an edge depends on node's degree at other end.

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 $\begin{cases} \&\end{cases}$  Possibility:  $B_{k1}$  decreases with k... hmmm.

 $\&B_{k1} \searrow$  is a plausible representation of a simple kind of social contagion.

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 $\ensuremath{ \begin{subarray}{ll} \& \& \ensuremath{ \begin{subarray}{ll} Cases 3, 4, 5, ...: \\ \ensuremath{ \begin{subarray}{ll} A & \ensuremath{$ 

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 $\&B_{k1} \searrow$  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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**Example:**  $B_{k1} = 1/k$ .

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 $\clubsuit$  Example:  $B_{k1} = 1/k$ .



$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1}$$

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**Basic Contagion** Models

Global spreading condition

#### Social Contagion Models

Network version All-to-all networks

#### Theory

Spreading possibility Spreading probability Physical explanation Final size





 $\clubsuit$  Example:  $B_{k1} = 1/k$ .



$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} = \sum_{k=1}^{\infty} (k-1) \bullet \frac{k P_k}{\langle k \rangle} \bullet \frac{1}{k}$$

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**Basic Contagion** Models

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 $\clubsuit$  Example:  $B_{k,1} = 1/k$ .



$$\begin{split} \mathbf{R} &= \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} = \sum_{k=1}^{\infty} (k-1) \bullet \frac{k P_k}{\langle k \rangle} \bullet \frac{1}{k} \\ &= \sum_{k=1}^{\infty} \frac{P_k}{\langle k \rangle} \bullet (k-1) \end{split}$$

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 $\clubsuit$  Example:  $B_{k1} = 1/k$ .



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Since R is always less than 1, no spreading can occur for this mechanism.

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**Basic Contagion** Models

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#### Social Contagion Models

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**Basic Contagion** Models

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$$\begin{split} \mathbf{R} &= \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} = \sum_{k=1}^{\infty} (k-1) \bullet \frac{k P_k}{\langle k \rangle} \bullet \frac{1}{k} \\ &= \sum_{k=1}^{\infty} \frac{P_k}{\langle k \rangle} \bullet (k-1) = 1 - \frac{1 - P_0}{\langle k \rangle} \end{split}$$

- Since R is always less than 1, no spreading can occur for this mechanism.
- Result is independent of degree distribution.

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Example:  $B_{k1} = H(\frac{1}{k} - \phi)$  where  $0 < \phi \le 1$  is a threshold and H is the Heaviside function lacksquare.

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- Example:  $B_{k1} = H(\frac{1}{k} \phi)$  where  $0 < \phi \le 1$  is a threshold and H is the Heaviside function  $\square$ .
- Infection only occurs for nodes with low degree.

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- $\bigotimes$  Example:  $B_{k1} = H(\frac{1}{k} \phi)$ 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.

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 $\bigotimes$  Example:  $B_{k1} = H(\frac{1}{k} - \phi)$ 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.



$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1}$$

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Example:  $B_{k1} = H(\frac{1}{k} - \phi)$  where  $0 < \phi \le 1$  is a threshold and H is the Heaviside function  $\mathbb{Z}$ .

Infection only occurs for nodes with low degree.

Call these nodes vulnerables: they flip when only one of their friends flips.



$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k\,1} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet H\left(\frac{1}{k} - \phi\right)$$

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Example:  $B_{k1} = H(\frac{1}{k} - \phi)$  where  $0 < \phi \le 1$  is a threshold and H is the Heaviside function  $\mathbb{Z}$ .

Infection only occurs for nodes with low degree.

Call these nodes vulnerables: they flip when only one of their friends flips.



$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} = \sum_{k=1}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet H \left(\frac{1}{k} - \phi\right)$$

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

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The uniform threshold model global spreading condition:

$$\mathbf{R} = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} (k-1) \bullet \frac{k P_k}{\langle k \rangle} > 1.$$

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The uniform threshold model global spreading condition:

$$\mathbf{R} = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} (k-1) \bullet \frac{k P_k}{\langle k \rangle} > 1.$$

 $\Leftrightarrow$  As  $\phi \to 1$ , all nodes become resilient and  $r \to 0$ .

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The uniform threshold model global spreading condition:

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

 $As \phi \rightarrow 1$ , all nodes become resilient and  $r \rightarrow 0$ .

As  $\phi \to 0$ , all nodes become vulnerable and the contagion condition matches up with the giant component condition.

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The uniform threshold model global spreading condition:

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 $As \phi \rightarrow 1$ , all nodes become resilient and  $r \rightarrow 0$ .

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.

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The uniform threshold model global spreading condition:

$$\mathbf{R} = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} (k-1) \bullet \frac{k P_k}{\langle k \rangle} > 1.$$

- $As \phi \rightarrow 1$ , all nodes become resilient and  $r \rightarrow 0$ .
- 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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# Virtual contagion: Corrupted Blood ☑, a 2005 virtual plague in World of Warcraft:



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## Some important models (recap from CSYS 300)



Tipping models—Schelling (1971)<sup>[11, 12, 13]</sup>

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# Some important models (recap from CSYS 300)



Tipping models—Schelling (1971) [11, 12, 13] Simulation on checker boards.

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# Some important models (recap from CSYS 300)



Tipping models—Schelling (1971) [11, 12, 13]

- Simulation on checker boards.
- Idea of thresholds.

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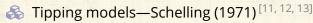
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## Some important models (recap from CSYS 300)



Simulation on checker boards.

ldea of thresholds.

Threshold models—Granovetter (1978) [8]

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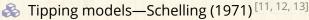
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# Some important models (recap from CSYS 300)



Simulation on checker boards.

ldea of thresholds.

A Threshold models—Granovetter (1978) [8]

Herding models—Bikhchandani et al. (1992) [1, 2]

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# Some important models (recap from CSYS 300)

- Tipping models—Schelling (1971) [11, 12, 13]
  - Simulation on checker boards.
  - ldea of thresholds.
- A Threshold models—Granovetter (1978) [8]
- A 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"

Duncan J. Watts, Proc. Natl. Acad. Sci., **99**, 5766–5771, 2002. [15] The PoCSverse Contagion 24 of 88

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Duncan J. Watts, Proc. Natl. Acad. Sci., 99, 5766-5771, 2002. [15]



Mean field Granovetter model → network model

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### Original work:



"A simple model of global cascades on random networks"

Duncan J. Watts, Proc. Natl. Acad. Sci., **99**, 5766–5771, 2002. [15]

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

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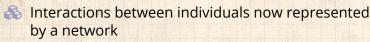
Social Contagion Models

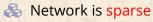
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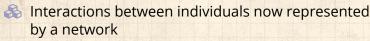
Network version

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Network is sparse

Individual i has  $k_i$  contacts

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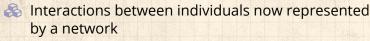
Social Contagion Models

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Network is sparse

Individual i has  $k_i$  contacts

Influence on each link is reciprocal and of unit weight

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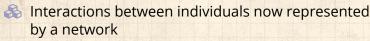
All-to-all networks

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Network is sparse

Influence on each link is reciprocal and of unit weight

 $\clubsuit$  Each individual i has a fixed threshold  $\phi_i$ 

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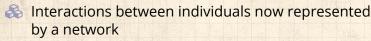
Network version All-to-all networks

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Network is sparse

 $\clubsuit$  Individual i has  $k_i$  contacts

Influence on each link is reciprocal and of unit weight

 $\red {\mathbb R}$  Each individual i has a fixed threshold  $\phi_i$ 

Individuals repeatedly poll contacts on network

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Interactions between individuals now represented by a network

Network is sparse

Individual i has  $k_i$  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

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Interactions between individuals now represented by a network

Network is sparse

Influence on each link is reciprocal and of unit weight

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Individuals repeatedly poll contacts on network

Synchronous, discrete time updating

A lndividual i becomes active when number of active contacts  $a_i \ge \phi_i k_i$ 

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Interactions between individuals now represented by a network

Network is sparse

Individual i has  $k_i$  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

A Individual i becomes active when number of active contacts  $a_i \geq \phi_i k_i$ 

Activation is permanent (SI)

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All nodes have threshold  $\phi = 0.2$ .

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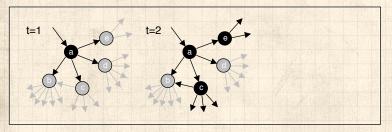
Social Contagion Models

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All nodes have threshold  $\phi = 0.2$ .

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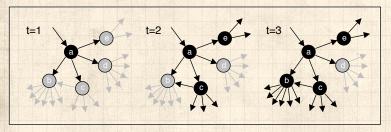
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All nodes have threshold  $\phi = 0.2$ .

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Vulnerables:

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#### Vulnerables:



Recall definition: individuals who can be activated by just one contact being active are vulnerables.

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#### Vulnerables:

Recall definition: individuals who can be activated by just one contact being active are vulnerables.

 $\clubsuit$  The vulnerability condition for node i:  $1/k_i \ge \phi_i$ .

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#### Vulnerables:

Recall definition: individuals who can be activated by just one contact being active are vulnerables.

 $\ensuremath{\&}$  The vulnerability condition for node i:  $1/k_i \geq \phi_i$ .

 $\Leftrightarrow$  Means # contacts  $k_i \leq \lfloor 1/\phi_i \rfloor$ .

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#### Vulnerables:

- Recall definition: individuals who can be activated by just one contact being active are vulnerables.
- $\ensuremath{ \Longrightarrow}$  The vulnerability condition for node i:  $1/k_i \geq \phi_i$ .
- $\red {\Bbb S}$  Means # contacts  $k_i \leq \lfloor 1/\phi_i \rfloor$ .
- Key: For global spreading events (cascades) on random networks, must have a global component of vulnerables [15]

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#### Vulnerables:

- Recall definition: individuals who can be activated by just one contact being active are vulnerables.
- $\ensuremath{ \begin{subarray}{c} \ensuremath{ \begin{subarray}{c$
- $\red{ }$  Means # contacts  $k_i \leq \lfloor 1/\phi_i \rfloor$ .
- Key: For global spreading events (cascades) on random networks, must have a global component of vulnerables [15]
- For a uniform threshold  $\phi$ , our global spreading condition tells us when such a component exists:

$$\mathbf{R} = \sum_{k=1}^{\lfloor \frac{1}{\phi} \rfloor} \frac{k P_k}{\langle k \rangle} \bullet (k-1) > 1.$$

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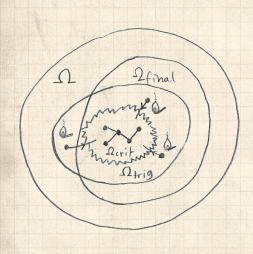
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# Example random network structure:



 $\Omega_{\rm crit}$  = critical mass = global vulnerable component

 $\Omega_{\mathrm{trig}}$  = triggering component

 $\Omega_{\text{final}} = \\ \text{potential} \\ \text{extent of} \\ \text{spread}$ 

 $\Omega$  = entire network

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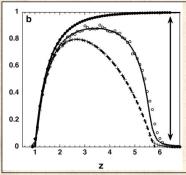
References



 $\Omega_{\mathrm{crit}} \subset \Omega_{\mathrm{trig}}; \ \Omega_{\mathrm{crit}} \subset \Omega_{\mathrm{final}}; \ \mathrm{and} \ \Omega_{\mathrm{trig}}, \Omega_{\mathrm{final}} \subset \Omega.$ 

# Global spreading events on random

networks [15]



$$z = \langle k \rangle$$



Top curve: final fraction infected if successful.

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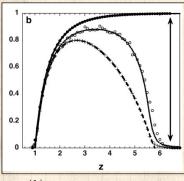
Theory

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# Global spreading events on random

networks [15]



$$z = \langle k \rangle$$



Top curve: final fraction infected if successful.

8

Bottom curve: fractional size of vulnerable subcomponent. [15]

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Basic Contagion Models

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Social Contagion Models

Network version

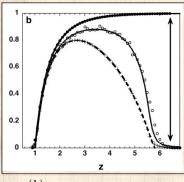
All-to-all networks

#### Theory

Spreading possibility
Spreading probability
Physical explanation
Final size



networks [15]



$$z = \langle k \rangle$$



Top curve: final fraction infected if successful.



Middle curve: chance of starting a global spreading event (cascade).



Bottom curve: fractional size of vulnerable subcomponent. [15]

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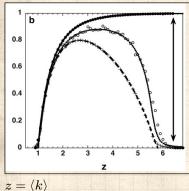
Network version All-to-all networks

Theory

Spreading probability Final size



networks [15]



8

Top curve: final fraction infected if successful.



Middle curve: chance of starting a global spreading event (cascade).



Bottom curve: fractional size of vulnerable subcomponent. [15]

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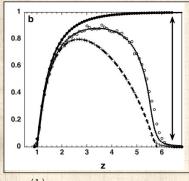
References



 $\Leftrightarrow$  Global spreading events occur only if size of vulnerable subcomponent > 0.



networks [15]



- - Top curve: final fraction infected if successful.
- Middle curve: chance of starting a global spreading event (cascade).
- - Bottom curve: fractional size of vulnerable subcomponent. [15]

$$z = \langle k \rangle$$



Global spreading events occur only if size of vulnerable subcomponent > 0.

System is robust-yet-fragile just below upper boundary [3, 4, 14]

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networks [15] b 0.8 0.6 0.4 0.2

Top curve: final fraction infected if successful.

Middle curve: chance of starting a global spreading event (cascade).



Bottom curve: fractional size of vulnerable subcomponent. [15]

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 $z = \langle k \rangle$ 

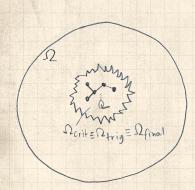
Global spreading events occur only if size of vulnerable subcomponent > 0.

System is robust-yet-fragile just below upper boundary [3, 4, 14]

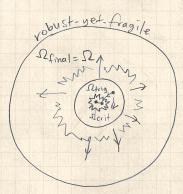


'Ignorance' facilitates spreading.





Above lower phase transition



Just below upper phase transition The PoCSverse Contagion 30 of 88

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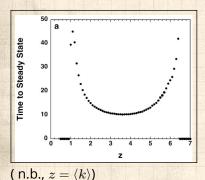
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Time taken for cascade to spread through network. [15]

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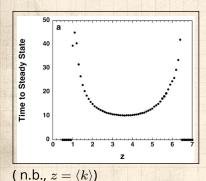
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Time taken for cascade to spread through network. [15]

Two phase transitions.

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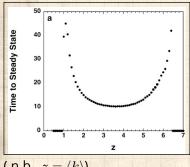
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Time taken for cascade to spread through network. [15]



Two phase transitions.

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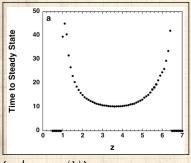
References

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



Largest vulnerable component = critical mass.





Time taken for cascade to spread through network. [15]



Two phase transitions.

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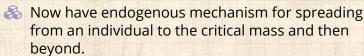
Spreading probability

References

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

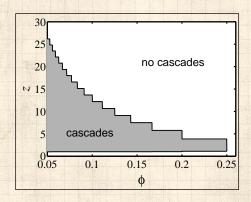


Largest vulnerable component = critical mass.





#### Cascade window for random networks



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

Outline of cascade window for random networks.

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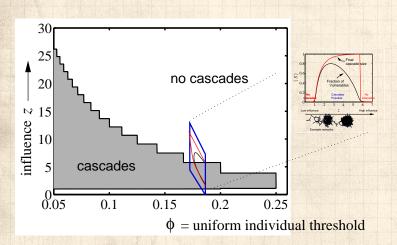
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#### Cascade window for random networks



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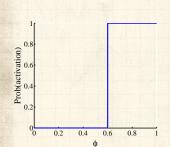
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### Granovetter's Threshold model—recap



Assumes deterministic response functions



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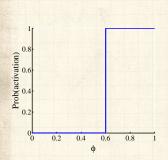
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### Granovetter's Threshold model—recap



Assumes deterministic response functions



 $\phi_*$  = threshold of an individual.

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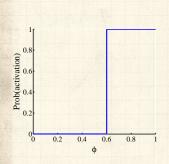
Network version All-to-all networks

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### Granovetter's Threshold model—recap



Assumes deterministic response functions



 $\phi_*$  = threshold of an individual.



 $\Leftrightarrow f(\phi_*) = distribution of$ thresholds in a population. The PoCSverse Contagion 35 of 88

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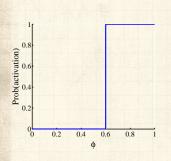
All-to-all networks

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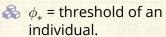
Spreading possibility Spreading probability Final size



### Granovetter's Threshold model—recap



Assumes deterministic response functions



 $\Re f(\phi_*)$  = distribution of thresholds in a population.

 $F(\phi_*)$  = cumulative distribution =  $\int_{\phi_*'=0}^{\phi_*} f(\phi_*') d\phi_*'$ 

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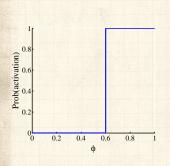
Network version All-to-all networks

Theory

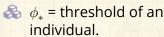
Spreading possibility Spreading probability Physical explanation Final size

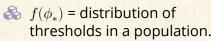


### Granovetter's Threshold model—recap



Assumes deterministic response functions





- $F(\phi_*)$  = cumulative distribution =  $\int_{\phi_*'=0}^{\phi_*} f(\phi_*') d\phi_*'$
- $\phi_t$  = fraction of people 'rioting' at time step t.

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 $\phi_* \leq \phi_t$ .

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



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

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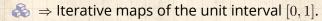




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$$\phi_{t+1} = \int_0^{\phi_t} f(\phi_*) \mathrm{d}\phi_* \, = \, F(\phi_*)|_0^{\phi_t} = F(\phi_t)$$



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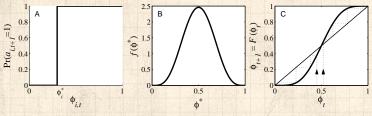
#### Theory

Spreading probability Final size





Action based on perceived behavior of others.



Two states: S and I

Recover now possible (SIS)

 $\Leftrightarrow \phi$  = fraction of contacts 'on' (e.g., rioting)

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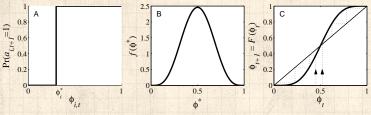
All-to-all networks

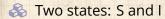
Theory

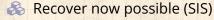
Spreading probability Final size



Action based on perceived behavior of others.







 $\Leftrightarrow \phi$  = fraction of contacts 'on' (e.g., rioting)

Discrete time, synchronous update (strong assumption!)

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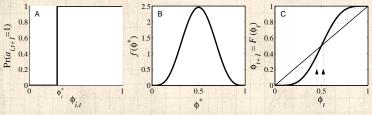
Network version
All-to-all networks

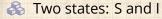
Theory

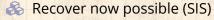
Spreading possibility Spreading probability Physical explanation Final size



Action based on perceived behavior of others.







 $\Leftrightarrow \phi$  = fraction of contacts 'on' (e.g., rioting)

Discrete time, synchronous update (strong assumption!)

This is a Critical mass model

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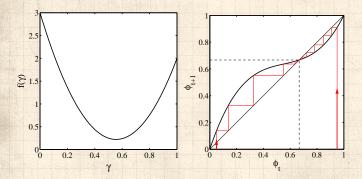
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Example of single stable state model

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Implications for collective action theory:

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## Implications for collective action theory:

1. Collective uniformity ⇒ individual uniformity

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### Implications for collective action theory:

- 1. Collective uniformity ⇒ individual uniformity
- 2. Small individual changes  $\Rightarrow$  large global changes

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### Implications for collective action theory:

- 1. Collective uniformity ⇒ individual uniformity
- 2. Small individual changes  $\Rightarrow$  large global changes

#### Next:

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### Implications for collective action theory:

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

#### Next:

Connect mean-field model to network model.

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### Implications for collective action theory:

- 1. Collective uniformity ⇒ individual uniformity
- 2. Small individual changes  $\Rightarrow$  large global changes

#### Next:

Connect mean-field model to network model.

Single seed for network model:  $1/N \rightarrow 0$ .

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## Implications for collective action theory:

- 1. Collective uniformity ⇒ individual uniformity
- 2. Small individual changes  $\Rightarrow$  large global changes

#### Next:

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

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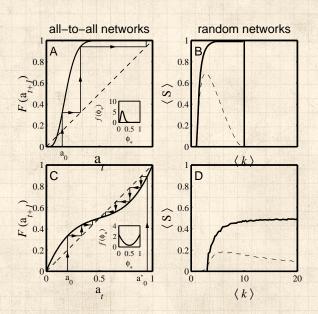
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### All-to-all versus random networks



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# Spreadworthiness: Cat videos

Bowling with Ragdolls:

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References

https://www.youtube.com/watch?v=XX-g2nmqL9Q?rel=0



Organic extreme outlier?



Success did not spread to other videos.



# Threshold contagion on random networks

Three key pieces to describe analytically:

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# Threshold contagion on random networks

### Three key pieces to describe analytically:

1. The fractional size of the largest subcomponent of vulnerable nodes,  $S_{\rm vuln}$ .

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# Threshold contagion on random networks

#### 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,  $P_{\mathsf{trig}} = S_{\mathsf{trig}}$ .

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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,  $P_{\rm trig} = S_{\rm trig}$ .
- 3. The expected final size of any successful spread, S.

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#### Three key pieces to describe analytically:

- 1. The fractional size of the largest subcomponent of vulnerable nodes,  $S_{\rm vuln}$ .
- 2. The chance of starting a global spreading event,  $P_{\rm trig} = S_{\rm trig}$ .
- 3. The expected final size of any successful spread, S.
  - n.b., the distribution of is almost always bimodal.

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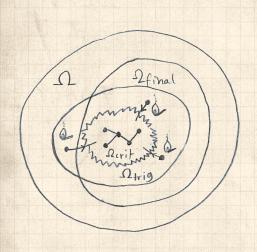
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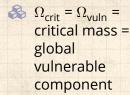
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### Example random network structure:





 $\Omega_{\mathrm{trig}}$  = triggering component

 $\Omega_{\text{final}} = \\ \text{potential} \\ \text{extent of} \\ \text{spread}$ 

 $\Omega$  = entire network

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 $\Omega_{\mathsf{crit}} \subset \Omega_{\mathsf{trig}}; \ \Omega_{\mathsf{crit}} \subset \Omega_{\mathsf{final}}; \ \mathsf{and} \ \Omega_{\mathsf{trig}}, \Omega_{\mathsf{final}} \subset \Omega.$ 

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First goal: Find the largest component of vulnerable nodes.

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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:

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

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$$F_{\pi}(x) = x F_{P}\left(F_{\rho}(x)\right) \text{ and } F_{\rho}(x) = x F_{R}\left(F_{\rho}(x)\right)$$

We'll find a similar result for the subset of nodes that are vulnerable. The PoCSverse Contagion 45 of 88

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- First goal: Find the largest component of vulnerable nodes.
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$$F_{\pi}(x) = x F_{P}\left(F_{\rho}(x)\right) \text{ and } F_{\rho}(x) = x F_{R}\left(F_{\rho}(x)\right)$$

- We'll find a similar result for the subset of nodes that are vulnerable.
- This is a node-based percolation problem.

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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:

$$F_{\pi}(x) = x F_{P}\left(F_{\rho}(x)\right) \text{ and } F_{\rho}(x) = x F_{R}\left(F_{\rho}(x)\right)$$

- 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

$$B_{k1} = \int_0^{1/k} f(\phi) \mathrm{d}\phi \,.$$

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We now have a generating function for the probability that a randomly chosen node is vulnerable and has degree k:

$$F_P^{(\mathrm{vuln})}(x) = \sum_{k=0}^\infty P_k B_{k1} x^k.$$

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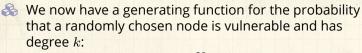
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$$F_P^{(\mathrm{vuln})}(x) = \sum_{k=0}^\infty P_k B_{k1} x^k.$$

The generating function for friends-of-friends distribution is similar to before:

$$F_R^{(\mathrm{vuln})}(x) = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} B_{k1} x^{k-1}$$

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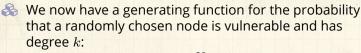
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FIIIdi Size





$$F_P^{(\mathrm{vuln})}(x) = \sum_{k=0}^\infty P_k B_{k1} x^k.$$

The generating function for friends-of-friends distribution is similar to before:

$$F_R^{(\mathrm{vuln})}(x) = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} B_{k1} x^{k-1}$$

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

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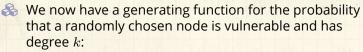
All-to-all networks

Theory

Spreading possibility
Spreading probability

Final size





$$F_P^{(\mathrm{vuln})}(x) = \sum_{k=0}^\infty P_k B_{k1} x^k.$$

The generating function for friends-of-friends distribution is similar to before:

$$F_R^{(\mathrm{vuln})}(x) = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} B_{k1} x^{k-1}$$

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

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Social Contagion Models

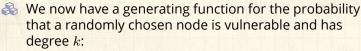
All-to-all network

Theory

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Detail: We still have the underlying degree distribution involved in the denominator.

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

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

$$F_{\pi}^{(\text{vuln})}(x) =$$

$$x F_P^{(\mathrm{vuln})} \left( F_\rho^{(\mathrm{vuln})}(x) \right)$$

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

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

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

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

$$F_{\rho}^{(\text{vuln})}(x) = x F_{R}^{(\text{vuln})} \left( F_{\rho}^{(\text{vuln})}(x) \right)$$

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

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

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

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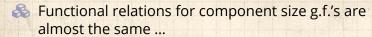
**Basic Contagion** Models

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

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

Can now solve as before to find

$$S_{\text{vuln}} = 1 - F_{\pi}^{(\text{vuln})}(1).$$

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Second goal: Find probability of triggering largest vulnerable component.

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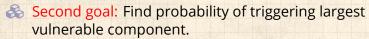
condition

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Assumption is first node is randomly chosen.

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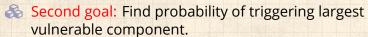
Social Contagion Models

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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:

$$\begin{split} F_{\pi}^{(\mathrm{trig})}(x) &= x \textcolor{red}{F_{P}} \left( F_{\rho}^{(\mathrm{vuln})}(x) \right) \\ F_{\rho}^{(\mathrm{vuln})}(x) &= 1 - F_{R}^{(\mathrm{vuln})}(1) + x F_{R}^{(\mathrm{vuln})} \left( F_{\rho}^{(\mathrm{vuln})}(x) \right) \end{split}$$

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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:

$$\begin{split} F_{\pi}^{(\mathrm{trig})}(x) &= x \textcolor{red}{F_{P}} \left( F_{\rho}^{(\mathrm{vuln})}(x) \right) \\ F_{\rho}^{(\mathrm{vuln})}(x) &= 1 - F_{R}^{(\mathrm{vuln})}(1) + x F_{R}^{(\mathrm{vuln})} \left( F_{\rho}^{(\mathrm{vuln})}(x) \right) \end{split}$$

 $\red Solve$  as before to find  $P_{\mathrm{trig}} = S_{\mathrm{trig}} = 1 - F_{\pi}^{(\mathrm{trig})}(1)$ .

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Possibility: binary indicator of phase. Global spreading events are either possible or can never happen.

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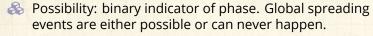
Social Contagion Models

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For random networks, global spreading possibility is understood as meaning a giant component of vulnerable nodes exists.

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- Possibility: binary indicator of phase. Global spreading events are either possible or can never happen.
- For random networks, global spreading possibility is understood as meaning a giant component of vulnerable nodes exists.
- Next: what's the probability that a randomly infected node will cause a global spreading event?

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- Possibility: binary indicator of phase. Global spreading events are either possible or can never happen.
- For random networks, global spreading possibility is understood as meaning a giant component of vulnerable nodes exists.
- Next: what's the probability that a randomly infected node will cause a global spreading event?
- $\clubsuit$  Call this  $P_{\text{trig}}$ .

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- Possibility: binary indicator of phase. Global spreading events are either possible or can never happen.
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- Next: what's the probability that a randomly infected node will cause a global spreading event?
- & Call this  $P_{\text{trig}}$ .
- As usual, it's all about edges and we need to first determine the probability that an infected edge leads to a global spreading event.

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- Possibility: binary indicator of phase. Global spreading events are either possible or can never happen.
- For random networks, global spreading possibility is understood as meaning a giant component of vulnerable nodes exists.
- Next: what's the probability that a randomly infected node will cause a global spreading event?
- & Call this  $P_{\text{trig}}$ .
- As usual, it's all about edges and we need to first determine the probability that an infected edge leads to a global spreading event.
- & Call this  $Q_{\text{trig}}$ .

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- Possibility: binary indicator of phase. Global spreading events are either possible or can never happen.
- For random networks, global spreading possibility is understood as meaning a giant component of vulnerable nodes exists.
- Next: what's the probability that a randomly infected node will cause a global spreading event?
- & Call this  $P_{\text{trig}}$ .
- As usual, it's all about edges and we need to first determine the probability that an infected edge leads to a global spreading event.
- & Call this  $Q_{\text{trig}}$ .
- Later: Generalize to more complex networks involving assortativity of all kinds.

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#### Probability an infected edge leads to a global spreading event:



 $Q_{\text{trig}}$  must satisfying a one-step recursion relation.

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# Probability an infected edge leads to a global spreading event:

 $\begin{cases} \&Q_{
m trig} & {
m must satisfying a one-step recursion relation.} \end{cases}$ 

Follow an infected edge and use three pieces:

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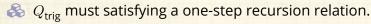
Network version All-to-all networks

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# Probability an infected edge leads to a global spreading event:



Follow an infected edge and use three pieces:

1. Probability of reaching a degree k node is  $Q_k = \frac{kP_k}{\langle k \rangle}$ .

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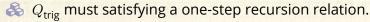
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## Probability an infected edge leads to a global spreading event:



Follow an infected edge and use three pieces:

- 1. Probability of reaching a degree k node is  $Q_k = \frac{kP_k}{\langle k \rangle}$ .
- 2. The node reached is vulnerable with probability  $B_{k+1}$ .

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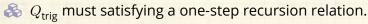
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# Probability an infected edge leads to a global spreading event:



Follow an infected edge and use three pieces:

1. Probability of reaching a degree k node is  $Q_k = \frac{kP_k}{\langle k \rangle}$ .

2. The node reached is vulnerable with probability  $B_{k1}$ .

3. At least one of the node's outgoing edges leads to a global spreading event = 1 - probability no edges do so =  $1 - (1 - Q_{\text{trio}})^{k-1}$ .

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# Probability an infected edge leads to a global spreading event:

 $\begin{cases} \&Q_{
m trig} & {
m must satisfying a one-step recursion relation.} \end{cases}$ 

Follow an infected edge and use three pieces:

- 1. Probability of reaching a degree k node is  $Q_k = \frac{kP_k}{\langle k \rangle}$ .
- 2. The node reached is vulnerable with probability  $B_{k1}$ .
- 3. At least one of the node's outgoing edges leads to a global spreading event = 1 probability no edges do so =  $1 (1 Q_{\rm trig})^{k-1}$ .

 $\red {\Bbb R}$  Put everything together and solve for  $Q_{{
m trig}}$ :

$$Q_{\mathrm{trig}} = \sum_{k} \frac{k P_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^{k-1} \right].$$

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$$Q_{\mathrm{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[1 - (1 - Q_{\mathrm{trig}})^{k-1}\right] = f(Q_{\mathrm{trig}}; P_k, B_{k1})$$

 $Q_{\text{trig}} = 0$  is always a solution.

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$$Q_{\mathrm{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[1 - (1 - Q_{\mathrm{trig}})^{k-1}\right] = f(Q_{\mathrm{trig}}; P_k, B_{k1})$$

- $\begin{cases} \&Q_{\mathsf{trig}}=0 \ \text{is always a solution.} \end{cases}$
- Spreading occurs if a second solution exists for which  $0 < Q_{\rm trig} \le 1$ .

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$$Q_{\mathrm{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[1 - (1 - Q_{\mathrm{trig}})^{k-1}\right] = f(Q_{\mathrm{trig}}; P_k, B_{k1})$$

- $\begin{cases} \&Q_{\mathsf{trig}}=0 \ \text{is always a solution.} \end{cases}$
- $\ \ \,$  Spreading occurs if a second solution exists for which  $0 < Q_{\rm trig} \leq 1.$
- $\ensuremath{\mathfrak{S}}$  Given  $P_k$  and  $B_{k1}$ , we can use any kind of root finder to solve for  $Q_{\text{trig}}$ , but ...

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$$Q_{\mathrm{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[1 - (1 - Q_{\mathrm{trig}})^{k-1}\right] = f(Q_{\mathrm{trig}}; P_k, B_{k1})$$

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- $\Leftrightarrow$  Given  $P_k$  and  $B_{k1}$ , we can use any kind of root finder to solve for  $Q_{\rm trig}$ , but ...
- & The function f increases monotonically with  $Q_{\text{trig}}$ .

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$$Q_{\mathrm{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[1 - (1 - Q_{\mathrm{trig}})^{k-1}\right] = f(Q_{\mathrm{trig}}; P_k, B_{k1})$$

- $\ \, \mathbf{\$} \,$  Spreading occurs if a second solution exists for which  $0 < Q_{\rm trig} \leq 1.$
- $\Leftrightarrow$  Given  $P_k$  and  $B_{k1}$ , we can use any kind of root finder to solve for  $Q_{\rm trig}$ , but ...
- & The function f increases monotonically with  $Q_{\mathsf{trig}}$ .
- We can therefore use an iterative cobwebbing approach to find the solution:  $Q_{\mathsf{trig}}^{(n+1)} = f(Q_{\mathsf{trig}}^{(n)}; P_k, B_{k1}).$

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$$Q_{\mathrm{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[1 - (1 - Q_{\mathrm{trig}})^{k-1}\right] = f(Q_{\mathrm{trig}}; P_k, B_{k1})$$

- $\begin{cases} \&Q_{
  m trig}=0 \ \mbox{is always a solution.} \end{cases}$
- $\ensuremath{\mathfrak{S}}$  Spreading occurs if a second solution exists for which  $0 < Q_{\rm trig} \leq 1.$
- $\Leftrightarrow$  Given  $P_k$  and  $B_{k1}$ , we can use any kind of root finder to solve for  $Q_{\rm trig}$ , but ...
- & The function f increases monotonically with  $Q_{\text{trig}}$ .
- We can therefore use an iterative cobwebbing approach to find the solution:  $Q_{\mathrm{trig}}^{(n+1)} = f(Q_{\mathrm{trig}}^{(n)}; P_k, B_{k1}).$
- $\ref{Start}$  Start with a suitably small seed  $Q_{\mathrm{trig}}^{(1)}>0$  and iterate while rubbing hands together.

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& Global spreading is possible if the fractional size  $S_{\text{vuln}}$  of the largest component of vulnerables is "giant".

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 Global spreading is possible if the fractional size  $S_{\text{vuln}}$ of the largest component of vulnerables is "giant".



Interpret  $S_{\text{vulp}}$  as the probability a randomly chosen node is vulnerable and that infecting it leads to a global spreading event:

$$S_{\mathrm{vuln}} = \sum_k P_k \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^k \right] > 0.$$

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 Global spreading is possible if the fractional size  $S_{\text{vuln}}$ of the largest component of vulnerables is "giant".



Interpret  $S_{\text{vulp}}$  as the probability a randomly chosen node is vulnerable and that infecting it leads to a global spreading event:

$$S_{\mathrm{vuln}} = \sum_k P_k \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^k \right] > 0.$$



 $\clubsuit$  Amounts to having  $Q_{\text{trig}} > 0$ .

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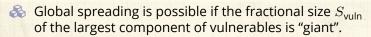
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Interpret  $S_{\text{vuln}}$  as the probability a randomly chosen node is vulnerable and that infecting it leads to a global spreading event:

$$S_{\mathrm{vuln}} = \sum_k P_k \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^k \right] > 0.$$

- $\red{\$}$  Amounts to having  $Q_{\rm trig}>0$ .
- Probability of global spreading differs only in that we don't care if the initial seed is vulnerable or not:

$$P_{\mathrm{trig}} = S_{\mathrm{trig}} = \sum_{k} P_{k} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^{k} \right]$$

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- & Global spreading is possible if the fractional size  $S_{\text{vuln}}$  of the largest component of vulnerables is "giant".
- Interpret  $S_{\rm vuln}$  as the probability a randomly chosen node is vulnerable and that infecting it leads to a global spreading event:

$$S_{\mathrm{vuln}} = \sum_k P_k \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^k \right] > 0.$$

- $\clubsuit$  Amounts to having  $Q_{\text{trig}} > 0$ .
- Probability of global spreading differs only in that we don't care if the initial seed is vulnerable or not:

$$P_{\mathrm{trig}} = S_{\mathrm{trig}} = \sum_{k} P_{k} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^{k} \right]$$

 $\clubsuit$  As for  $S_{
m vuln}$ ,  $P_{
m trig}$  is non-zero when  $Q_{
m trig}>0$ .

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 $\aleph$  We found that  $F_{\rho}^{(\text{vuln})}(1)$ —the probability that a random edge leads to a finite vulnerable component—satisfies

$$F_{\rho}^{(\mathrm{vuln})}(1) = 1 - F_R^{(\mathrm{vuln})}(1) + 1 \cdot F_R^{(\mathrm{vuln})}\left(F_{\rho}^{(\mathrm{vuln})}(1)\right).$$

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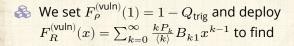
### Theory

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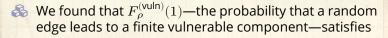
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$$F_{\rho}^{(\mathrm{vuln})}(1) = 1 - F_R^{(\mathrm{vuln})}(1) + 1 \cdot F_R^{(\mathrm{vuln})}\left(F_{\rho}^{(\mathrm{vuln})}(1)\right).$$

$$1 - Q_{\rm trig} = 1 - \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle} B_{k1} + \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle} B_{k1} \left( 1 - Q_{\rm trig} \right)^{k-1}. \label{eq:trig}$$

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We found that  $F_{\rho}^{(\text{vuln})}(1)$ —the probability that a random edge leads to a finite vulnerable component—satisfies

$$F_{\rho}^{(\mathrm{vuln})}(1) = 1 - F_R^{(\mathrm{vuln})}(1) + 1 \cdot F_R^{(\mathrm{vuln})} \left( F_{\rho}^{(\mathrm{vuln})}(1) \right).$$

 $\label{eq:weights} \begin{array}{l} \text{\&We set } F_{\rho}^{(\text{vuln})}(1) = 1 - Q_{\text{trig}} \text{ and deploy} \\ F_{R}^{(\text{vuln})}(x) = \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle} B_{k1} x^{k-1} \text{ to find} \end{array}$ 

$$1 - Q_{\rm trig} = 1 - \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} B_{k1} + \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} B_{k1} \left( 1 - Q_{\rm trig} \right)^{k-1}. \label{eq:trig}$$

Some breathless algebra it all matches:

$$Q_{\mathrm{trig}} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ 1 - \left( 1 - Q_{\mathrm{trig}} \right)^{k-1} \right].$$

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## Fractional size of the largest vulnerable component:



The generating function approach gave  $S_{\text{yuln}} = 1 - F_{\pi}^{(\text{vuln})}(1)$  where

$$F_{\pi}^{(\mathrm{vuln})}(1) = 1 - F_P^{(\mathrm{vuln})}(1) + 1 \cdot F_P^{(\mathrm{vuln})} \left( F_{\rho}^{(\mathrm{vuln})}(1) \right).$$

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# Fractional size of the largest vulnerable component:

The generating function approach gave  $S_{
m vuln} = 1 - F_\pi^{
m (vuln)}(1)$  where

$$F_\pi^{(\mathrm{vuln})}(1) = 1 - F_P^{(\mathrm{vuln})}(1) + 1 \cdot F_P^{(\mathrm{vuln})}\left(F_\rho^{(\mathrm{vuln})}(1)\right).$$

 $\label{eq:power_power} \& \text{Again using } F_{\rho}^{(\text{vuln})}(1) = 1 - Q_{\text{trig}} \text{ along with } \\ F_{P}^{(\text{vuln})}(x) = \sum_{k=0}^{\infty} P_k B_{k1} x^k \text{, we have:}$ 

$$1-S_{\mathrm{vuln}} = 1 - \sum_{k=0}^{\infty} P_k B_{k1} + \sum_{k=0}^{\infty} P_k B_{k1} \left(1 - Q_{\mathrm{trig}}\right)^k. \label{eq:spectrum}$$

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## Fractional size of the largest vulnerable component:

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Again using  $F_{\rho}^{(\text{vuln})}(1) = 1 - Q_{\text{trig}}$  along with  $F_{P}^{(\text{vuln})}(x) = \sum_{k=0}^{\infty} P_k B_{k1} x^k$ , we have:

$$1-S_{\mathrm{vuln}} = 1 - \sum_{k=0}^{\infty} P_k B_{k1} + \sum_{k=0}^{\infty} P_k B_{k1} \left(1 - Q_{\mathrm{trig}}\right)^k. \label{eq:suln}$$

Excited scrabbling about gives us, as before:

$$S_{\mathrm{vuln}} = \sum_{k=0}^{\infty} P_k B_{k1} \left[ 1 - \left( 1 - Q_{\mathrm{trig}} \right)^k \right]. \label{eq:svuln}$$

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Slight adjustment to the vulnerable component calculation.

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Slight adjustment to the vulnerable component calculation.

$$\Re S_{\mathsf{trig}} = 1 - F_{\pi}^{(\mathsf{trig})}(1)$$
 where

$$F_{\pi}^{(\mathrm{trig})}(1) = 1 \cdot F_{P}\left(F_{\rho}^{(\mathrm{vuln})}(1)\right).$$

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- Slight adjustment to the vulnerable component calculation.
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We play these cards:  $F_{
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$$1 - S_{\mathsf{trig}} = 1 + \sum_{k=0}^{\infty} P_k \left( 1 - Q_{\mathsf{trig}} \right)^k.$$

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$$1 - S_{\mathsf{trig}} = 1 + \sum_{k=0}^{\infty} P_k \left( 1 - Q_{\mathsf{trig}} \right)^k.$$

More scruffing around brings happiness:

$$S_{\rm trig} = \sum_{k=0}^{\infty} P_k \left[ 1 - \left( 1 - Q_{\rm trig} \right)^k \right]. \label{eq:Strig}$$

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& Earlier, we showed the global spreading condition follows from the gain ratio  $\mathbf{R} > 1$ :

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} > 1.$$

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We would very much like to see that  ${\bf R}>1$  matches up with  $Q_{\rm trig}>0$ .

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- It really would be just so totally awesome.

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- Must come from our basic edge triggering probability equation:

$$Q_{\mathrm{trig}} = \sum_k \frac{k P_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^{k-1} \right].$$

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 $\ \ \, \ \ \, \ \ \, \ \, \ \,$  When does this equation have a solution  $0 < Q_{\rm trig} \le 1?$ 

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Earlier, we showed the global spreading condition follows from the gain ratio  $\mathbf{R} > 1$ :

$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} > 1.$$

- We would very much like to see that  ${\bf R}>1$  matches up with  $Q_{\rm trig}>0$ .
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- Must come from our basic edge triggering probability equation:

$$Q_{\mathrm{trig}} = \sum_k \frac{k P_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ 1 - (1 - Q_{\mathrm{trig}})^{k-1} \right].$$

- $\red{\$}$  When does this equation have a solution  $0 < Q_{\mathrm{trig}} \leq 1$ ?
- $\red{\$}$  We need to find out what happens as  $Q_{\rm trig} o 0.^{[9]}$

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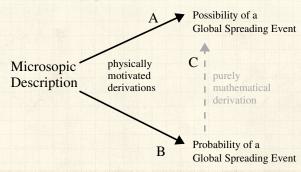
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## What we're doing:



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$$Q_{\mathrm{trig}} = \sum_{k} \frac{k P_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ 1 - \left( 1 - (k-1) Q_{\mathrm{trig}} + \ldots \right) \right]$$

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$$Q_{\mathrm{trig}} = \sum_{k} \frac{kP_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \mathbb{1} + \left( \mathbb{1} + (k-1)Q_{\mathrm{trig}} + \ldots \right) \right]$$

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$$\begin{split} Q_{\mathrm{trig}} &= \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \cancel{1} + \left( \cancel{1} + (k-1) Q_{\mathrm{trig}} + \ldots \right) \right] \\ \\ &\Rightarrow Q_{\mathrm{trig}} = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet (k-1) Q_{\mathrm{trig}} \end{split}$$

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$$\begin{split} Q_{\mathrm{trig}} &= \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \cancel{1} + \left( \cancel{1} + (k-1) Q_{\mathrm{trig}} + \ldots \right) \right] \\ \\ &\Rightarrow Q_{\mathrm{trig}} = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet (k-1) Q_{\mathrm{trig}} \\ \\ &\Rightarrow 1 = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} \end{split}$$

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 $lap{8}$  For  $Q_{
m trig} 
ightarrow 0^+$ , equation tends towards

$$\begin{split} Q_{\mathrm{trig}} &= \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \cancel{1} + \left( \cancel{1} + (k-1) Q_{\mathrm{trig}} + \ldots \right) \right] \\ &\Rightarrow Q_{\mathrm{trig}} = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet (k-1) Q_{\mathrm{trig}} \\ &\Rightarrow 1 = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} \end{split}$$

3 Only defines the phase transition points (i.e.,  $\mathbf{R} = 1$ ).

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 $\Leftrightarrow$  For  $Q_{\text{trig}} \to 0^+$ , equation tends towards

$$\begin{split} Q_{\mathrm{trig}} &= \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \cancel{1} + \left( \cancel{1} + (k-1) Q_{\mathrm{trig}} + \ldots \right) \right] \\ \\ &\Rightarrow Q_{\mathrm{trig}} = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet (k-1) Q_{\mathrm{trig}} \\ \\ &\Rightarrow 1 = \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} \end{split}$$

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Inequality?

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 $\mbox{\&}$  Again take  $Q_{\rm trig} \rightarrow 0^+$ , but keep next higher order term:

$$Q_{\mathrm{trig}} = \sum_{k} \frac{k P_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ 1 - \left( 1 - (k-1) Q_{\mathrm{trig}} + \binom{k-1}{2} Q_{\mathrm{trig}}^2 \right) \right]$$

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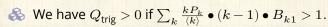
Again take  $Q_{\text{trig}} \rightarrow 0^+$ , but keep next higher order term:

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 $\Leftrightarrow$  We have  $Q_{\text{trig}} > 0$  if  $\sum_{k} \frac{kP_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} > 1$ .

Again take  $Q_{\text{trig}} \to 0^+$ , but keep next higher order term:

$$\begin{split} Q_{\mathrm{trig}} &= \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \cancel{1} + \left( \cancel{1} + (k-1) Q_{\mathrm{trig}} - \binom{k-1}{2} Q_{\mathrm{trig}}^{2} \right) \right] \\ \Rightarrow Q_{\mathrm{trig}} &= \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet B_{k1} \bullet \left[ (k-1) Q_{\mathrm{trig}} - \binom{k-1}{2} Q_{\mathrm{trig}}^{2} \right] \\ \Rightarrow \sum_{k} \frac{k P_{k}}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} &= 1 + \sum_{k} \frac{k P_{k}}{\langle k \rangle} B_{k1} \binom{k-1}{2} Q_{\mathrm{trig}} \end{split}$$



Repeat: Above is a mathematical connection between two physically derived equations.

& Again take  $Q_{\mathsf{trig}} \to 0^+$ , but keep next higher order term:

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- Repeat: Above is a mathematical connection between two physically derived equations.
- From this connection, we don't know anything about a gain ratio R or how to arrange the pieces.

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Third goal: Find expected fractional size of spread.

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Third goal: Find expected fractional size of spread. Not obvious even for uniform threshold problem.

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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.

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Problem solved for infinite seed case by Gleeson and Cahalane:

"Seed size strongly affects cascades on random networks," Phys. Rev. E, 2007. [7]

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Developed further by Gleeson in "Cascades on correlated and modular random networks," Phys. Rev. E, 2008. [6]



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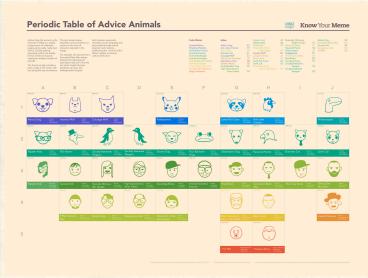
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Idea:

Randomly turn on a fraction  $\phi_0$  of nodes at time t=0

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### Idea:



Randomly turn on a fraction  $\phi_0$  of nodes at time t=0



Capitalize on local branching network structure of random networks (again)

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### Idea:



Randomly turn on a fraction  $\phi_0$  of nodes at time t=0



Capitalize on local branching network structure of random networks (again)



Now think about what must happen for a specific node i to become active at time t:

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Randomly turn on a fraction  $\phi_0$  of nodes at time t=0



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Now think about what must happen for a specific node i to become active at time t:

• t=0: i is one of the seeds (prob =  $\phi_0$ )

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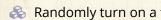
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### Idea:



Randomly turn on a fraction  $\phi_0$  of nodes at time t=0

Capitalize on local branching network structure of random networks (again)

Now think about what must happen for a specific node i to become active at time t:

- t=0: i is one of the seeds (prob =  $\phi_0$ )
- t = 1: i was not a seed but enough of i's friends switched on at time t = 0 so that i's threshold is now exceeded.

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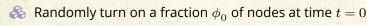
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### Idea:



- Capitalize on local branching network structure of random networks (again)
- Now think about what must happen for a specific node *i* to become active at time *t*:
  - t=0: i is one of the seeds (prob =  $\phi_0$ )
  - t=1: i was not a seed but enough of i's friends switched on at time t=0 so that i's threshold is now exceeded.
  - t=2: enough of i's friends and friends-of-friends switched on at time t=0 so that i's threshold is now exceeded.

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### Idea:

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  - Randomly turn on a fraction  $\phi_0$  of nodes at time t=0
- Capitalize on local branching network structure of random networks (again)
- Now think about what must happen for a specific node *i* to become active at time *t*:
  - t=0: i is one of the seeds (prob =  $\phi_0$ )
  - t=1: i was not a seed but enough of i's friends switched on at time t=0 so that i's threshold is now exceeded.
  - t=2: enough of i's friends and friends-of-friends switched on at time t=0 so that i's threshold is now exceeded.
  - t = n: enough nodes within n hops of i switched on at t = 0 and their effects have propagated to reach i.

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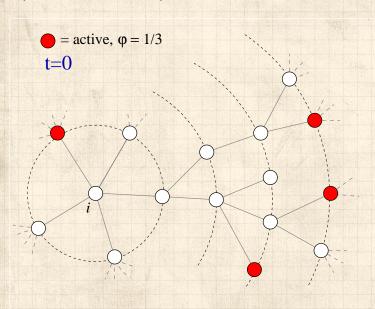
Network version All-to-all networks

Theory

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Spreading probability
Physical explanation

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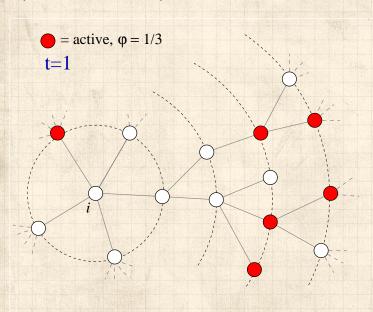
Network version All-to-all networks

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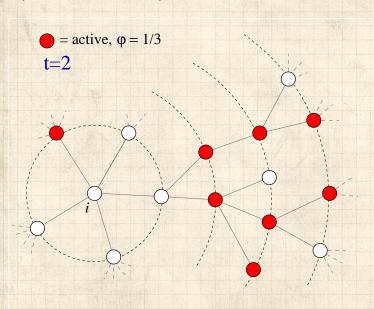
Network version All-to-all networks

#### Theory

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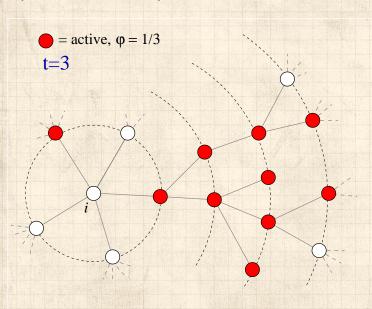
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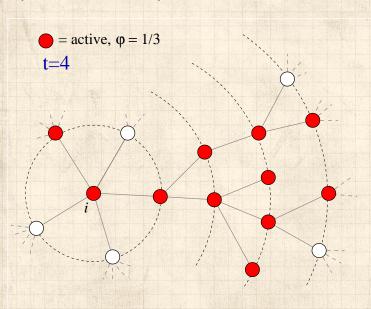
Network version All-to-all networks

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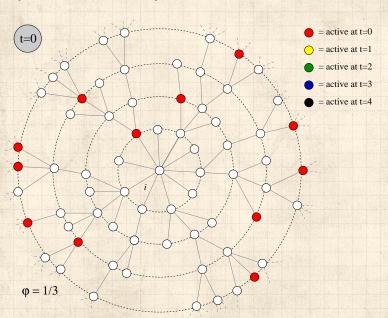
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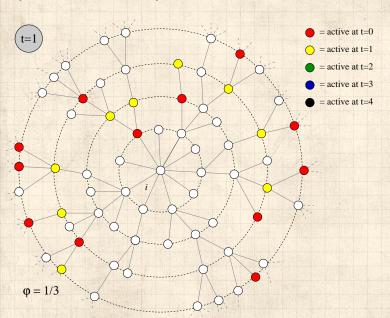
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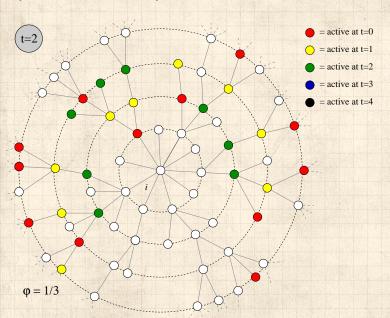
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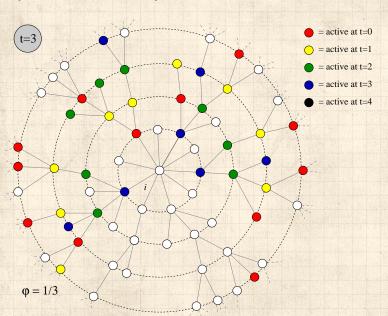
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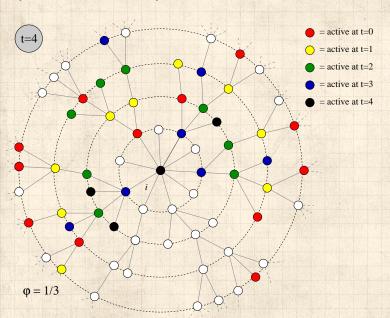
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### Notes:



Calculations presume nodes do not become inactive (strong restriction, liftable)

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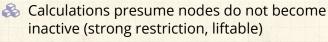
Theory

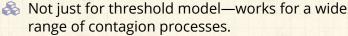
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#### Notes:





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#### Notes:

- Calculations presume nodes do not become inactive (strong restriction, liftable)
- 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.

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#### Notes:

- Calculations presume nodes do not become inactive (strong restriction, liftable)
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- 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).

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#### Notes:

- Calculations presume nodes do not become inactive (strong restriction, liftable)
- 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).
- $\clubsuit$  Even more, we can compute: **Pr**(specific node i switches on at time t).

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#### Notes:

- Calculations presume nodes do not become inactive (strong restriction, liftable)
- 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).
- Even more, we can compute: Pr(specific node i)switches on at time t).
- Asynchronous updating can be handled too.

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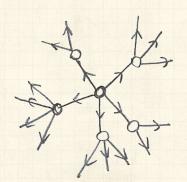
Spreading probability

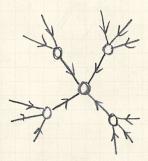
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#### Pleasantness:

Taking off from a single seed story is about expansion away from a node.





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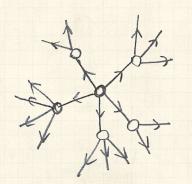
Spreading possibility Spreading probability Physical explanation

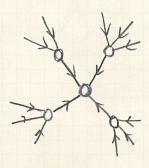
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#### Pleasantness:

- 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:

 $\phi_{k,t} = \mathbf{Pr}(\text{a degree } k \text{ node is active at time } t).$ 

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#### Notation:

 $\phi_{k,t} = \mathbf{Pr}(\text{a degree } k \text{ node is active at time } t).$ 



Notation:  $B_{k,i} = \mathbf{Pr}$  (a degree k node becomes active if j neighbors are active).

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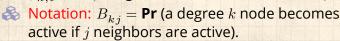
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#### Notation:

 $\phi_{k,t} = \mathbf{Pr}(\mathbf{a} \text{ degree } k \text{ node is active at time } t).$ 



 $\mbox{\&}$  Our starting point:  $\phi_{k,0} = \phi_0$ .

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 $\phi_{k,t} = \mathbf{Pr}(\mathbf{a} \text{ degree } k \text{ node is active at time } t).$ 

Notation:  $B_{kj} = \mathbf{Pr}$  (a degree k node becomes active if j neighbors are active).

 $\red {}$  Our starting point:  $\phi_{k,0}=\phi_0.$ 

 $\binom{k}{j}\phi_0^{j}(1-\phi_0)^{k-j}=\Pr\left(j\text{ of a degree }k\text{ node's neighbors were seeded at time }t=0\right).$ 

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Notation:

 $\phi_{k,t} = \mathbf{Pr}(\mathbf{a} \text{ degree } k \text{ node is active at time } t).$ 

Notation:  $B_{kj} = \mathbf{Pr}$  (a degree k node becomes active if j neighbors are active).

 $\red { }$  Our starting point:  $\phi_{k,0}=\phi_0.$ 

 $(k \choose j) \phi_0^j (1 - \phi_0)^{k-j} = \mathbf{Pr} (j \text{ of a degree } k \text{ node's neighbors were seeded at time } t = 0).$ 

Probability a degree k node was a seed at t = 0 is  $\phi_0$  (as above).

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Notation:

 $\phi_{k,t} = \mathbf{Pr}(\mathbf{a} \text{ degree } k \text{ node is active at time } t).$ 

Notation:  $B_{kj} = \mathbf{Pr}$  (a degree k node becomes active if j neighbors are active).

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Probability a degree k node was not a seed at t = 0 is  $(1 - \phi_0)$ .

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 $\phi_{k,t} = \mathbf{Pr}(\mathbf{a} \text{ degree } k \text{ node is active at time } t).$ 

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 $\red { }$  Our starting point:  $\phi_{k,0}=\phi_0.$ 

 $(k \choose j) \phi_0^j (1 - \phi_0)^{k-j} = \mathbf{Pr} (j \text{ of a degree } k \text{ node's neighbors were seeded at time } t = 0).$ 

Probability a degree k node was a seed at t = 0 is  $\phi_0$  (as above).

Probability a degree k node was not a seed at t = 0 is  $(1 - \phi_0)$ .

Combining everything, we have:

$$\phi_{k,1} = \phi_0 + (1 - \phi_0) \sum_{j=0}^k \binom{k}{j} \phi_0^j (1 - \phi_0)^{k-j} B_{kj}.$$

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 For general t, we need to know the probability an edge coming into a degree k node at time t is active.

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For general t, we need to know the probability an edge coming into a degree k node at time t is active.

 $\red > Notation$ : call this probability  $heta_t$ .

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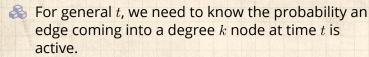
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 $\triangle$  Notation: call this probability  $\theta_t$ .

 $\clubsuit$  We already know  $\theta_0 = \phi_0$ .

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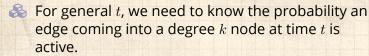
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 $lap{Notation:}$  call this probability  $\theta_t$ .

 $\red { } ext{ } ext{We already know } heta_0 = \phi_0.$ 

3 Story analogous to t = 1 case. For specific node i:

$$\phi_{i,\,t+1} = \frac{\phi_0}{\phi_0} + \frac{(1-\phi_0)}{\sum_{j=0}^{k_i} \binom{k_i}{j} \theta_t^{\,j} (1-\theta_t)^{k_i-j} B_{k_i j}.$$

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For general t, we need to know the probability an edge coming into a degree k node at time t is active.

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

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

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

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For general t, we need to know the probability an edge coming into a degree k node at time t is active.

 $\red {8}$  Notation: call this probability  $\theta_t$ .

 $\red {\$}$  We already know  $heta_0 = \phi_0$ .

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

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

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

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

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

$$\theta_1 = \phi_0 +$$

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

- $\stackrel{kP_k}{\langle k \rangle} = Q_k$  = **Pr** (edge connects to a degree k node).
- $\sum_{j=0}^{k-1}$  piece gives **Pr** (degree node k activates if j of its k-1 incoming neighbors are active).
- $\ \ \phi_0$  and  $(1-\phi_0)$  terms account for state of node at time t=0.

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- $\stackrel{kP_k}{\langle k \rangle} = Q_k$  = **Pr** (edge connects to a degree k node).
- $\sum_{j=0}^{k-1}$  piece gives **Pr** (degree node k activates if j of its k-1 incoming neighbors are active).
- $\prescript{\&} \phi_0$  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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Two pieces: edges first, and then nodes

1. 
$$\theta_{t+1} = \underbrace{\phi_0}_{\text{exogenous}}$$

$$+(1-\phi_0)\underbrace{\sum_{k=1}^{\infty}\frac{kP_k}{\langle k\rangle}\sum_{j=0}^{k-1}\binom{k-1}{j}\theta_t^{\ j}(1-\theta_t)^{k-1-j}B_{kj}}_{\text{social effects}}$$

with  $\theta_0 = \phi_0$ .

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

$$\underbrace{\phi_0}_{\text{exogenous}} + (1 - \phi_0) \underbrace{\sum_{k=0}^{\infty} P_k \sum_{j=0}^k \binom{k}{j} \theta_t^{\,j} (1 - \theta_t)^{k-j} B_{kj}}_{\text{social effects}}.$$

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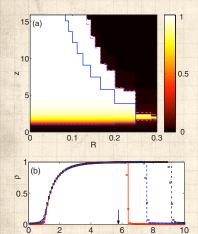
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# Comparison between theory and simulations



From Gleeson and Cahalane [7]

Pure random networks with simple threshold responses

R = uniform threshold(our  $\phi_*$ ); z = averagedegree;  $\rho = \phi$ ;  $q = \theta$ ;  $N = 10^5$ .

 $\phi_0 = 10^{-3}, 0.5 \times 10^{-2},$ and  $10^{-2}$ .

Cascade window is for  $\phi_0 = 10^{-2}$  case.

Sensible expansion of cascade window as  $\phi_0$ increases.

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Retrieve cascade condition for spreading from a single seed in limit  $\phi_0 \to 0$ .

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 $\ \, \hbox{$ \ \, $ \ \,$ 

 $\red {f R}$  Depends on map  $heta_{t+1} = G( heta_t;\phi_0)$ .

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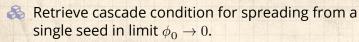
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 $\red {\Bbb S}$  Depends on map  $heta_{t+1} = G( heta_t;\phi_0).$ 

First: if self-starters are present, some activation is assured:

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

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

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Retrieve cascade condition for spreading from a single seed in limit  $\phi_0 \to 0$ .

 $\triangle$  Depends on map  $\theta_{t+1} = G(\theta_t; \phi_0)$ .

First: if self-starters are present, some activation is assured:

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

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

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

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

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Insert assignment question

### In words:



 $\Re$  If  $G(0; \phi_0) > 0$ , spreading must occur because some nodes turn on for free.

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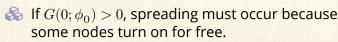
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#### In words:



 $\Re$  If G has an unstable fixed point at  $\theta = 0$ , then cascades are also always possible.

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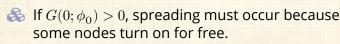
#### Theory

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Final size



#### In words:



### Non-vanishing seed case:

& Cascade condition is more complicated for  $\phi_0 > 0$ .

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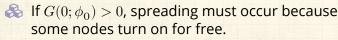
Theory

Spreading possibility Spreading probability Physical explanation

Final size



#### In words:



### Non-vanishing seed case:

 $\ensuremath{\&}$  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.

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#### In words:

- $\Leftrightarrow$  If  $G(0; \phi_0) > 0$ , spreading must occur because some nodes turn on for free.

### Non-vanishing seed case:

- $\ensuremath{\&}$  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.
- $\begin{cases} \ragged Fricky point: $G$ depends on $\phi_0$, so as we change $\phi_0$, we also change $G$.$

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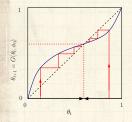
Network version All-to-all networks

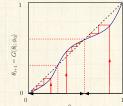
Theory

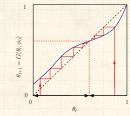
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Given  $\theta_0(=\phi_0)$  ,  $\theta_\infty$  will be the nearest stable fixed point, either above or below.

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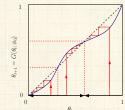
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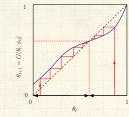
Spreading possibility Spreading probability Physical explanation Final size

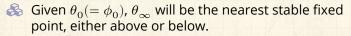
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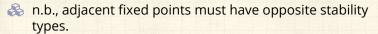












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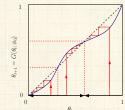
Theory

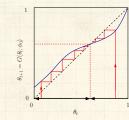
Spreading possibility
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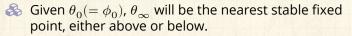
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n.b., adjacent fixed points must have opposite stability types.

 $\ensuremath{\&}$  Important: Actual form of G depends on  $\phi_0$ .

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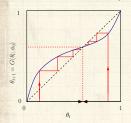
All-to-all network

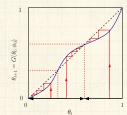
Theory

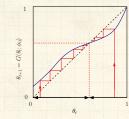
Spreading possibility
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- Given  $\theta_0(=\phi_0)$ ,  $\theta_\infty$  will be the nearest stable fixed point, either above or below.
- n.b., adjacent fixed points must have opposite stability types.

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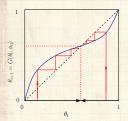
Theory

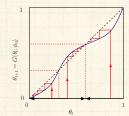
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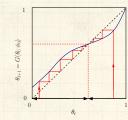
References



## General fixed point story:







- 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:  $\phi_t$  can only increase monotonically so  $\phi_0$  must shape G so that  $\phi_0$  is at or above an unstable fixed point.
- $\clubsuit$  First reason:  $\phi_1 \ge \phi_0$ .

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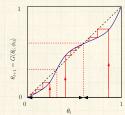
References

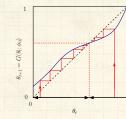
Final size



## General fixed point story:







- 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.
- $\ensuremath{\&}$  Important: Actual form of G depends on  $\phi_0$ .
- $\clubsuit$  First reason:  $\phi_1 \ge \phi_0$ .
- $\S$  Second:  $G'(\theta; \phi_0) \ge 0, 0 \le \theta \le 1$ .

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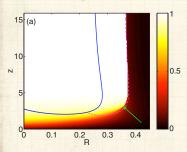
All-to-all networks

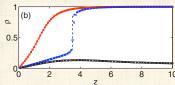
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Now allow thresholds to be distributed according to a Gaussian with mean R.



R = 0.2, 0.362, and0.38;  $\sigma = 0.2$ .

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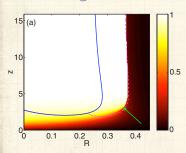
Spreading possibility Spreading probability

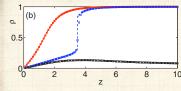
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References

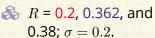
From Gleeson and Cahalane [7]







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



 $\phi_0 = 0$  but some nodes have thresholds  $\leq 0$  so effectively  $\phi_0 > 0$ .

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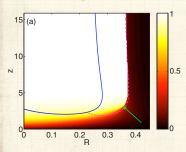
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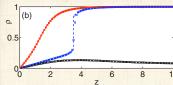
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References

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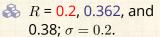






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Now allow thresholds to be distributed according to a Gaussian with mean R.



 $\phi_0 = 0$  but some nodes have thresholds  $\leq 0$  so effectively  $\phi_0 > 0$ .

Now see a (nasty) discontinuous phase transition for low  $\langle k \rangle$ . The PoCSverse Contagion 78 of 88

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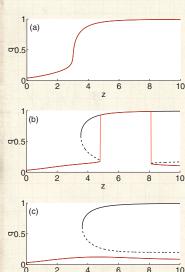
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From Gleeson and Cahalane [7]

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

- n.b.: 0 is not a fixed point here:  $\theta_0 = 0$  always takes off.
- Top to bottom: R = 0.35, 0.371, and 0.375.
- Saddle node bifurcations appear and merge (b and c).

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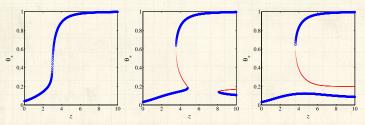
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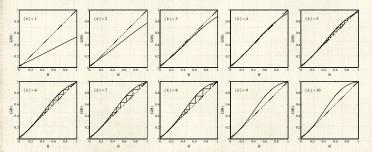
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## What's happening:



 $\ \, \ \,$  Fixed points slip above and below the  $\theta_{t+1}=\theta_t$  line:



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Synchronous update

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### Synchronous update



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

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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  $\alpha$ .

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### Synchronous update

#### Asynchronous updates

- $\red {\Bbb S}$  Update nodes with probability lpha.
- As  $\alpha \to 0$ , updates become effectively independent.

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### Synchronous update

#### Asynchronous updates

- $\ensuremath{\mathfrak{S}}$  Update nodes with probability  $\alpha$ .
- As  $\alpha \to 0$ , updates become effectively independent.
- $\red {8}$  Now can talk about  $\phi(t)$  and  $\theta(t)$ .

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Solid dive into understanding contagion on generalized random networks.

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- Solid dive into understanding contagion on generalized random networks.
- Threshold model leads to idea of vulnerables and a critical mass. [16, 8]

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- Much can be generalized for more realistic kinds of networks: degree-correlated, modular, bipartite, ...

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- The single seed contagion condition and triggering probability can be fully developed using a physical story. [5, 9]

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- Much can be generalized for more realistic kinds of networks: degree-correlated, modular, bipartite, ...
- The single seed contagion condition and triggering probability can be fully developed using a physical story. [5, 9]
- Many connections to other kinds of models: Voter models, Ising models, ...

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# Neural reboot (NR):

Pangolin happiness:

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References

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