

# Contagion

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Principles of Complex Systems, Vols. 1, 2, & 3D  
CSYS/MATH 300, 303, & 394, 2022-2023 | @pocsvox

Prof. Peter Sheridan Dodds | @peterdodds

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



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Basic Contagion  
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Global spreading  
condition

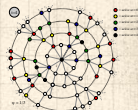
Social Contagion  
Models

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All-to-all networks

Theory

Spreading possibility  
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Final size

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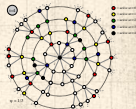
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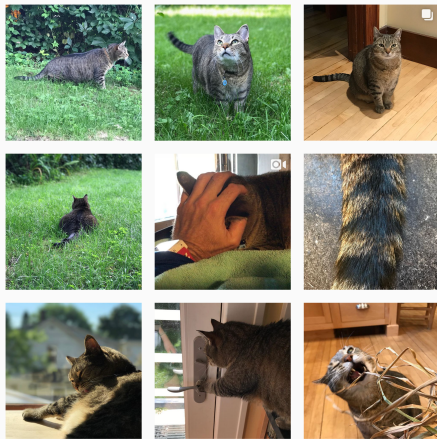
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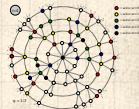
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## Global spreading condition

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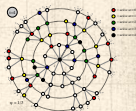
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# Contagion models

Some large questions concerning network contagion:

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## Basic Contagion Models

Global spreading condition

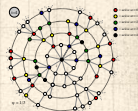
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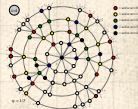
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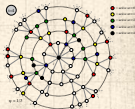
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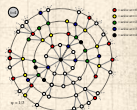
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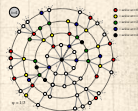
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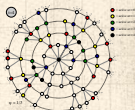
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**Next up:** We'll look at some fundamental kinds of spreading on generalized random networks.

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## Basic Contagion Models

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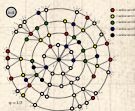
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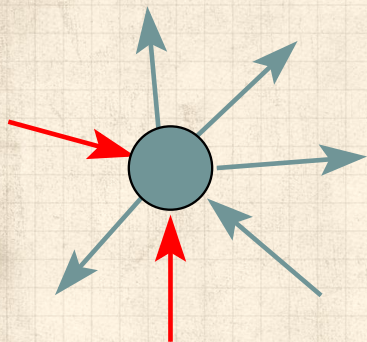
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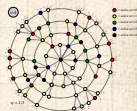
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**General spreading mechanism:**

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

■ uninfected  
■ infected



# Spreading mechanisms

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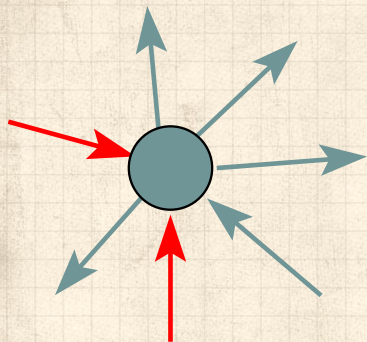
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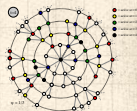
State of node  $i$  depends on history of  $i$  and  $i$ 's neighbors' states.



**Doses** of entity may be stochastic and history-dependent.

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# Spreading mechanisms

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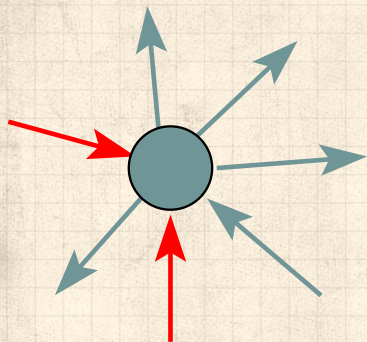
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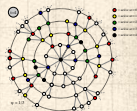


**Doses** of entity may be stochastic and history-dependent.



May have **multiple, interacting entities** spreading at once.

■ uninfected  
■ infected



# Spreading on Random Networks



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

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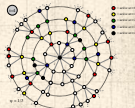
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# Spreading on Random Networks

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

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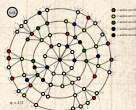
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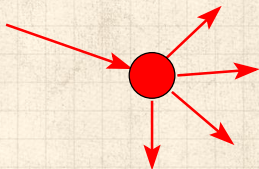


# Spreading on Random Networks

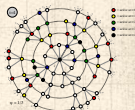
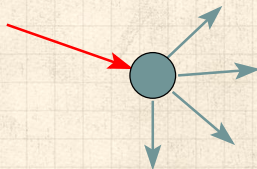
For random networks, we know local structure is pure branching.

Successful spreading is  $\therefore$  contingent on **single edges** infecting nodes.

Success



Failure:

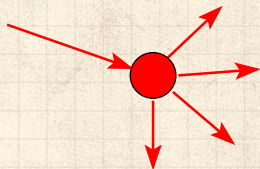


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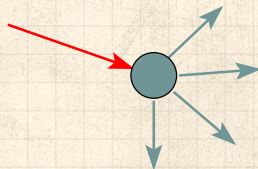
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Focus on **binary** case with edges and nodes either infected or not.

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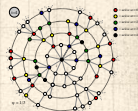
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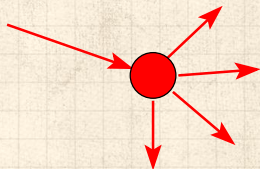


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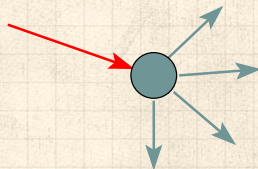
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**First big question:** for a given network and contagion process, can global spreading from a single seed occur?

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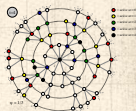
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# Global spreading condition



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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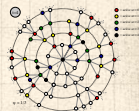
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Define  $B_{k1}$  as the probability that a node of degree  $k$  is infected by a single infected edge.

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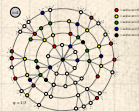
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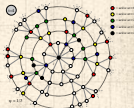
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 Define  $B_{k1}$  as the probability that a node of degree  $k$  is infected by a single infected edge.



$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{kP_k}{\langle k \rangle}$$

prob. of  
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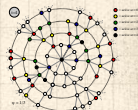


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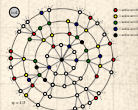


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$$\mathbf{R} = \sum_{k=0}^{\infty} \frac{k P_k}{\langle k \rangle} \bullet \underbrace{(k-1)}_{\substack{\# \text{ outgoing} \\ \text{infected} \\ \text{edges}}} \bullet \underbrace{B_{k1}}_{\substack{\text{Prob. of} \\ \text{infection}}}$$

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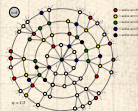
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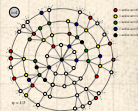
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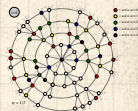
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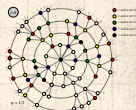
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Our global spreading condition is then:

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



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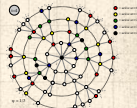


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Case 1:



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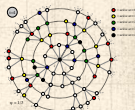
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 **Case 1:** If  $B_{k1} = 1$



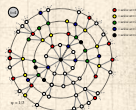
# Global spreading condition

 Our global spreading condition is then:

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

 **Case 1:** If  $B_{k1} = 1$  then

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



# Global spreading condition

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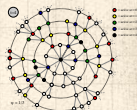
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 **Good:** This is just our giant component condition again.



# Global spreading condition



## Case 2:

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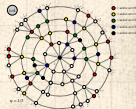
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# Global spreading condition

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

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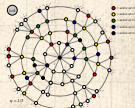
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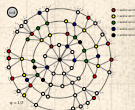
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 A fraction  $(1-\beta)$  of edges do not transmit infection.

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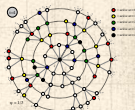
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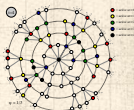
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 Analogous phase transition to giant component case but **critical value** of  $\langle k \rangle$  is **increased**.




# Global spreading condition

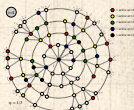
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 Aka bond percolation .






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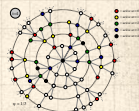
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 Aka bond percolation .


 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 question from assignment 9 






# Global spreading condition


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

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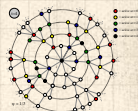
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Insert question from assignment 9 

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



# Global spreading condition



Cases 3, 4, 5, ...:

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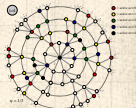
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# Global spreading condition

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

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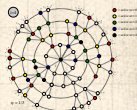
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# Global spreading condition

-  **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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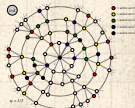
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# Global spreading condition

- Cases 3, 4, 5, ...: Now allow  $B_{k1}$  to depend on  $k$
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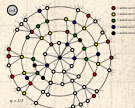
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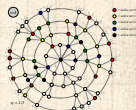
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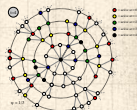
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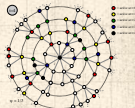
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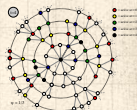
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- $B_{k1} \searrow$  is a plausible representation of a simple kind of social contagion.

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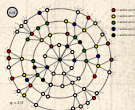
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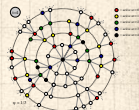
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- Possibility:  $B_{k1}$  decreases with  $k$ ... hmmm.
- $B_{k1} \searrow$  is a plausible representation of a simple kind of social contagion.
- The story:  
More well connected people are harder to influence.





# Global spreading condition



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

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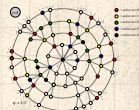
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# Global spreading condition



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



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

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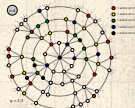
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# Global spreading condition



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



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

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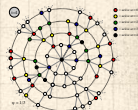
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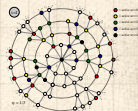
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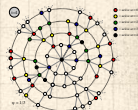
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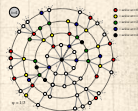
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# Global spreading condition



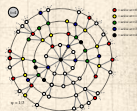
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Since  $\mathbf{R}$  is always less than 1, no spreading can occur for this mechanism.



# Global spreading condition



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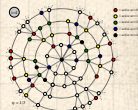
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Decay of  $B_{k1}$  is too fast.





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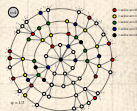


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 Since  $\mathbf{R}$  is always less than 1, no spreading can occur for this mechanism.

 Decay of  $B_{k1}$  is too fast.


 Result is independent of degree distribution.



# Global spreading condition



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

where  $0 < \phi \leq 1$  is a **threshold** and  $H$  is the Heaviside function .

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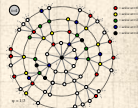
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Infection only occurs for nodes with **low** degree.

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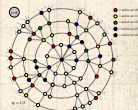
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# Global spreading condition



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where  $0 < \phi \leq 1$  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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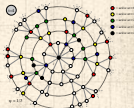
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# Global spreading condition



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

where  $0 < \phi \leq 1$  is a **threshold** and  $H$  is the Heaviside function ↗.



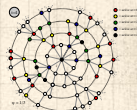
Infection only occurs for nodes with **low** degree.



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




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



# Global spreading condition



**Example:**  $B_{k1} = H\left(\frac{1}{k} - \phi\right)$

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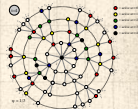
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$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{kP_k}{\langle k \rangle} \cdot (k-1) \cdot B_{k1} = \sum_{k=1}^{\infty} \frac{kP_k}{\langle k \rangle} \cdot (k-1) \cdot H\left(\frac{1}{k} - \phi\right)$$



# Global spreading condition



**Example:**  $B_{k1} = H\left(\frac{1}{k} - \phi\right)$

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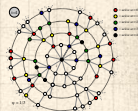


Call these nodes **vulnerables**:  
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$$\mathbf{R} = \sum_{k=1}^{\infty} \frac{kP_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} = \sum_{k=1}^{\infty} \frac{kP_k}{\langle k \rangle} \bullet (k-1) \bullet H\left(\frac{1}{k} - \phi\right)$$

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



# Global spreading condition



The uniform threshold model global spreading condition:

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

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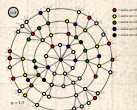
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# Global spreading condition

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

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

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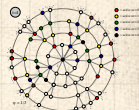
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 As  $\phi \rightarrow 1$ , all nodes become resilient and  $r \rightarrow 0$ .

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

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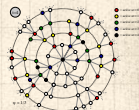
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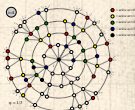


# Global spreading condition

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

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

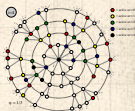


# Global spreading condition

- The uniform threshold model global spreading condition:

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

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



# Virtual contagion: Corrupted Blood, a 2005 virtual plague in World of Warcraft:



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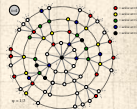
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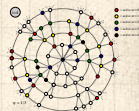
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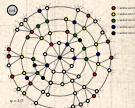
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Some important models (recap from CSYS 300)

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



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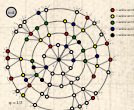
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 Simulation on checker boards.





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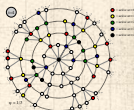
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-  Simulation on checker boards.
-  Idea of thresholds.



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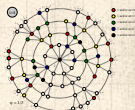
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-  Idea of thresholds.

 Threshold models—Granovetter (1978) [8]



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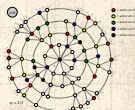
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- 🧱 Threshold models—Granovetter (1978) [8]
- 🧱 Herding models—Bikhchandani et al. (1992) [1, 2]



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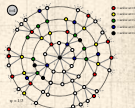
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  - 🧱 Idea of thresholds.
- 🧱 Threshold models—Granovetter (1978) [8]
- 🧱 Herding models—Bikhchandani et al. (1992) [1, 2]
  - 🧱 Social learning theory, Informational cascades,...



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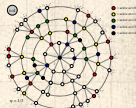
References

Original work:



"A simple model of global cascades on  
random networks" 

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



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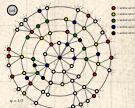


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Mean field Granovetter model → network model



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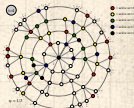


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 Mean field Granovetter model → network model

 Individuals now have a limited view of the world



# Threshold model on a network

 Interactions between individuals now represented by a network

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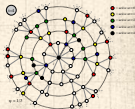
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# Threshold model on a network

 Interactions between individuals now represented by a network

 Network is **sparse**

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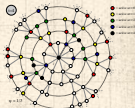
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# Threshold model on a network

- Interactions between individuals now represented by a network
- Network is **sparse**
- Individual  $i$  has  $k_i$  contacts

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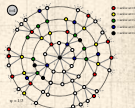
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# Threshold model on a network

- 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**

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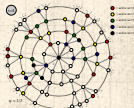
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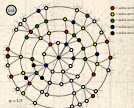
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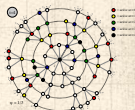
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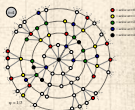
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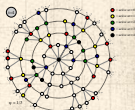
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- Individuals repeatedly poll contacts on network
- Synchronous, discrete time updating
- Individual  $i$  becomes active when number of active contacts  $a_i \geq \phi_i k_i$
- Activation is permanent (SI)

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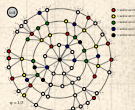
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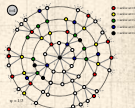
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 All nodes have threshold  $\phi = 0.2$ .



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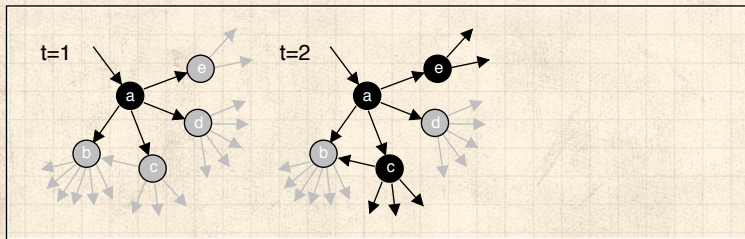
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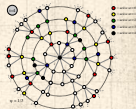
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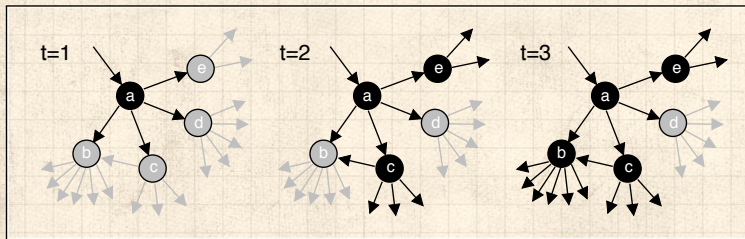
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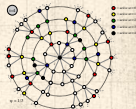
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# The most gullible

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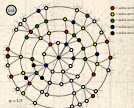
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# The most gullible

## Vulnerables:

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

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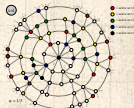
Spreading possibility

Spreading probability

Physical explanation


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# The most gullible

## Vulnerables:

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

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

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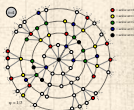
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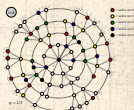
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# The most gullible

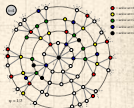
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 **Key:** For global spreading events (cascades) on random networks, must have a *global component of vulnerables* <sup>[15]</sup>








# The most gullible


## Vulnerables:

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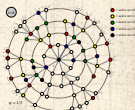
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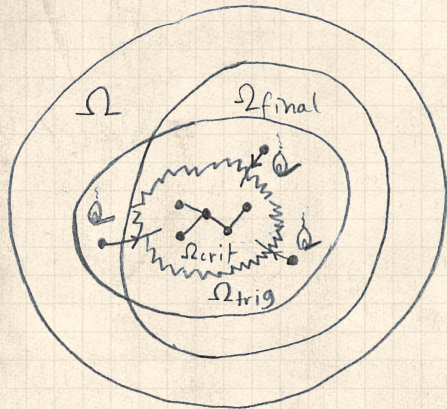
 **Key:** For global spreading events (cascades) on random networks, must have a *global component of vulnerables* <sup>[15]</sup>


 For a uniform threshold  $\phi$ , our global spreading condition tells us when such a component exists:


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





# Example random network structure:



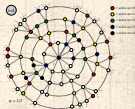
  $\Omega_{\text{crit}}$  = critical mass = global vulnerable component

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

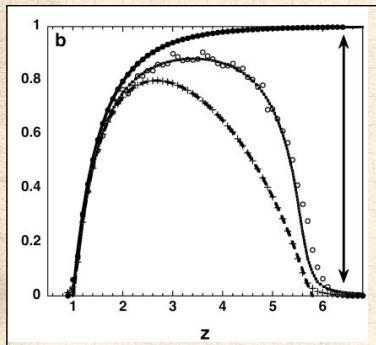
  $\Omega_{\text{final}}$  = potential extent of spread

  $\Omega$  = entire network

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



# Global spreading events on random networks <sup>[15]</sup>



$$z = \langle k \rangle$$



**Top curve:** final fraction infected if successful.

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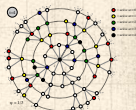
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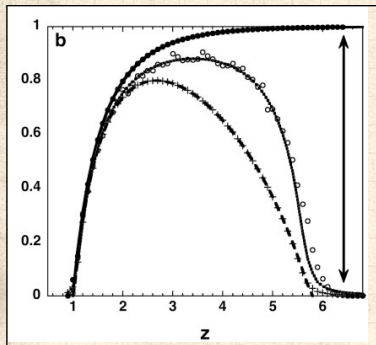
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# Global spreading events on random networks <sup>[15]</sup>



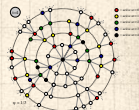
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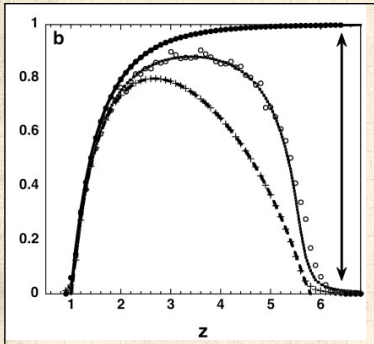
**Top curve:** final fraction infected if successful.



**Bottom curve:** fractional size of vulnerable subcomponent. <sup>[15]</sup>



# Global spreading events on random networks <sup>[15]</sup>



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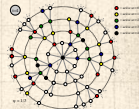
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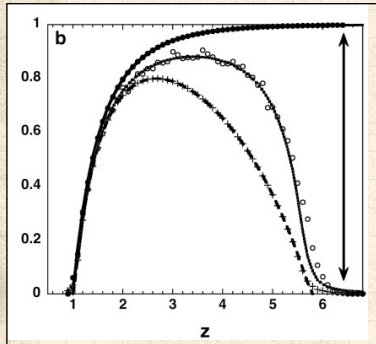
**Middle curve:** chance of starting a global spreading event (cascade).



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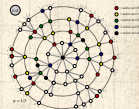
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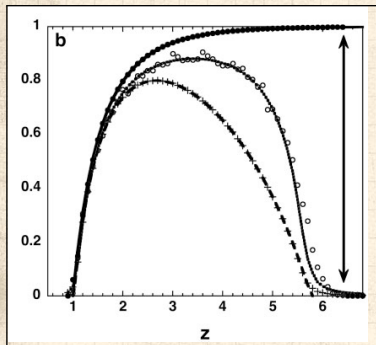
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Global spreading events occur only if size of vulnerable subcomponent  $> 0$ .





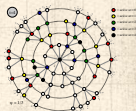
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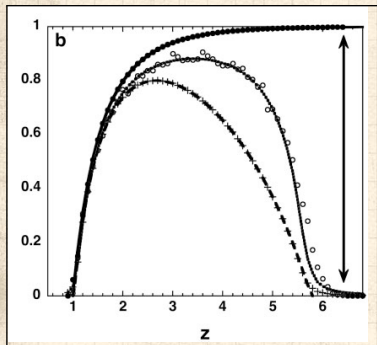
$$z = \langle k \rangle$$

-  **Top curve:** final fraction infected if successful.
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-  Global spreading events occur only if size of vulnerable subcomponent  $> 0$ .
-  System is robust-yet-fragile just below upper boundary <sup>[3, 4, 14]</sup>





# Global spreading events on random networks <sup>[15]</sup>





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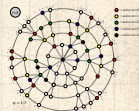
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 'Ignorance' facilitates spreading.





# Cascades on random networks

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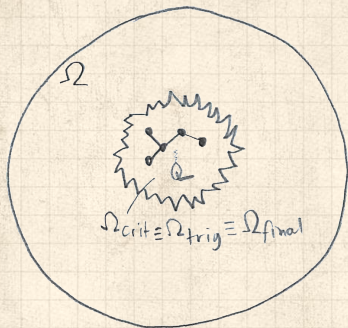
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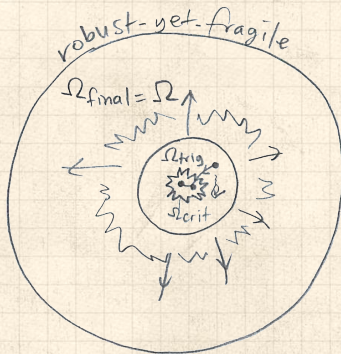
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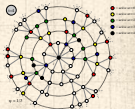
References



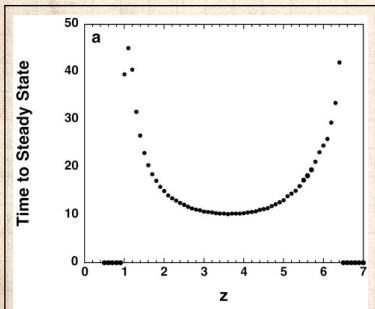
Above lower phase  
transition



Just below upper  
phase transition



# Cascades on random networks



Time taken for cascade to spread through network. <sup>[15]</sup>

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

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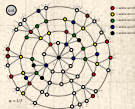
Social Contagion  
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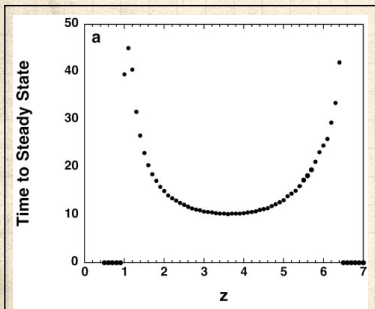
Theory

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# Cascades on random networks



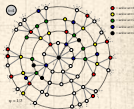
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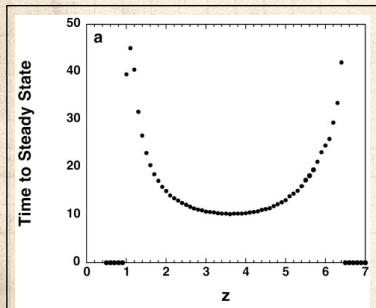
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Two phase transitions.



# Cascades on random networks



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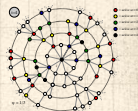


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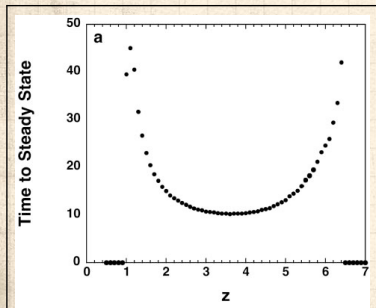
(n.b.,  $z = \langle k \rangle$ )



Largest vulnerable component = **critical mass**.



# Cascades on random networks



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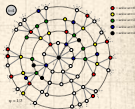
(n.b.,  $z = \langle k \rangle$ )



Largest vulnerable component = **critical mass**.



Now have endogenous mechanism for spreading from an individual to the critical mass and then beyond.



# Cascade window for random networks

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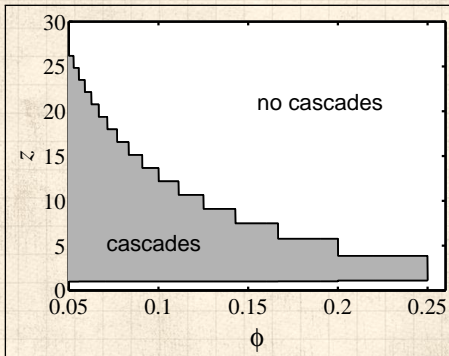
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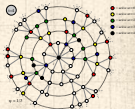
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 Outline of cascade window for random networks.



# Cascade window for random networks

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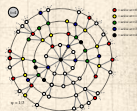
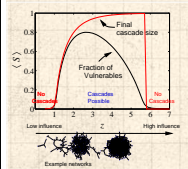
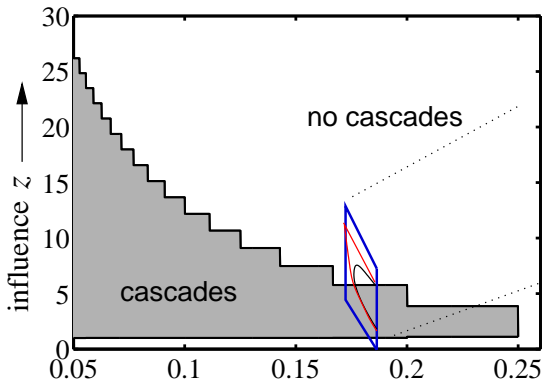
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$\phi$  = uniform individual threshold

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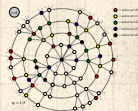
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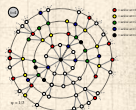
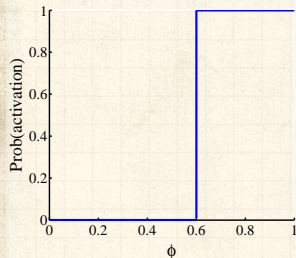
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## Granovetter's Threshold model—recap



Assumes deterministic  
response functions



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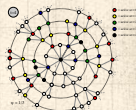
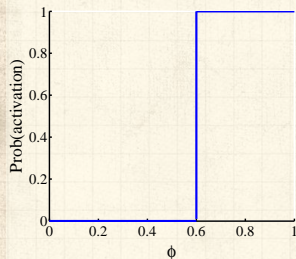
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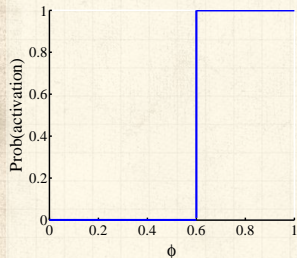
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$\phi_*$  = threshold of an  
individual.



## Granovetter's Threshold model—recap



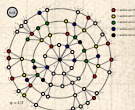
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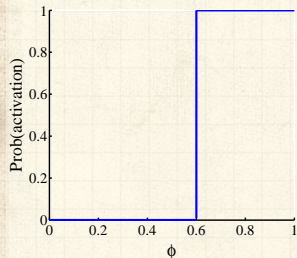
$\phi_*$  = threshold of an individual.





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





## Granovetter's Threshold model—recap

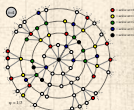


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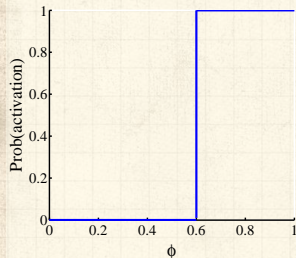
  $\phi_*$  = threshold of an individual.


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


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





## Granovetter's Threshold model—recap




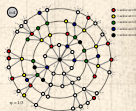
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  $\phi_t$  = fraction of people 'rioting' at time step  $t$ .



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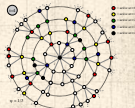
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At time  $t + 1$ , fraction rioting = fraction with

$$\phi_* \leq \phi_t.$$



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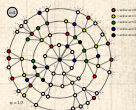


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# Social Sciences—Threshold models

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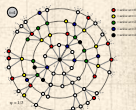
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$\Rightarrow$  Iterative maps of the unit interval  $[0, 1]$ .





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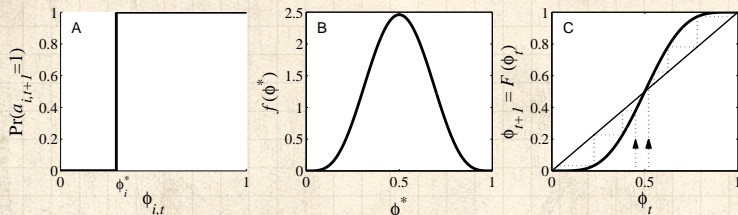
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Action based on perceived behavior of others.



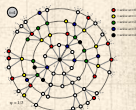
Two states: S and I



Recover now possible (SIS)



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



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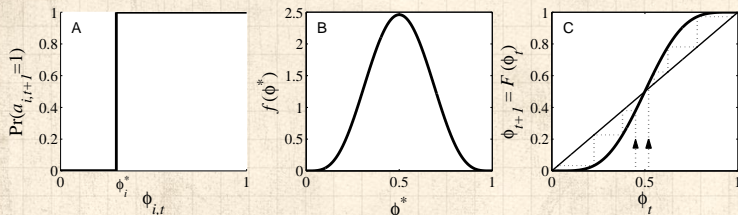
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Two states: S and I



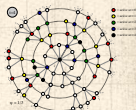
Recover now possible (SIS)



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Discrete time, synchronous update (strong assumption!)



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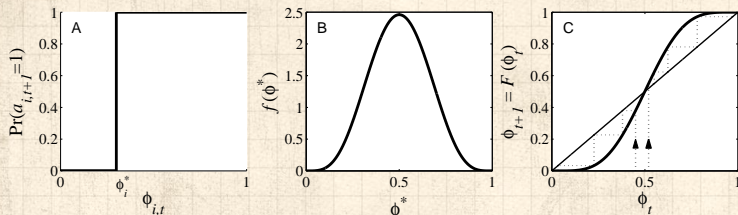
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Two states: S and I



Recover now possible (SIS)



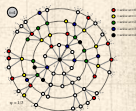
$\phi$  = fraction of contacts 'on' (e.g., rioting)



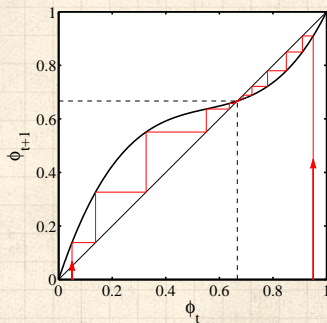
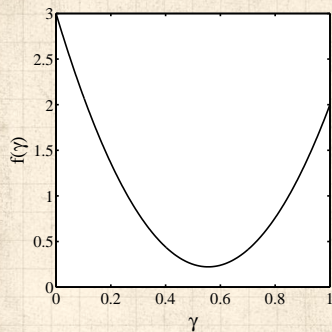
Discrete time, synchronous update (strong assumption!)



This is a **Critical mass model**



# Social Sciences—Threshold models



Example of single stable state model

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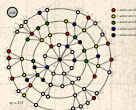
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Implications for collective action theory:

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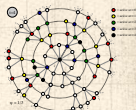
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# Social Sciences—Threshold models

Implications for collective action theory:

1. Collective uniformity  $\nRightarrow$  individual uniformity

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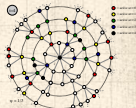
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# Social Sciences—Threshold models

## Implications for collective action theory:

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

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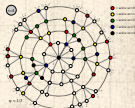
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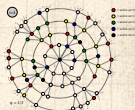
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Implications for collective action theory:

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

Next:





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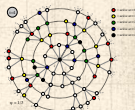
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Connect mean-field model to network model.



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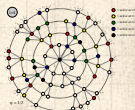
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 Single seed for network model:  $1/N \rightarrow 0$ .



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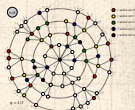
References

## Implications for collective action theory:

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

## Next:

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



# All-to-all versus random networks

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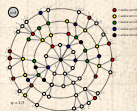
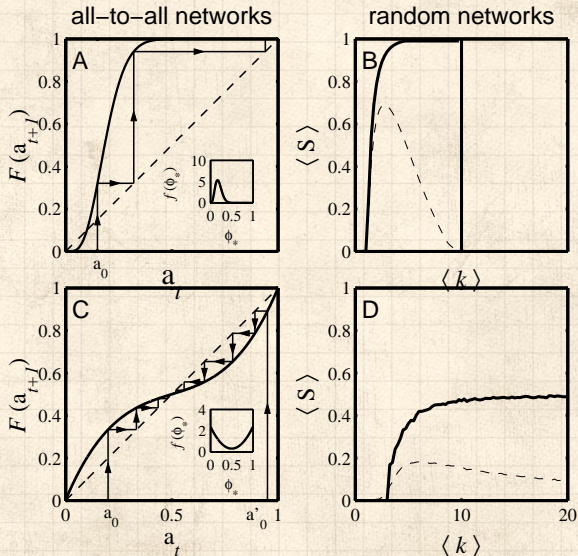
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# Spreadworthiness: Cat videos

## Bowling with Ragdolls:

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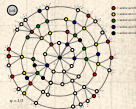
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<https://www.youtube.com/watch?v=XX-g2nmqL9Q?rel=0> 

 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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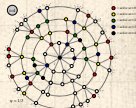
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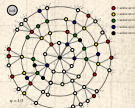
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Three key pieces to describe analytically:

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



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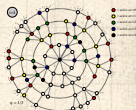
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Three key pieces to describe analytically:

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2. The chance of starting a global spreading event,

$$P_{\text{trig}} = S_{\text{trig}}.$$





# Threshold contagion on random networks

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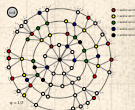
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2. The chance of starting a global spreading event,  $P_{\text{trig}} = S_{\text{trig}}$ .
3. The expected final size of any successful spread,  $S$ .



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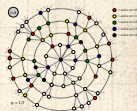
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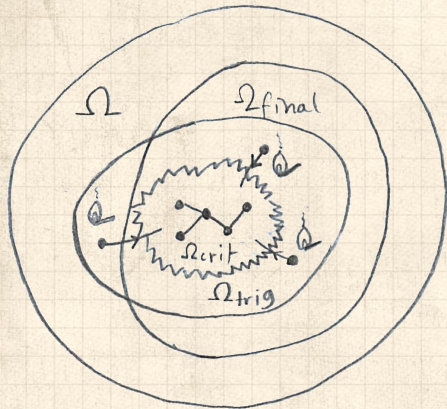
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3. The expected final size of any successful spread,  $S$ .
  - 📦 n.b., the distribution of  $S$  is almost always bimodal.





# Example random network structure:



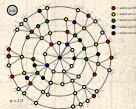
  $\Omega_{crit} = \Omega_{vuln} =$   
critical mass =  
global  
vulnerable  
component

  $\Omega_{trig} =$   
triggering  
component

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

  $\Omega =$  entire  
network

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



# Outline

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## Global spreading condition

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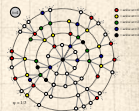
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# Threshold contagion on random networks



**First goal:** Find the largest component of vulnerable nodes.

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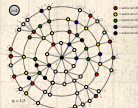
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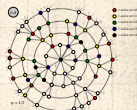


# Threshold contagion on random networks


 **First goal:** Find the largest component of vulnerable nodes.


 Recall that for finding the giant component's size, we had to solve:

$$F_{\pi}(x) = xF_P(F_{\rho}(x)) \quad \text{and} \quad F_{\rho}(x) = xF_R(F_{\rho}(x))$$




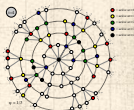
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 We'll find a similar result for the subset of nodes that are vulnerable.





# Threshold contagion on random networks

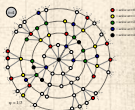
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 This is a node-based percolation problem.







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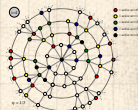
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 We'll find a similar result for the subset of nodes that are vulnerable.


 This is a node-based percolation problem.

 For a general monotonic threshold distribution  $f(\phi)$ , a degree  $k$  node is vulnerable with probability

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



# Threshold contagion on random networks

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

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

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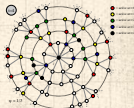
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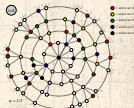
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 The generating function for friends-of-friends distribution is similar to before:


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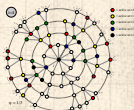
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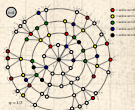
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## Threshold contagion on random networks

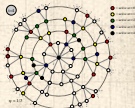
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 Detail: We still have the underlying degree distribution involved in the denominator.



# Threshold contagion on random networks



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

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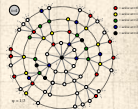
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# Threshold contagion on random networks



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

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

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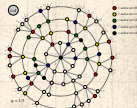
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# Threshold contagion on random networks



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

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

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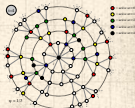
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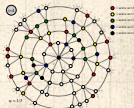
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$$F_{\rho}^{(\text{vuln})}(x) = \underbrace{1 - F_R^{(\text{vuln})}(1)}_{\substack{\text{first node} \\ \text{is not} \\ \text{vulnerable}}} + x F_R^{(\text{vuln})}(F_{\rho}^{(\text{vuln})}(x))$$

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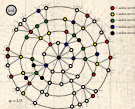
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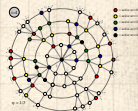
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Functional relations for component size g.f.'s are almost the same ...

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

$$F_{\rho}^{(\text{vuln})}(x) = \underbrace{1 - F_R^{(\text{vuln})}(1)}_{\substack{\text{first node} \\ \text{is not} \\ \text{vulnerable}}} + x F_R^{(\text{vuln})}(F_{\rho}^{(\text{vuln})}(x))$$

Can now solve as before to find

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

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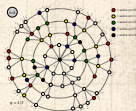
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# Threshold contagion on random networks

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

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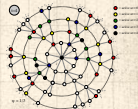
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# Threshold contagion on random networks

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 Assumption is **first node** is **randomly chosen**.

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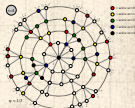
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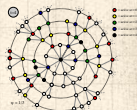
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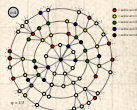
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 Solve as before to find  $P_{\text{trig}} = S_{\text{trig}} = 1 - F_{\pi}^{(\text{trig})}(1)$ .



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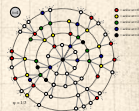
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# Physical derivation of possibility and probability of global spreading:

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

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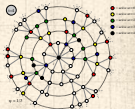
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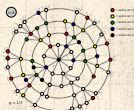
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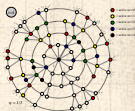
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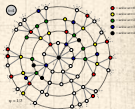
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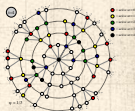
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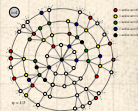
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🧱 Call this  $Q_{\text{trig}}$ .





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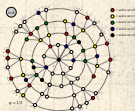
🧱 Next: what's the probability that a randomly infected node will cause a global spreading event?

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🧱 Call this  $Q_{\text{trig}}$ .

🧱 Later: Generalize to more complex networks involving assortativity of all kinds.



Probability an infected edge leads to a global spreading event:

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

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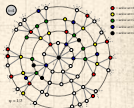
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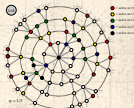
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 Follow an infected edge and use three pieces:



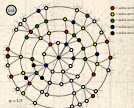
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$$Q_k = \frac{k P_k}{\langle k \rangle}.$$



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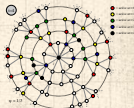
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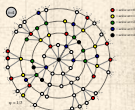
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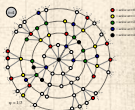
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 Put everything together and solve for  $Q_{\text{trig}}$ :

$$Q_{\text{trig}} = \sum_k \frac{k P_k}{\langle k \rangle} \cdot B_{k1} \cdot [1 - (1 - Q_{\text{trig}})^{k-1}].$$



## Good things about our equation for $Q_{\text{trig}}$ :

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

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

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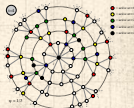
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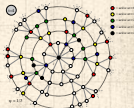
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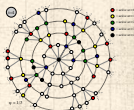
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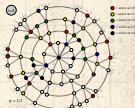
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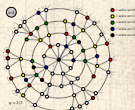
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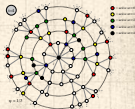
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-  Start with a suitably small seed  $Q_{\text{trig}}^{(1)} > 0$  and iterate while rubbing hands together.





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

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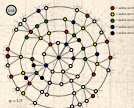
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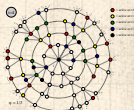


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Interpret  $S_{\text{vuln}}$  as the probability a randomly chosen node is vulnerable and that infecting it leads to a global spreading event:

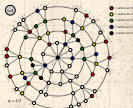
$$S_{\text{vuln}} = \sum_k P_k \cdot B_{k1} \cdot [1 - (1 - Q_{\text{trig}})^k] > 0.$$



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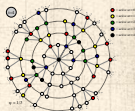


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Probability of global spreading differs only in that we don't care if the initial seed is vulnerable or not:

$$P_{\text{trig}} = S_{\text{trig}} = \sum_k P_k \cdot [1 - (1 - Q_{\text{trig}})^k]$$



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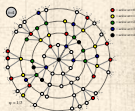
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As for  $S_{\text{vuln}}$ ,  $P_{\text{trig}}$  is non-zero when  $Q_{\text{trig}} > 0$ .



## Connection to generating function results:

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

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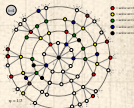
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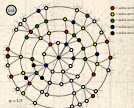
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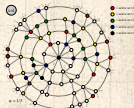
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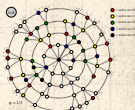
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🧱 Some breathless algebra it all matches:

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



# Fractional size of the largest vulnerable component:



The generating function approach gave

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

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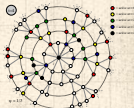
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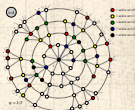
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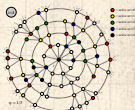
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🧱 Excited scrabbling about gives us, as before:

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# Triggering probability for single-seed global spreading events:



Slight adjustment to the vulnerable component calculation.

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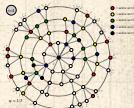
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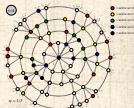
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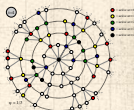
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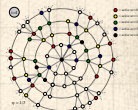
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## Connection to simple gain ratio argument:

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

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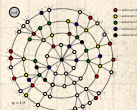
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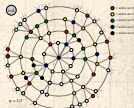
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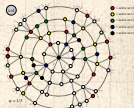
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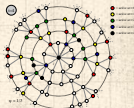
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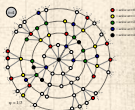
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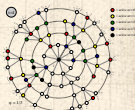
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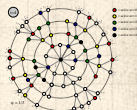
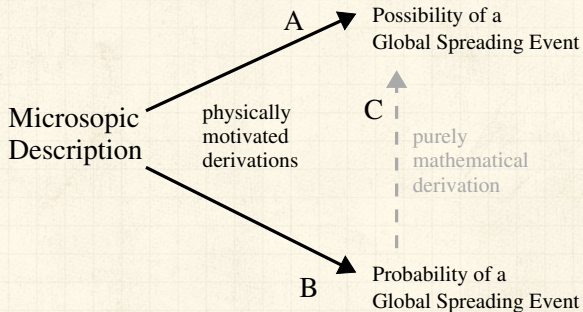
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- When does this equation have a solution  $0 < Q_{\text{trig}} \leq 1$ ?

- We need to find out what happens as  $Q_{\text{trig}} \rightarrow 0$ . [9]



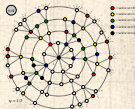
## What we're doing:





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

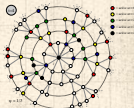
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For  $Q_{\text{trig}} \rightarrow 0^+$ , equation tends towards

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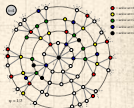




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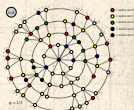


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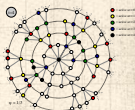
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Only defines the phase transition points (i.e.,  $\mathbf{R} = 1$ ).



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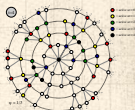
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Inequality?




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

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





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$$Q_{\text{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ \lambda + \left( \lambda + (k-1)Q_{\text{trig}} - \binom{k-1}{2} Q_{\text{trig}}^2 \right) \right]$$

$$\Rightarrow Q_{\text{trig}} = \sum_k \frac{kP_k}{\langle k \rangle} \bullet B_{k1} \bullet \left[ (k-1)Q_{\text{trig}} - \binom{k-1}{2} Q_{\text{trig}}^2 \right]$$

$$\Rightarrow \sum_k \frac{kP_k}{\langle k \rangle} \bullet (k-1) \bullet B_{k1} = 1 + \sum_k \frac{kP_k}{\langle k \rangle} B_{k1} \binom{k-1}{2} Q_{\text{trig}}$$

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

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

 From this connection, we don't know anything about a gain ratio  $\mathbf{R}$  or how to arrange the pieces.

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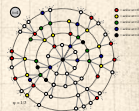
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# Threshold contagion on random networks

 **Third goal:** Find expected fractional size of spread.

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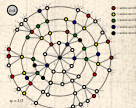
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# Threshold contagion on random networks

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 Not obvious even for uniform threshold problem.

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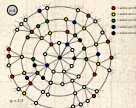
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# Threshold contagion on random networks

- Third goal: Find expected fractional size of spread.
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- Difficulty is in figuring out if and when nodes that need  $\geq 2$  hits switch on.

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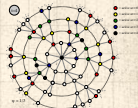
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- Problem solved for infinite seed case by Gleeson and Cahalane:  
"Seed size strongly affects cascades on random networks," Phys. Rev. E, 2007. [7]

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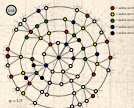
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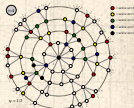
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 Developed further by Gleeson in “Cascades on correlated and modular random networks,” Phys. Rev. E, 2008. [6]








# Expected size of spread

Idea:

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

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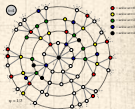
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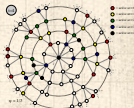
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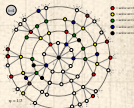
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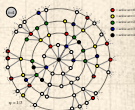
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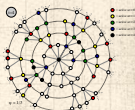
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# Expected size of spread

## Idea:

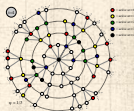
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# Expected size of spread

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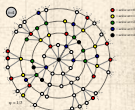
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  - $t = 2$ : enough of  $i$ 's friends and friends-of-friends switched on at time  $t = 0$  so that  $i$ 's threshold is now exceeded.



# Expected size of spread

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  - $t = 2$ : 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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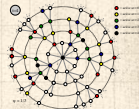
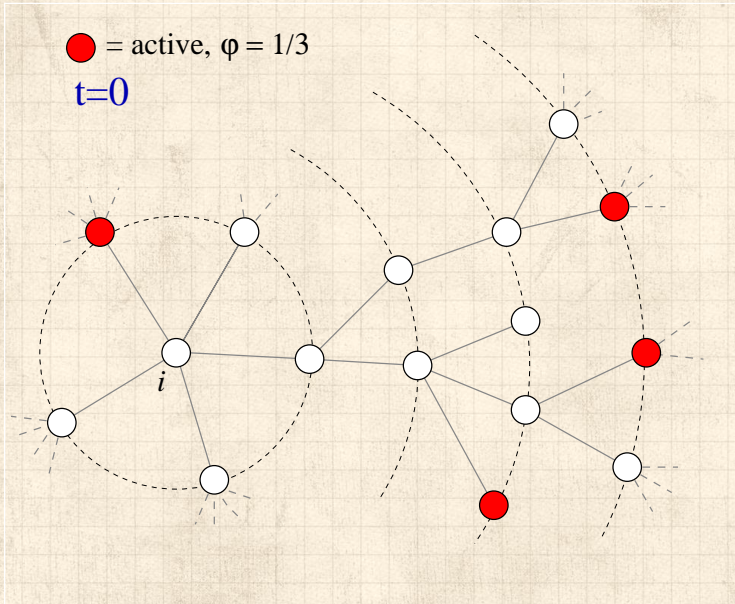
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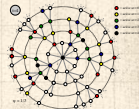
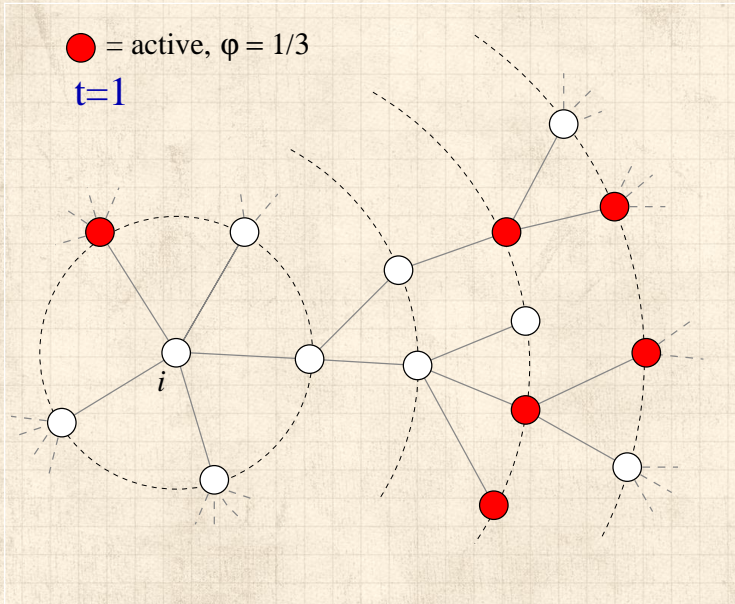
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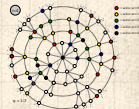
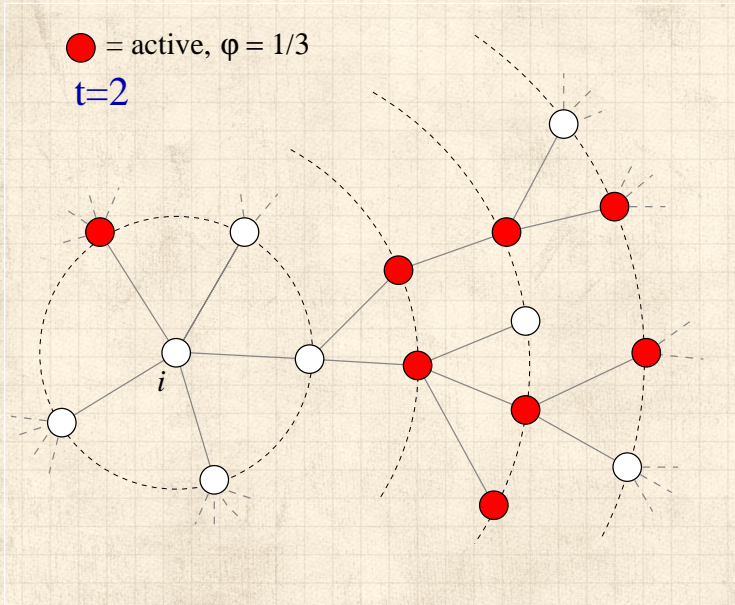
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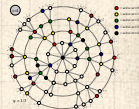
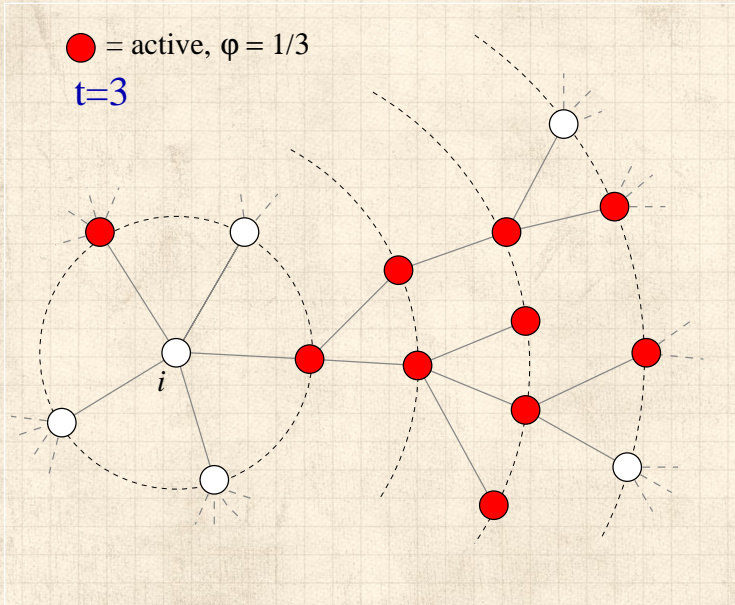
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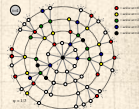
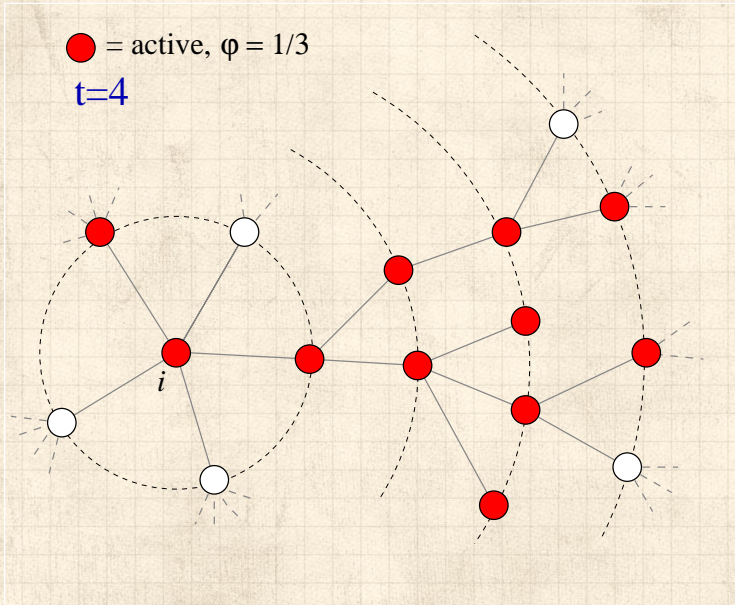
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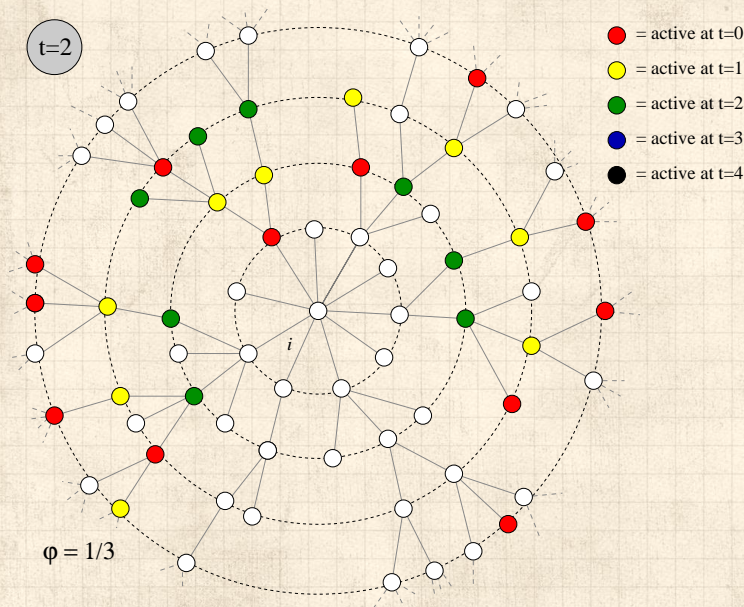
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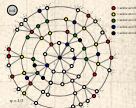
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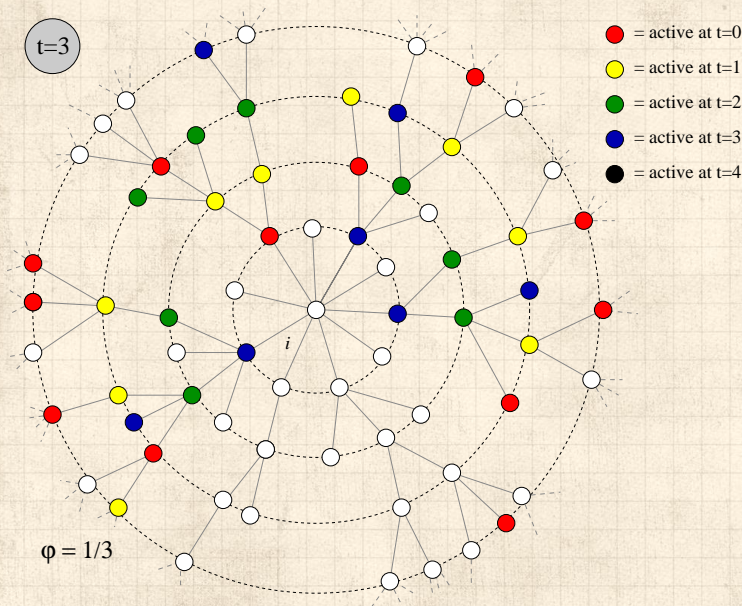
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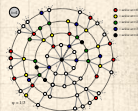
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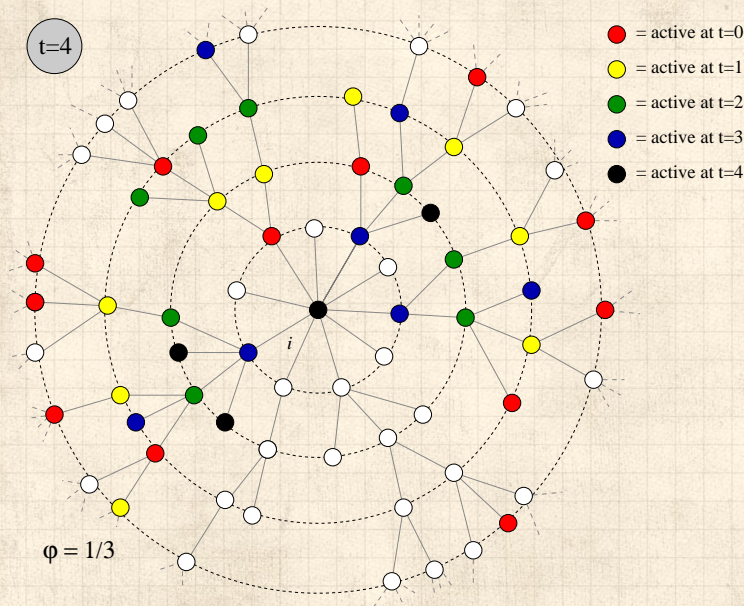
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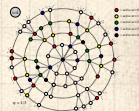
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## Notes:

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

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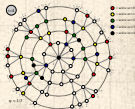
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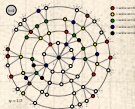
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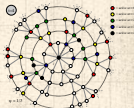
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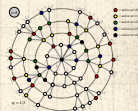
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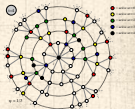
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- We can in fact determine  $\Pr(\text{node of degree } k \text{ switches on at time } t)$ .
- Even more, we can compute:  $\Pr(\text{specific node } i \text{ switches on at time } t)$ .
- Asynchronous updating can be handled too.

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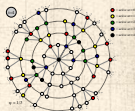
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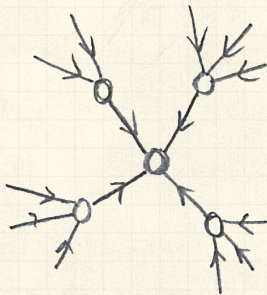
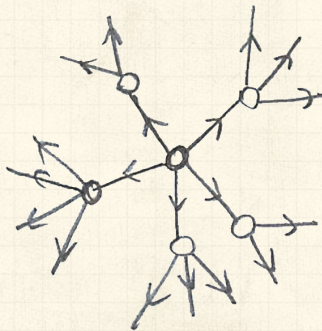
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# Expected size of spread

Pleasantness:

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



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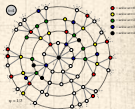
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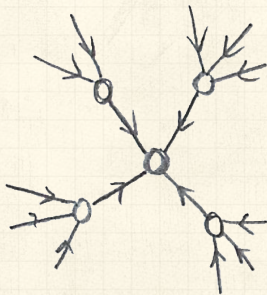
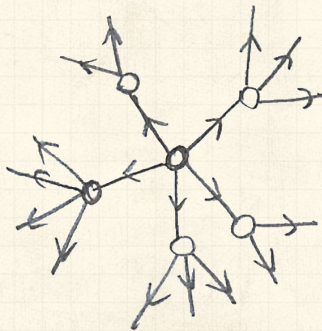




# Expected size of spread

## 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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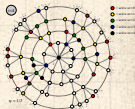
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# Expected size of spread



## Notation:

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

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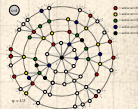
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# Expected size of spread



**Notation:**

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



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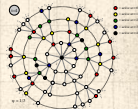
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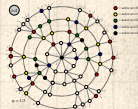
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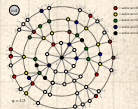
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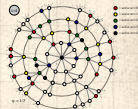
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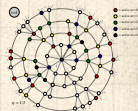
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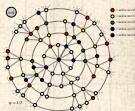


Probability a degree  $k$  node was not a seed at  $t = 0$  is  $(1 - \phi_0)$ .




Combining everything, we have:

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





# Expected size of spread

 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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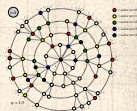
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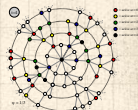
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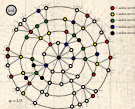


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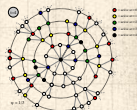
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Story analogous to  $t = 1$  case. For specific node  $i$ :

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



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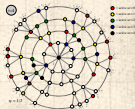
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Average over all nodes with degree  $k$  to obtain expression for  $\phi_{t+1}$ :

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



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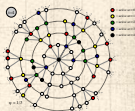
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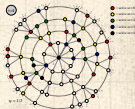
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So we need to compute  $\theta_t$ ... massive excitement...





# Expected size of spread


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

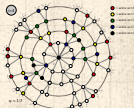
  $\theta_1 = \phi_0 +$

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

  $\frac{k P_k}{\langle k \rangle} = Q_k = \mathbf{Pr}$  (edge connects to a degree  $k$  node).

  $\sum_{j=0}^{k-1}$  piece gives  $\mathbf{Pr}$  (degree node  $k$  activates if  $j$  of its  $k - 1$  incoming neighbors are active).


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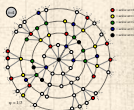
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 See this all generalizes to give  $\theta_{t+1}$  in terms of  $\theta_t \dots$



# Expected size of spread

Two pieces: edges first, and then nodes

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

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

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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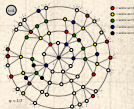
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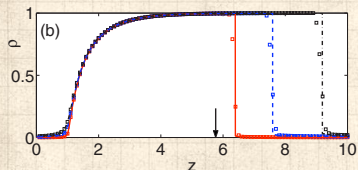
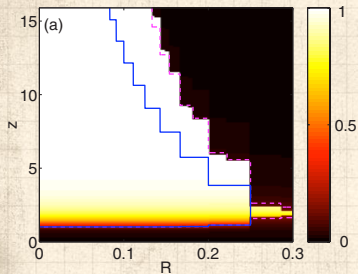
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# Comparison between theory and simulations



Pure random networks with simple threshold responses



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



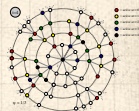
$\phi_0 = 10^{-3}$ ,  $0.5 \times 10^{-2}$ , and  $10^{-2}$ .



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



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



From Gleeson and Cahalane [7]

# Notes:



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

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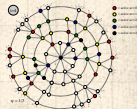
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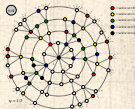
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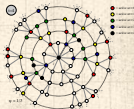


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- Depends on map  $\theta_{t+1} = G(\theta_t; \phi_0)$ .
- First: if self-starters are present, some activation is assured:

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

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



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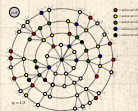
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meaning  $B_{k0} > 0$  for at least one value of  $k \geq 1$ .


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

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



# Notes:

## In words:

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

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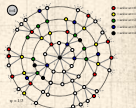
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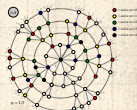
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- 🧱 Cascade condition is more complicated for  $\phi_0 > 0$ .

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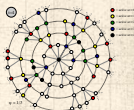
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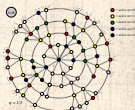
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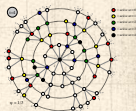
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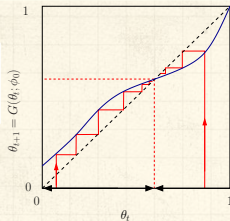
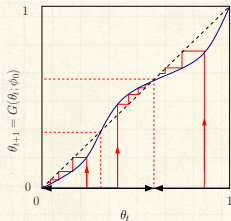
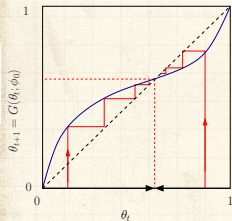
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- 🧱 Tricky point:  $G$  depends on  $\phi_0$ , so as we change  $\phi_0$ , we also change  $G$ .



# General fixed point story:



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

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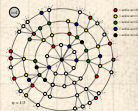
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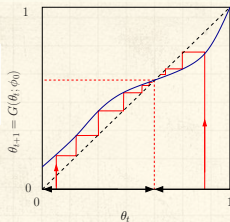
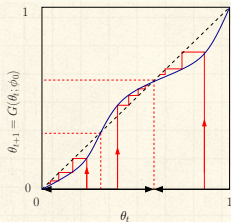
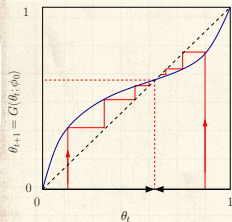
Final size

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:** Actual form of  $G$  depends on  $\phi_0$ .

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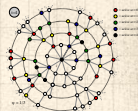
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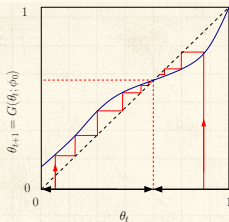
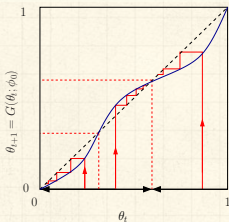
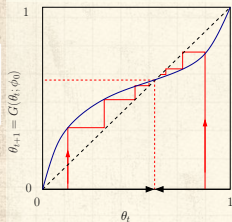
Spreading possibility  
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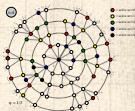


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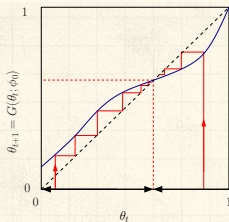
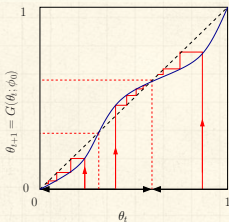
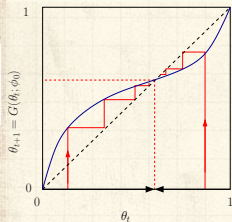
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 **Important:**  $\phi_t$  can only increase monotonically so  $\phi_0$  must shape  $G$  so that  $\phi_0$  is at or above an unstable fixed point.





# General fixed point story:



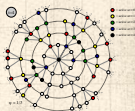
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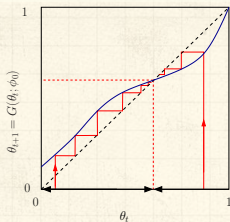
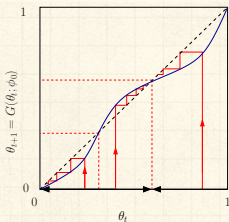
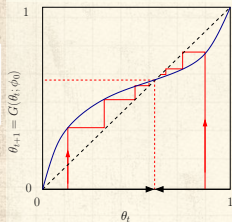
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First reason:  $\phi_1 \geq \phi_0$ .





# General fixed point story:





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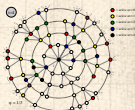
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 **Important:** Actual form of  $G$  depends on  $\phi_0$ .

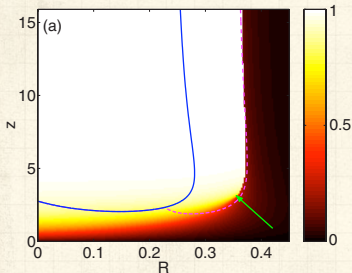
 **Important:**  $\phi_t$  can only increase monotonically so  $\phi_0$  must shape  $G$  so that  $\phi_0$  is at or above an unstable fixed point.

 First reason:  $\phi_1 \geq \phi_0$ .

 Second:  $G'(\theta; \phi_0) \geq 0, 0 \leq \theta \leq 1$ .



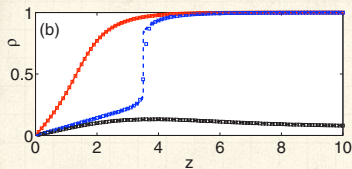
## Interesting behavior:



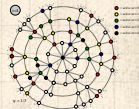
Now allow thresholds  
to be distributed  
according to a  
Gaussian with mean  $R$ .



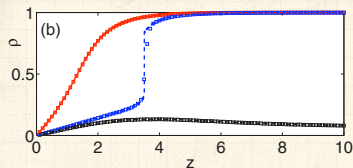
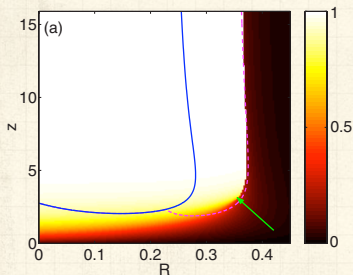
$R = 0.2, 0.362,$  and  
 $0.38; \sigma = 0.2.$



From Gleeson and  
Cahalane [7]



## Interesting behavior:



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

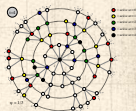


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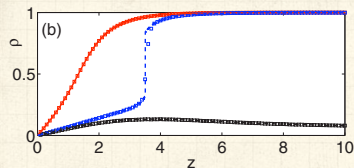
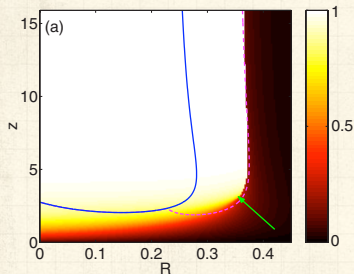


$\phi_0 = 0$  but some nodes have thresholds  $\leq 0$  so effectively  $\phi_0 > 0.$

From Gleeson and Cahalane [7]



## Interesting behavior:



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



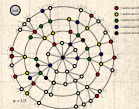
$R = 0.2, 0.362,$  and  $0.38; \sigma = 0.2.$



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

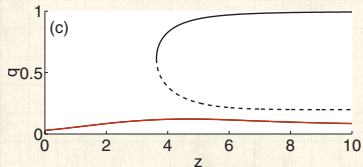
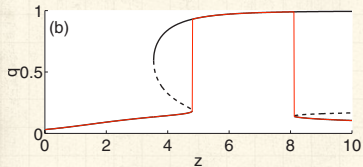
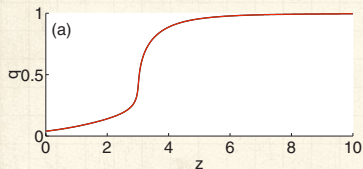


Now see a (nasty) discontinuous phase transition for low  $\langle k \rangle.$



From Gleeson and Cahalane [7]

## Interesting behavior:



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



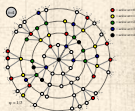
n.b.: 0 is not a fixed point here:  $\theta_0 = 0$  always takes off.



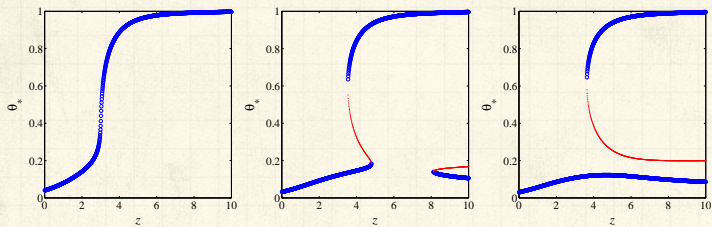
Top to bottom:  $R = 0.35, 0.371,$  and  $0.375$ .



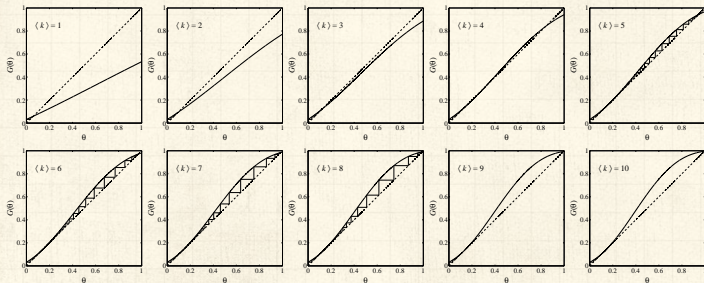
Saddle node bifurcations appear and merge (b and c).



# What's happening:



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



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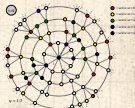
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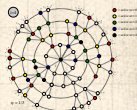
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# Time-dependent solutions

## Synchronous update

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

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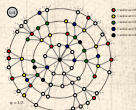
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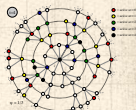
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 Update nodes with probability  $\alpha$ .



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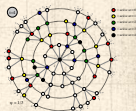
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 Update nodes with probability  $\alpha$ .

 As  $\alpha \rightarrow 0$ , updates become effectively independent.



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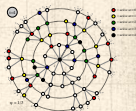
Done: Evolution of  $\phi_t$  and  $\theta_t$  given exactly by the maps we have derived.

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Update nodes with probability  $\alpha$ .

As  $\alpha \rightarrow 0$ , updates become effectively independent.

Now can talk about  $\phi(t)$  and  $\theta(t)$ .



# Nutshell:

 Solid dive into understanding contagion on generalized random networks.

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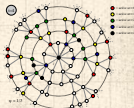
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# Nutshell:

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- ❏ Generating function approaches provided first breakthroughs and gave possibility and probability of spreading. [10, 16]

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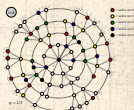
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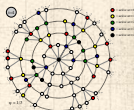
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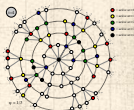
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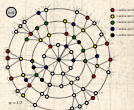
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- 🧱 Many connections to other kinds of models: Voter models, Ising models, ...

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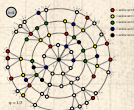
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# Neural reboot (NR):

Pangolin happiness:

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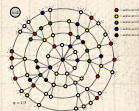
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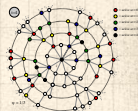


<https://www.youtube.com/watch?v=LMIYjkG4onM?rel=0>



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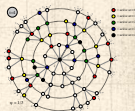
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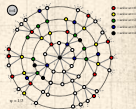
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