

# Assortativity and Mixing

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

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Assortativity by  
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Contagion

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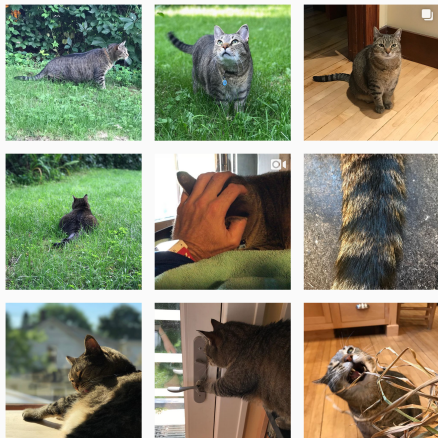
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 On Instagram at [pratchett\\_the\\_cat](https://www.instagram.com/pratchett_the_cat) 

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## Basic idea:



Random networks with arbitrary degree distributions cover much territory but do not represent all networks.

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  2. demographics (age, gender, etc.)
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- ❏ We speak of mixing patterns, correlations, biases...
- ❏ Networks are still random at base but now have more global structure.
- ❏ Build on work by Newman <sup>[5, 6]</sup>, and Boguñá and Serano. <sup>[1]</sup>.

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# General mixing between node categories



Assume types of nodes are countable, and are assigned numbers 1, 2, 3, ....

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 Requirements:

$$\sum_{\nu} e_{\mu\nu} = 1, \quad \sum_{\nu} e_{\mu\nu} = a_{\mu}, \quad \text{and} \quad \sum_{\mu} e_{\mu\nu} = b_{\nu}.$$

# Notes:



Varying  $e_{\mu\nu}$  allows us to move between the following:

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Varying  $e_{\mu\nu}$  allows us to move between the following:

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Basic story: level of assortativity reflects the degree to which nodes are connected to nodes within their group.


# Correlation coefficient:

 Quantify the level of assortativity with the following **assortativity coefficient** <sup>[6]</sup>:

$$r = \frac{\sum_{\mu} e_{\mu\mu} - \sum_{\mu} a_{\mu} b_{\mu}}{1 - \sum_{\mu} a_{\mu} b_{\mu}} = \frac{\text{Tr} \mathbf{E} - \|E^2\|_1}{1 - \|E^2\|_1}$$


where  $\|\cdot\|_1$  is the 1-norm = sum of a matrix's entries.

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- When  $e_{\mu\mu} = a_{\mu} b_{\mu}$ , we have  $r = 0$ . ✓



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$$r_{\min} = \frac{-\|E^2\|_1}{1 - \|E^2\|_1}$$

where  $-1 \leq r_{\min} < 0$ .

# Watch your step

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NuhnuhNuhnuhNuhnuhNuhnuhNuhnuhNuhnuh

...



# Scalar quantities



Now consider nodes defined by a scalar integer quantity.

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- This is the observed normalized deviation from randomness in the product  $jk$ .

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 **Important:** Must separately define  $P_0$  as the  $\{e_{jk}\}$  contain no information about isolated nodes.


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 Directed networks still fine but we will assume from here on that  $e_{jk} = e_{kj}$ .

# Degree-degree correlations

 Notation reconciliation for undirected networks:

$$r = \frac{\sum_{j,k} jk(e_{jk} - R_j R_k)}{\sigma_R^2}$$


where, as before,  $R_k$  is the probability that a randomly chosen edge leads to a node of degree  $k + 1$ , and

$$\sigma_R^2 = \sum_j j^2 R_j - \left[ \sum_j j R_j \right]^2.$$



# Degree-degree correlations

Error estimate for  $r$ :

 Remove edge  $i$  and recompute  $r$  to obtain  $r_i$ .

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## Error estimate for $r$ :

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-  Repeat for all edges and compute using the jackknife method  <sup>[3]</sup>


$$\sigma_r^2 = \sum_i (r_i - r)^2.$$

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$$\sigma_r^2 = \sum_i (r_i - r)^2.$$

-  Mildly sneaky as variables need to be independent for us to be truly happy and edges are correlated...

# Measurements of degree-degree correlations

|               | Group | Network                   | Type       | Size $n$  | Assortativity $r$ | Error $\sigma_r$ |
|---------------|-------|---------------------------|------------|-----------|-------------------|------------------|
| Social        | a     | Physics coauthorship      | undirected | 52 909    | 0.363             | 0.002            |
|               | a     | Biology coauthorship      | undirected | 1 520 251 | 0.127             | 0.0004           |
|               | b     | Mathematics coauthorship  | undirected | 253 339   | 0.120             | 0.002            |
|               | c     | Film actor collaborations | undirected | 449 913   | 0.208             | 0.0002           |
|               | d     | Company directors         | undirected | 7 673     | 0.276             | 0.004            |
|               | e     | Student relationships     | undirected | 573       | -0.029            | 0.037            |
|               | f     | Email address books       | directed   | 16 881    | 0.092             | 0.004            |
| Technological | g     | Power grid                | undirected | 4 941     | -0.003            | 0.013            |
|               | h     | Internet                  | undirected | 10 697    | -0.189            | 0.002            |
|               | i     | World Wide Web            | directed   | 269 504   | -0.067            | 0.0002           |
|               | j     | Software dependencies     | directed   | 3 162     | -0.016            | 0.020            |
| Biological    | k     | Protein interactions      | undirected | 2 115     | -0.156            | 0.010            |
|               | l     | Metabolic network         | undirected | 765       | -0.240            | 0.007            |
|               | m     | Neural network            | directed   | 307       | -0.226            | 0.016            |
|               | n     | Marine food web           | directed   | 134       | -0.263            | 0.037            |
|               | o     | Freshwater food web       | directed   | 92        | -0.326            | 0.031            |

 Social networks tend to be assortative (homophily)

 Technological and biological networks tend to be disassortative

# Hot lava

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"I like it" 

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# Spreading on degree-correlated networks



Next: Generalize our work for random networks to degree-correlated networks.

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# Spreading on degree-correlated networks

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1. find the giant component size.



# Spreading on degree-correlated networks

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 Next: Generalize our work for random networks to degree-correlated networks.

 As before, by allowing that a node of degree  $k$  is activated by one neighbor with probability  $B_{k1}$ , we can handle various problems:

1. find the giant component size.
2. find the probability and extent of spread for simple disease models.

# Spreading on degree-correlated networks

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 Next: Generalize our work for random networks to degree-correlated networks.

 As before, by allowing that a node of degree  $k$  is activated by one neighbor with probability  $B_{k1}$ , we can handle various problems:

1. find the giant component size.
2. find the probability and extent of spread for simple disease models.
3. find the probability of spreading for simple threshold models.

# Spreading on degree-correlated networks



**Goal:** Find  $f_{n,j} = \Pr$  an edge emanating from a degree  $j + 1$  node leads to a finite active subcomponent of size  $n$ .

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 Repeat: a node of degree  $k$  is in the game with probability  $B_{k1}$ .

 Define  $\vec{B}_1 = [B_{k1}]$ .

 **Plan:** Find the generating function

$$F_j(x; \vec{B}_1) = \sum_{n=0}^{\infty} f_{n,j} x^n.$$

# Spreading on degree-correlated networks



Recursive relationship:

$$F_j(x; \vec{B}_1) = x^0 \sum_{k=0}^{\infty} \frac{e_{jk}}{R_j} (1 - B_{k+1,1}) \\ + x \sum_{k=0}^{\infty} \frac{e_{jk}}{R_j} B_{k+1,1} [F_k(x; \vec{B}_1)]^k .$$

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 **First term** = **Pr** (that the first node we reach is not in the game).

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 **First term** = **Pr** (that the first node we reach is not in the game).

 **Second term** involves **Pr** (we hit an active node which has  $k$  outgoing edges).

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 **First term** = **Pr** (that the first node we reach is not in the game).

 **Second term** involves **Pr** (we hit an active node which has  $k$  outgoing edges).

 Next: find average size of active components reached by following a link from a degree  $j + 1$  node =  $F'_j(1; \vec{B}_1)$ .



# Spreading on degree-correlated networks

 Differentiate  $F_j(x; \vec{B}_1)$ , set  $x = 1$ , and rearrange.

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# Spreading on degree-correlated networks

 Differentiate  $F_j(x; \vec{B}_1)$ , set  $x = 1$ , and rearrange.

 We use  $F_k(1; \vec{B}_1) = 1$  which is true when no giant component exists.

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# Spreading on degree-correlated networks


 Differentiate  $F_j(x; \vec{B}_1)$ , set  $x = 1$ , and rearrange.

 We use  $F_k(1; \vec{B}_1) = 1$  which is true when no giant component exists. We find:


$$R_j F'_j(1; \vec{B}_1) = \sum_{k=0}^{\infty} e_{jk} B_{k+1,1} + \sum_{k=0}^{\infty} k e_{jk} B_{k+1,1} F'_k(1; \vec{B}_1).$$

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 Rearranging and introducing a sneaky  $\delta_{jk}$ :

$$\sum_{k=0}^{\infty} (\delta_{jk} R_k - k B_{k+1,1} e_{jk}) F'_k(1; \vec{B}_1) = \sum_{k=0}^{\infty} e_{jk} B_{k+1,1}.$$

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 In matrix form, we have


$$\mathbf{A}_{\mathbf{E}, \vec{B}_1} \vec{F}'(1; \vec{B}_1) = \mathbf{E} \vec{B}_1$$

where

$$\begin{aligned} [\mathbf{A}_{\mathbf{E}, \vec{B}_1}]_{j+1, k+1} &= \delta_{jk} R_k - k B_{k+1, 1} e_{jk}, \\ [\vec{F}'(1; \vec{B}_1)]_{k+1} &= F'_k(1; \vec{B}_1), \\ [\mathbf{E}]_{j+1, k+1} &= e_{jk}, \text{ and } [\vec{B}_1]_{k+1} = B_{k+1, 1}. \end{aligned}$$



# Spreading on degree-correlated networks

 So, in principle at least:

$$\vec{F}'(1; \vec{B}_1) = \mathbf{A}_{\mathbf{E}, \vec{B}_1}^{-1} \mathbf{E} \vec{B}_1.$$

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 Now: as  $\vec{F}'(1; \vec{B}_1)$ , the average size of an active component reached along an edge, increases, we move towards a transition to a giant component.

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 Right at the transition, the average component size explodes.

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 Exploding inverses of matrices occur when their determinants are 0.

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 Right at the transition, the average component size explodes.

 Exploding inverses of matrices occur when their determinants are 0.

 The condition is therefore:

$$\det \mathbf{A}_{\mathbf{E}, \vec{B}_1} = 0$$

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# Spreading on degree-correlated networks



General condition details:

$$\det \mathbf{A}_{\mathbf{E}, \vec{B}_1} = \det [\delta_{jk} R_{k-1} - (k-1) B_{k,1} e_{j-1, k-1}] = 0.$$

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 The above collapses to our standard contagion condition when  $e_{jk} = R_j R_k$  (see next slide). [2]

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 Bonusville: We'll find a much better version of this set of conditions later...



# Retrieving the cascade condition for uncorrelated networks

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# Spreading on degree-correlated networks

We'll next find two more pieces:

1.  $P_{\text{trig}}$ , the probability of starting a cascade

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# Spreading on degree-correlated networks

We'll next find two more pieces:

1.  $P_{\text{trig}}$ , the probability of starting a cascade
2.  $S$ , the expected extent of activation given a small seed.

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# Spreading on degree-correlated networks

We'll next find two more pieces:

1.  $P_{\text{trig}}$ , the probability of starting a cascade
2.  $S$ , the expected extent of activation given a small seed.

Triggering probability:

 Generating function:

$$H(x; \vec{B}_1) = x \sum_{k=0}^{\infty} P_k \left[ F_{k-1}(x; \vec{B}_1) \right]^k .$$

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 Generating function:

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 Generating function for vulnerable component size is more complicated.

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# Spreading on degree-correlated networks



Want probability of **not reaching** a finite component.

$$\begin{aligned} P_{\text{trig}} = S_{\text{trig}} &= 1 - H(1; \vec{B}_1) \\ &= 1 - \sum_{k=0}^{\infty} P_k \left[ F_{k-1}(1; \vec{B}_1) \right]^k. \end{aligned}$$

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Last piece: we have to compute  $F_{k-1}(1; \vec{B}_1)$ .

# Spreading on degree-correlated networks

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-  Last piece: we have to compute  $F_{k-1}(1; \vec{B}_1)$ .

-  Nastier (nonlinear)—we have to solve the recursive expression we started with when  $x = 1$ :

$$F_j(1; \vec{B}_1) = \sum_{k=0}^{\infty} \frac{e_{jk}}{R_j} (1 - B_{k+1,1}) + \sum_{k=0}^{\infty} \frac{e_{jk}}{R_j} B_{k+1,1} \left[ F_k(1; \vec{B}_1) \right]^k.$$

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- Iterative methods should work here.



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# Spreading on degree-correlated networks

 **Truly final piece:** Find final size using approach of Gleeson <sup>[4]</sup>, a generalization of that used for uncorrelated random networks.

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# Spreading on degree-correlated networks

-  **Truly final piece:** Find final size using approach of Gleeson <sup>[4]</sup>, a generalization of that used for uncorrelated random networks.
-  Need to compute  $\theta_{j,t}$ , the probability that an edge leading to a degree  $j$  node is infected at time  $t$ .

# Spreading on degree-correlated networks

-  **Truly final piece:** Find final size using approach of Gleeson<sup>[4]</sup>, a generalization of that used for uncorrelated random networks.
-  Need to compute  $\theta_{j,t}$ , the probability that an edge leading to a degree  $j$  node is infected at time  $t$ .
-  Evolution of edge activity probability:

$$\theta_{j,t+1} = G_j(\vec{\theta}_t) = \phi_0 + (1 - \phi_0) \times$$

$$\sum_{k=1}^{\infty} \frac{e_{j-1,k-1}}{R_{j-1}} \sum_{i=0}^{k-1} \binom{k-1}{i} \theta_{k,t}^i (1 - \theta_{k,t})^{k-1-i} B_{ki}.$$

# Spreading on degree-correlated networks

- Truly final piece: Find final size using approach of Gleeson<sup>[4]</sup>, a generalization of that used for uncorrelated random networks.
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- Overall active fraction's evolution:

$$\phi_{t+1} = \phi_0 + (1 - \phi_0) \sum_{k=0}^{\infty} P_k \sum_{i=0}^k \binom{k}{i} \theta_{k,t}^i (1 - \theta_{k,t})^{k-i} B_{ki}.$$



# Spreading on degree-correlated networks



As before, these equations give the actual evolution of  $\phi_t$  for synchronous updates.

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As before, these equations give the actual evolution of  $\phi_t$  for synchronous updates.



Contagion condition follows from  $\vec{\theta}_{t+1} = \vec{G}(\vec{\theta}_t)$ .

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Expand  $\vec{G}$  around  $\vec{\theta}_0 = \vec{0}$ .

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Expand  $\vec{G}$  around  $\vec{\theta}_0 = \vec{0}$ .

$$\theta_{j,t+1} = G_j(\vec{0}) + \sum_{k=1}^{\infty} \frac{\partial G_j(\vec{0})}{\partial \theta_{k,t}} \theta_{k,t} + \frac{1}{2!} \sum_{k=1}^{\infty} \frac{\partial^2 G_j(\vec{0})}{\partial \theta_{k,t}^2} \theta_{k,t}^2 + \dots$$

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If  $G_j(\vec{0}) \neq 0$  for at least one  $j$ , always have some infection.



If  $G_j(\vec{0}) = 0 \forall j$ , want largest eigenvalue

$$\left[ \frac{\partial G_j(\vec{0})}{\partial \theta_{k,t}} \right] > 1.$$



# Spreading on degree-correlated networks



As before, these equations give the actual evolution of  $\phi_t$  for synchronous updates.



Contagion condition follows from  $\vec{\theta}_{t+1} = \vec{G}(\vec{\theta}_t)$ .



Expand  $\vec{G}$  around  $\vec{\theta}_0 = \vec{0}$ .

$$\theta_{j,t+1} = G_j(\vec{0}) + \sum_{k=1}^{\infty} \frac{\partial G_j(\vec{0})}{\partial \theta_{k,t}} \theta_{k,t} + \frac{1}{2!} \sum_{k=1}^{\infty} \frac{\partial^2 G_j(\vec{0})}{\partial \theta_{k,t}^2} \theta_{k,t}^2 + \dots$$



If  $G_j(\vec{0}) \neq 0$  for at least one  $j$ , always have some infection.



If  $G_j(\vec{0}) = 0 \forall j$ , want largest eigenvalue

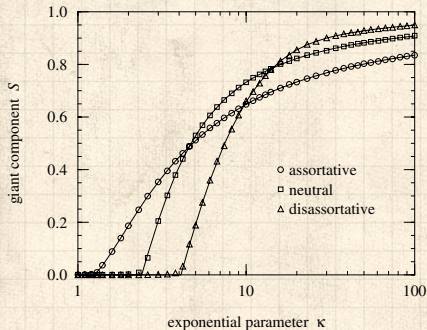
$$\left[ \frac{\partial G_j(\vec{0})}{\partial \theta_{k,t}} \right] > 1.$$



Condition for spreading is therefore dependent on eigenvalues of this matrix:

$$\frac{\partial G_j(\vec{0})}{\partial \theta_{k,t}} = \frac{e_{j-1,k-1}}{R_{j-1}} (k-1) B_{k1}$$

# How the giant component changes with assortativity:



from Newman, 2002 [5]



More assortative networks percolate for lower average degrees



But disassortative networks end up with higher extents of spreading.

# Toy guns don't pretend blow up things ...

# Splsshht

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