

Three Lectures on Networks

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lecture 3: null models and statistical inference for network structure

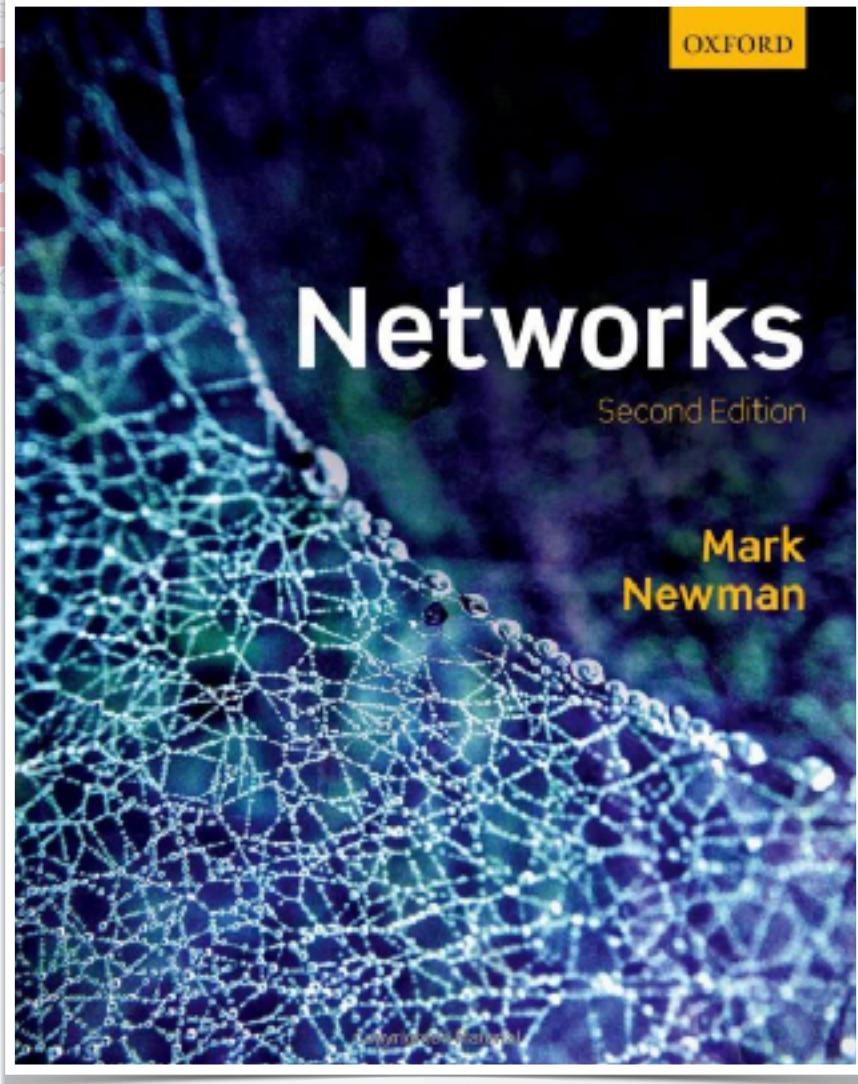


Mark Newman

Professor of Physics
University of Michigan

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<http://www-personal.umich.edu/~mejn/>





University of Colorado **Boulder**

Network Analysis and Modeling

Instructor: Aaron Clauset or Daniel B. Larremore

This graduate-level course will examine modern techniques for analyzing and modeling the structure and dynamics of complex networks. The focus will be on statistical algorithms and methods, and both lectures and assignments will emphasize model interpretability and understanding the processes that generate real data. Applications will be drawn from computational biology and computational social science. No biological or social science training is required. (Note: this is not a scientific computing course, but there will be plenty of computing for science.)

Full lectures notes online (~150 pages in PDF)

<https://aaronclauset.github.io/courses/5352/>



University of Colorado **Boulder**

Biological Networks

Instructor: Aaron Clauset

This undergraduate-level course examines the computational representation and analysis of biological phenomena through the structure and dynamics of networks, from molecules to species. Attention focuses on algorithms for clustering network structures, predicting missing information, modeling flows, regulation, and spreading-process dynamics, examining the evolution of network structure, and developing intuition for how network structure and dynamics relate to biological phenomena.

Full lectures notes online (~150 pages in PDF)

<https://aaronclauset.github.io/courses/3352/>

Software

R

Python

Matlab

★ NetworkX [python]

★ igraph [python, R, c++]

graph-tool [python, c++]

GraphLab [python, c++]

Standalone editors

UCI-Net

NodeXL

Gephi

Pajek

Network Workbench

Cytoscape

yEd graph editor

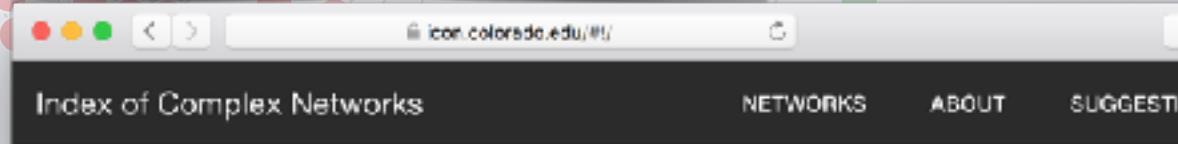
Graphviz

Network data sets



Colorado Index of Complex Networks

icon.colorado.edu



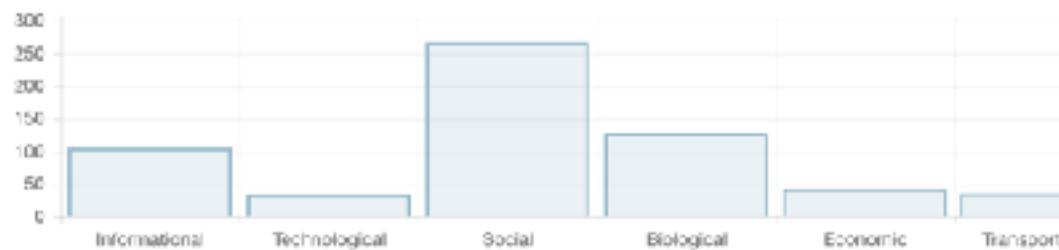
The Colorado Index of Complex Networks (ICON)

ICON is a comprehensive index of research-quality network data sets from all domains of networks, including social, web, information, biological, ecological, connectome, transportation, and technological networks.

Each network record in the index is annotated with and searchable or browsable by its graph properties, description, size, etc., and many records include links to multiple networks. The contents of ICON are curated by volunteer experts from Prof. Aaron Clauset's research group at the University of Colorado Boulder.

Click on the NETWORKS tab above to get started.

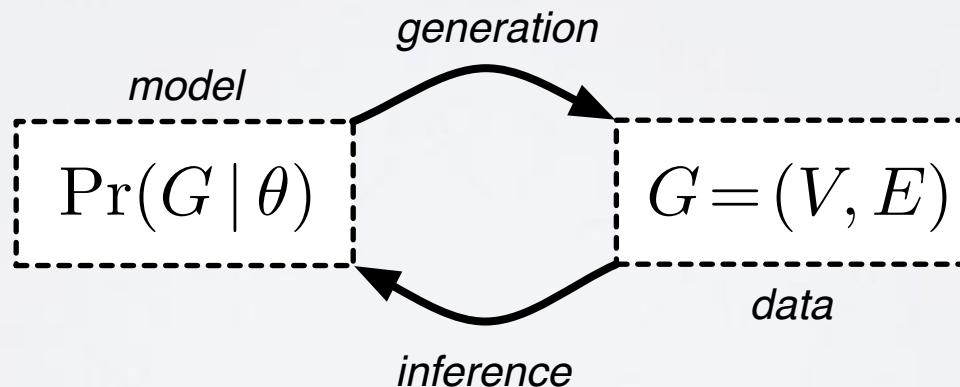
Entries found: 609 Networks found: 4419



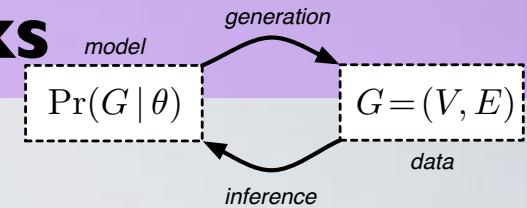
1. defining a network
2. describing a network
- 3. null models and statistical inference for networks**

generative models for complex networks

- define a parametric probability distribution over networks $\Pr(G | \theta)$
- **generation** : given θ , draw G from this distribution
- **inference** : given G , choose θ that makes G likely



generative models for complex networks



general form

$$\Pr(G \mid \theta) = \prod_{ij} \Pr(A_{ij} \mid \theta)$$

edge generation function

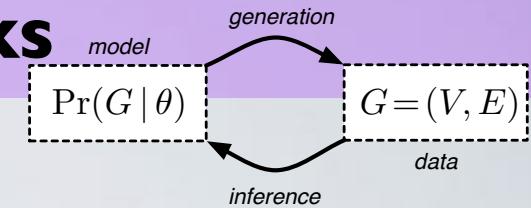
assumptions about “structure” go into $\Pr(A_{ij} \mid \theta)$

$$\text{consistency } \lim_{n \rightarrow \infty} \Pr(\hat{\theta} \neq \theta) = 0$$

requires that edges be conditionally independent*

3 main classes of these models

generative models for complex networks



I. Random graph models (unstructured)

- **edge density** (Erdős-Rényi)

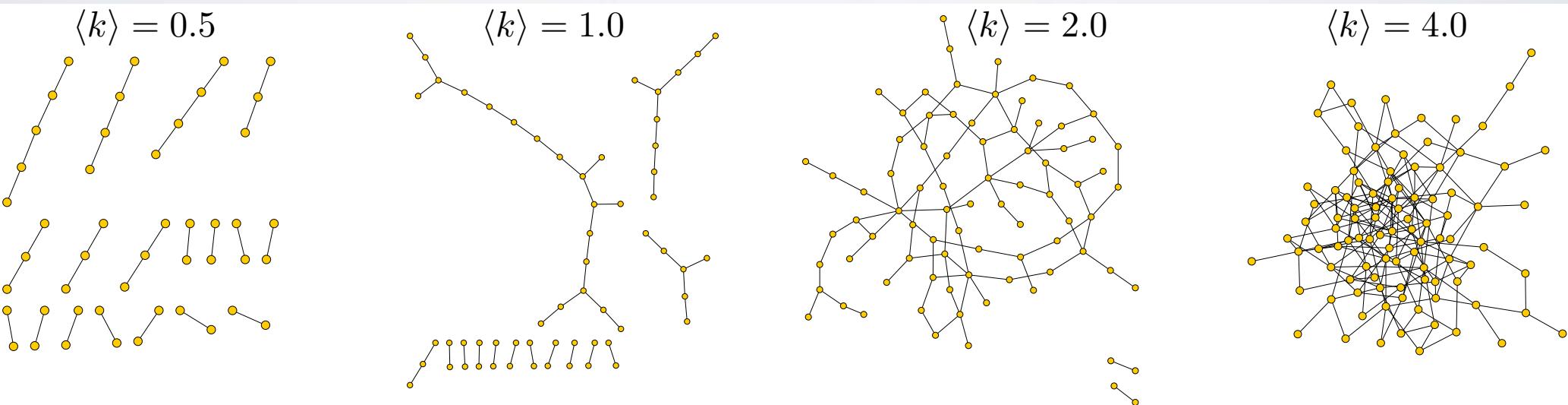
edges are iid $\Pr(A_{ij}) = p$

"homogeneous" random graphs

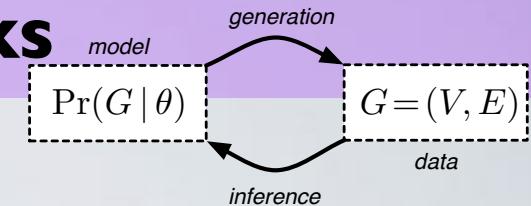
- **degree-based** (Chung-Lu & configuration)

edges independent, conditioned on degree $\Pr(A_{ij}) \propto k_i k_j$

"heterogeneous" random graphs

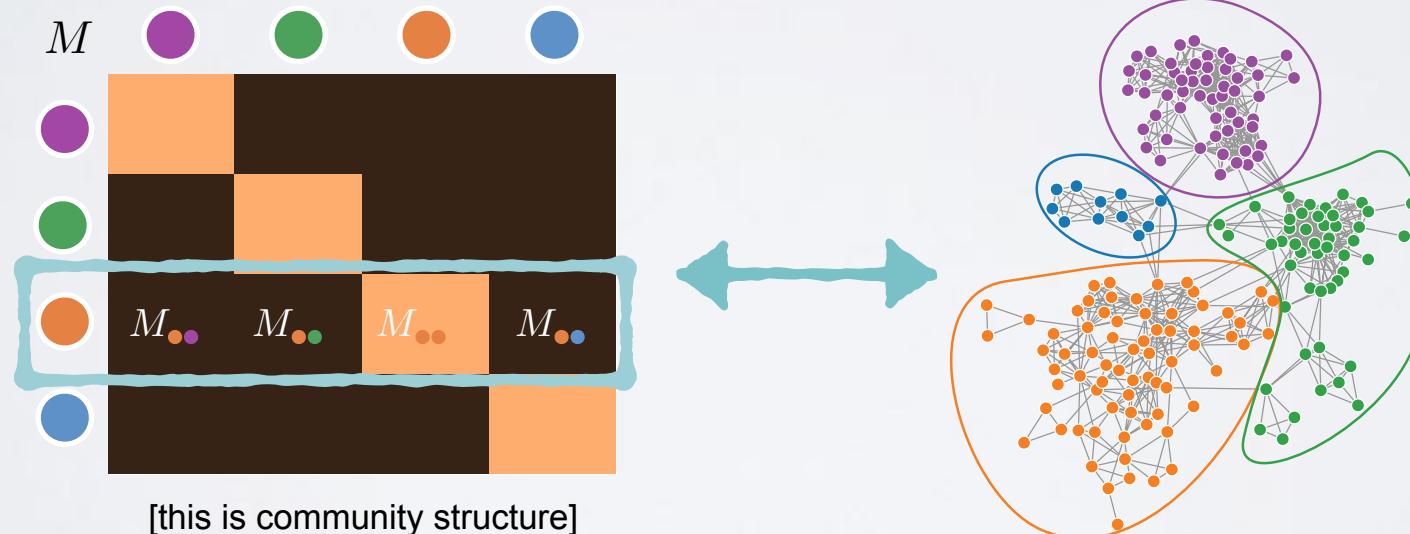


generative models for complex networks

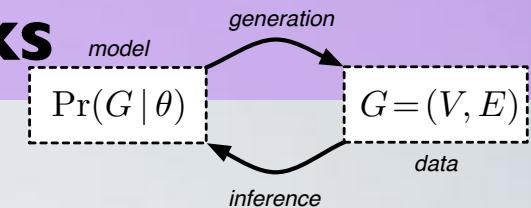


2. Stochastic block models (community structure)

- k groups of nodes: $\Pr(A_{ij} | M, z)$ depends only on the types z_i, z_j of the pair i, j
- M is a mixing matrix : $\Pr(i \rightarrow j) = M_{z_i, z_j}$

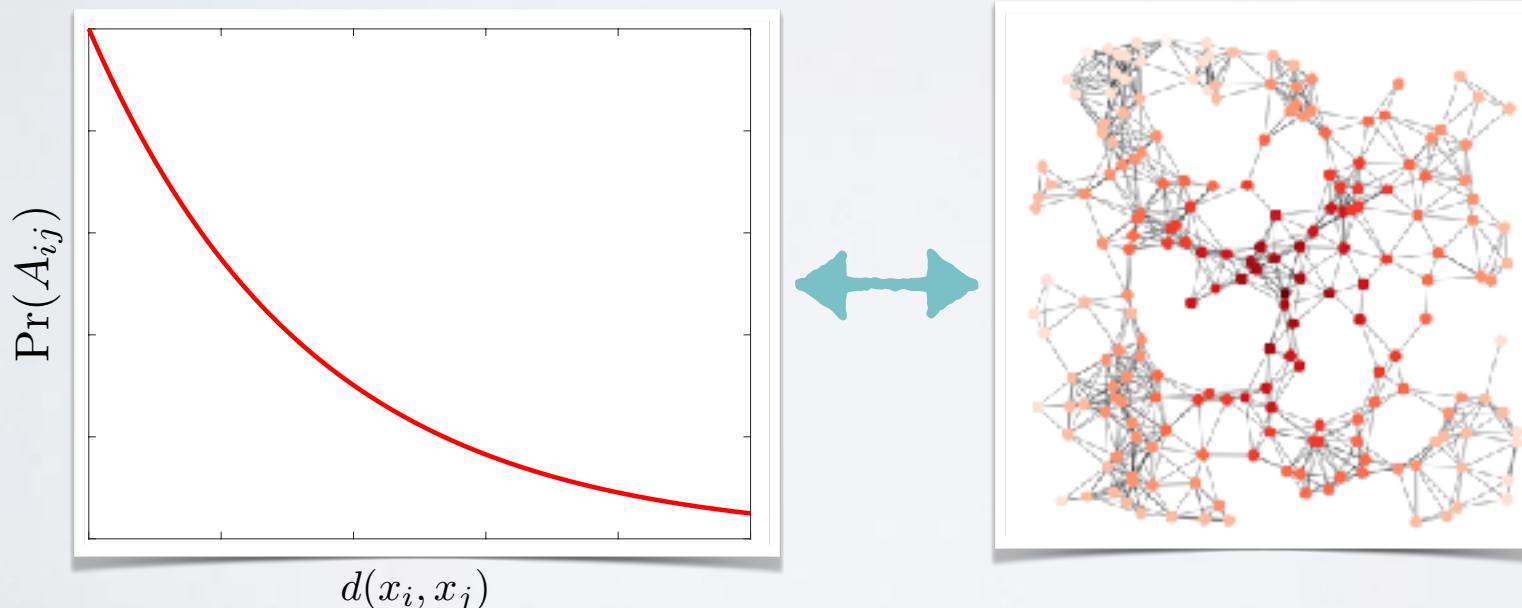


generative models for complex networks

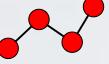
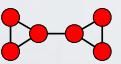


3. Latent space models (random geometric graphs)

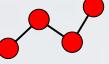
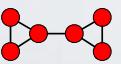
- nodes have position in latent space $x_i \in \mathbb{S}$
- $\Pr(A_{ij} | d(x_i, x_j))$ depends on distance $d(x_i, x_j)$ of the pair i, j



what patterns should we expect?

feature	real networks
	degree distribution
	clustering coefficient
	diameter
	large-scale structure

what patterns should we expect?

feature	real networks
	heavy tailed
	social: higher non-social: lower
	small, like $O(\ln n)$
	communities, dense core, hierarchies, etc.

Erdos-Renyi random graphs

denoted $G(n, p)$

where edges are iid $\Pr(A_{ij}) = p = \frac{c}{n - 1}$



mean degree

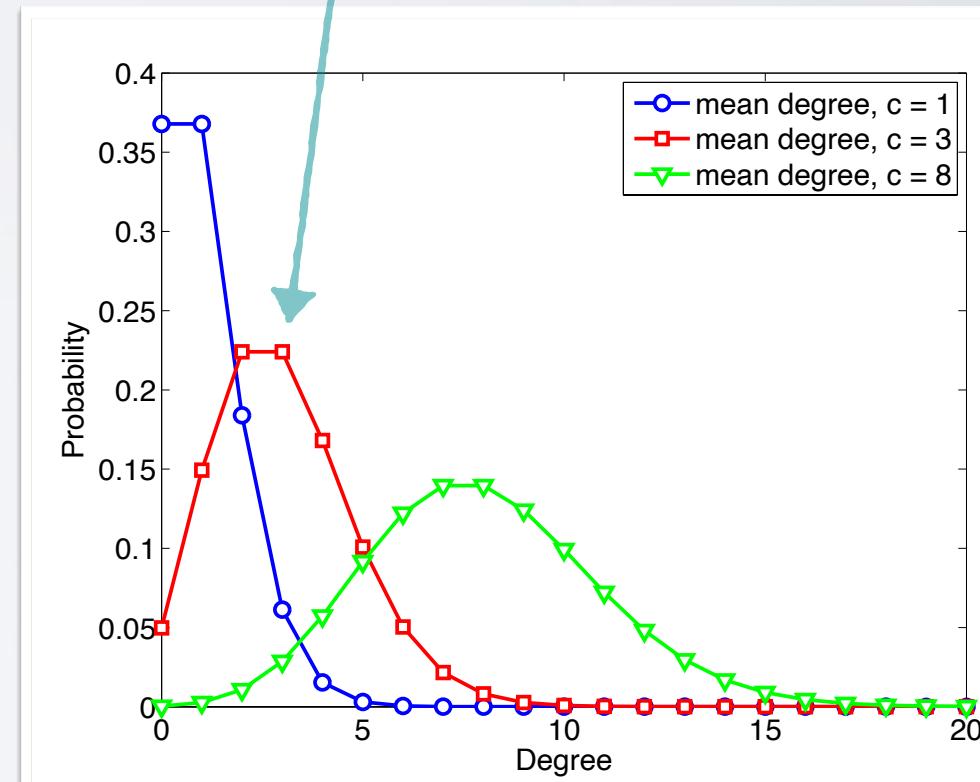
comments:

- highly *unrealistic* model (all edges iid)
- but, useful for building intuition & doing math
- the most well-studied random graph model
- warm up for more realistic models

degree distribution

mean degree: $\langle k \rangle = c = (n - 1)p$

degree distribution: $\Pr(k) = e^{-c} \frac{c^k}{k!}$ } Poisson distribution



degree distribution

mean degree: $\langle k \rangle = c = (n - 1)p$

degree distribution: $\Pr(k) = e^{-c} \frac{c^k}{k!}$ } Poisson distribution

clustering coefficient:

$$C = \frac{3 \times \text{\#triangles}}{\text{\#connected triples}}$$

$$= \frac{\binom{n}{3} p^3}{\binom{n}{3} p^2} = p = \frac{c}{n - 1} = \underbrace{O(n^{-1})}_{\text{asymptotically, zero clustering}}$$

asymptotically,
zero clustering

degree distribution

mean degree: $\langle k \rangle = c = (n - 1)p$

degree distribution: $\Pr(k) = e^{-c} \frac{c^k}{k!}$ } Poisson distribution

clustering coefficient: $C = O(n^{-1})$ } asymptotically, zero

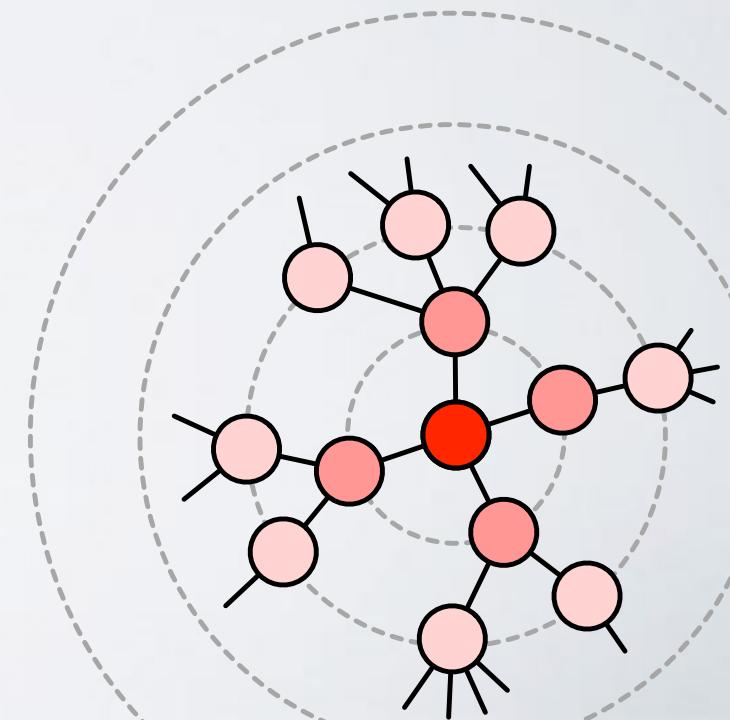
diameter: $G(n, p)$ is locally tree-like

mean number of vertices within s steps is c^s

all n vertices within ℓ steps

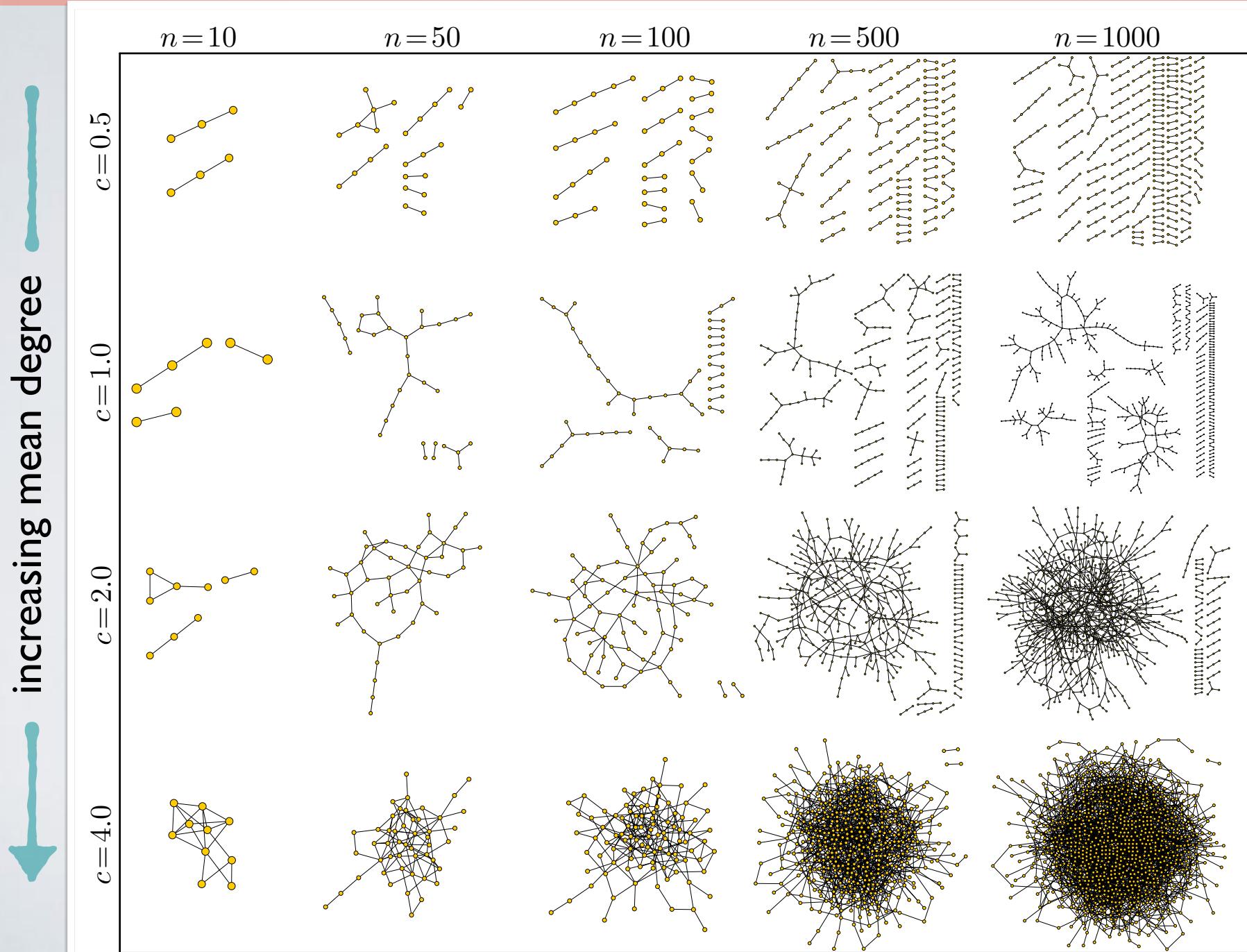
thus, diameter is $\underbrace{\ell = O(\ln n)}$

a "small" world

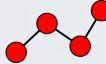
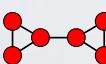


* this argument can be made more formal, but yields the same asymptotic result

examples of ER random graphs



how are we doing?

feature		$G(n, p)$	real networks
 degree distribution		Poisson	heavy tailed
 clustering coefficient		$O(n^{-1})$	social: higher non-social: lower
 diameter		$O(\ln n)$	small
 large-scale structure		none	communities, dense core, hierarchies, etc.

degree-based random graphs

configuration model : a random graph conditioned on having the specified degree sequence $\{k_1, k_2, \dots, k_n\}$

$$\Pr(i \rightarrow j) = \frac{k_i k_j}{2m}$$

* Fosdick et al. SIAM Review 60, 315-355 (2018)

* Chung & Lu, Ann. Comb. 6, 125-145 (2002) specifies a model that produces a simple graph with a given degree sequence in expectation

degree-based random graphs

configuration model : a random graph conditioned on having the specified degree sequence $\{k_1, k_2, \dots, k_n\}$

$$\Pr(i \rightarrow j) = \frac{k_i k_j}{2m}$$

double-edge swap algorithm:*

start with a graph G

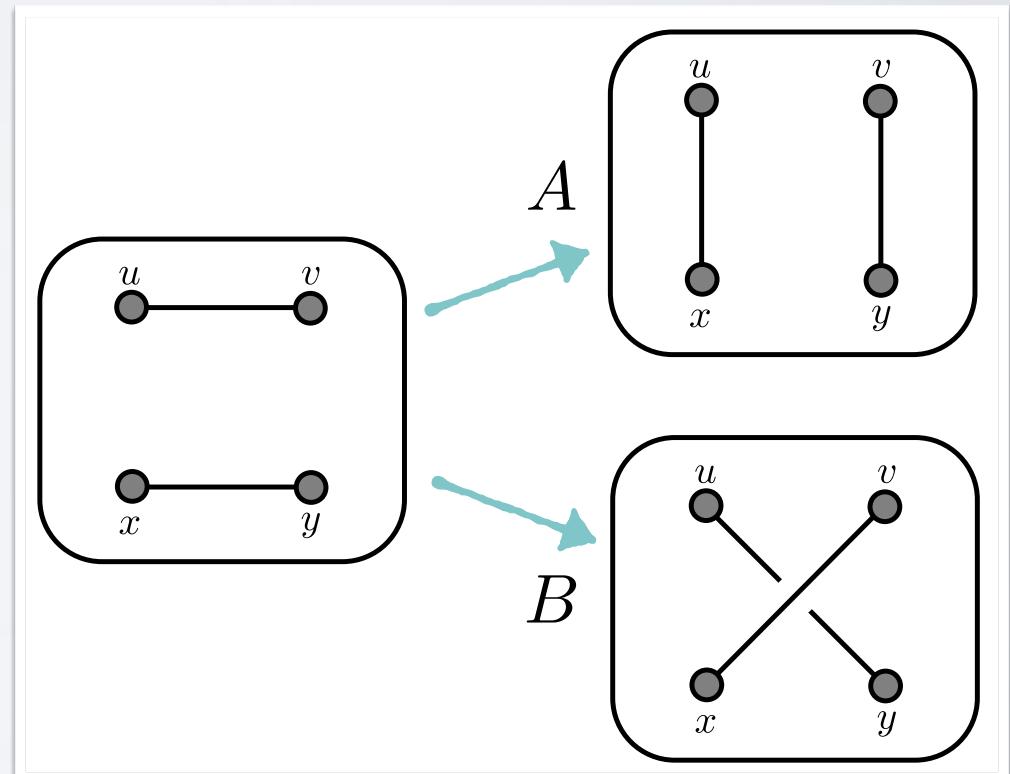
choose $\{(u, v), (x, y)\}$ uniformly

rearrange to A or B

repeat until convergence

degree preserving

record a G every $2m$



* we use the MCMC from Fosdick et al. SIAM Review (2018) [covers technical details]

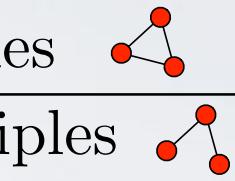
* we choose sampling gap and convergence time via Dutta et al. Preprint (2022)

degree-based random graphs

configuration model : a random graph conditioned on having the specified degree sequence $\{k_1, k_2, \dots, k_n\}$

clustering coefficient: $C = \frac{3 \times \# \text{triangles}}{\#\text{connected triples}}$

$$= \frac{1}{n} \frac{[\langle k^2 \rangle - \langle k \rangle]^2}{\langle k \rangle^3} = \underbrace{O(n^{-1})}_{\text{asymptotically, zero clustering}}$$



degree-based random graphs

configuration model : a random graph conditioned on having the specified degree sequence $\{k_1, k_2, \dots, k_n\}$

clustering coefficient: $C = O(n^{-1})$ } asymptotically, zero

diameter: also locally tree-like (if variance of degrees is finite)

following similar argument as ER graphs  $\underbrace{\ell = O(\ln n)}$
a "small" world

degree-based random graphs

the standard **null model** for empirical patterns

defines a probability distribution $\Pr(G \mid \vec{k})$

if $f(G_o)$ is "typical" within $\Pr(f(G) \mid \vec{k})$

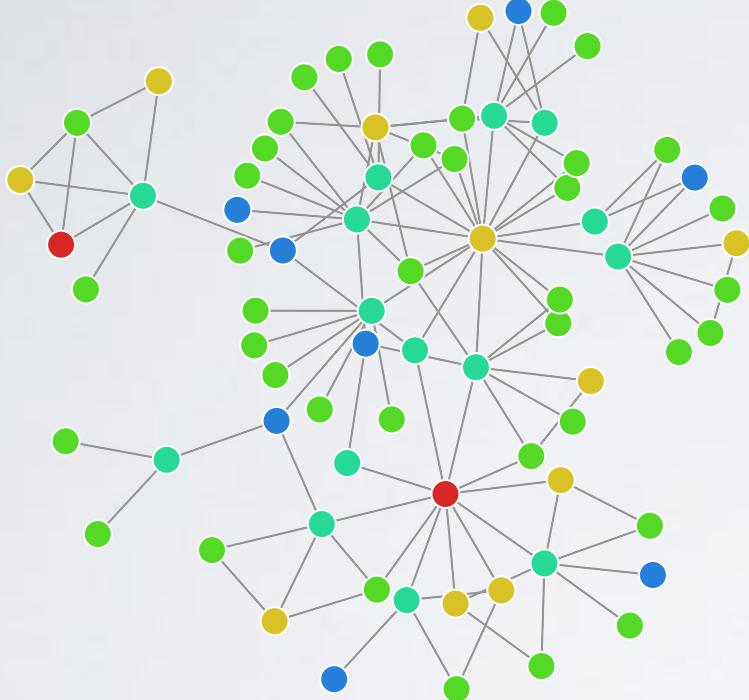
then we say that \vec{k} "explains" $f(G_o)$

e.g. from an empirical G_o
or a preferred $\Pr(k)$

* when a $\Pr(k)$ is drawn from a power-law distribution, we call these "power-law random graphs", which are a popular model for mathematical calculations

degree-based random graphs

the standard **null model** for empirical patterns



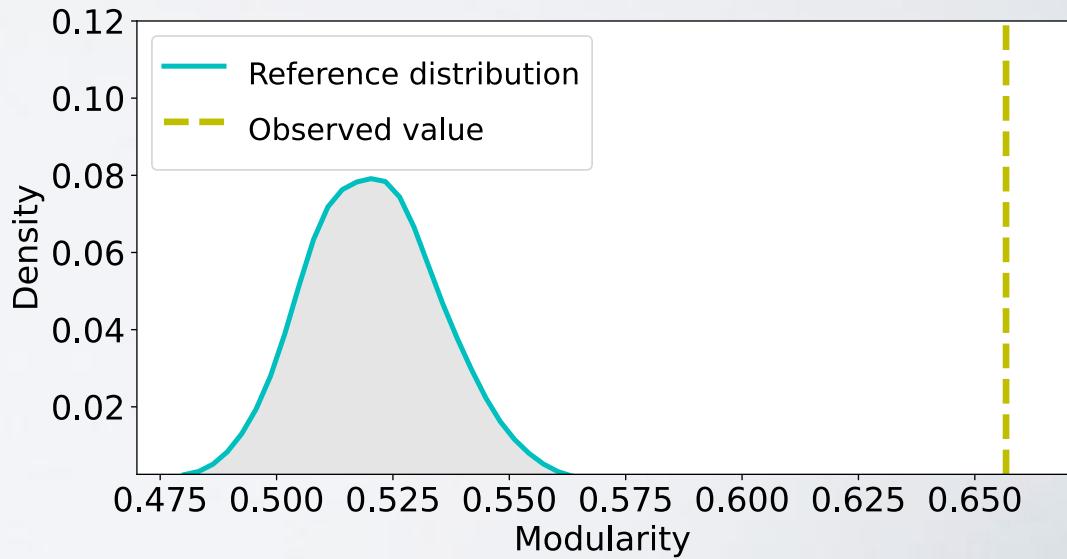
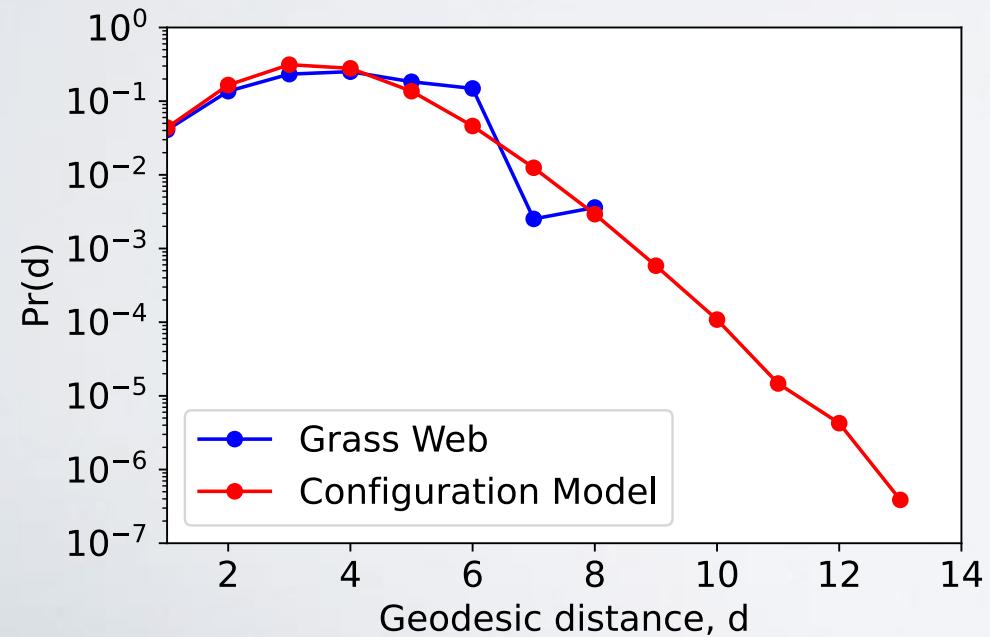
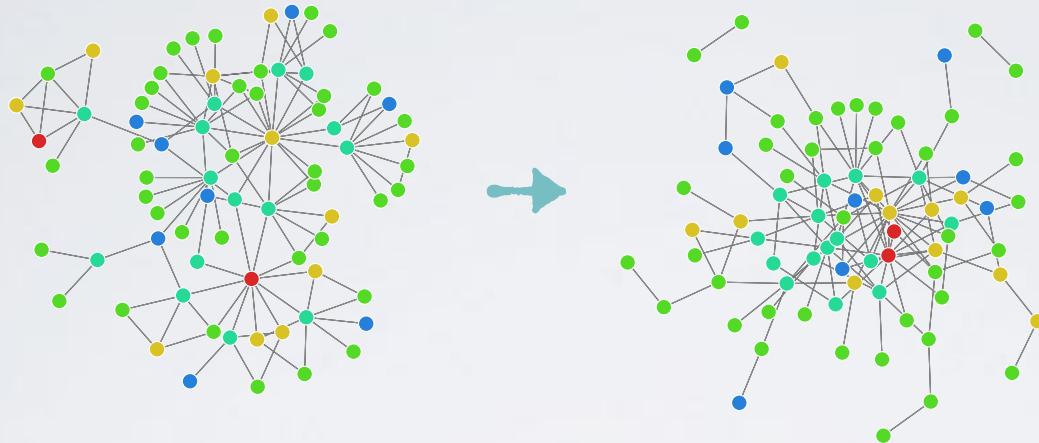
grassland species



grassland species
(randomized)

degree-based random graphs

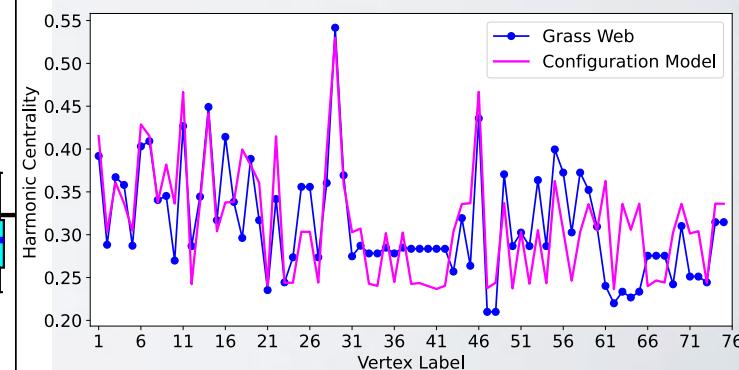
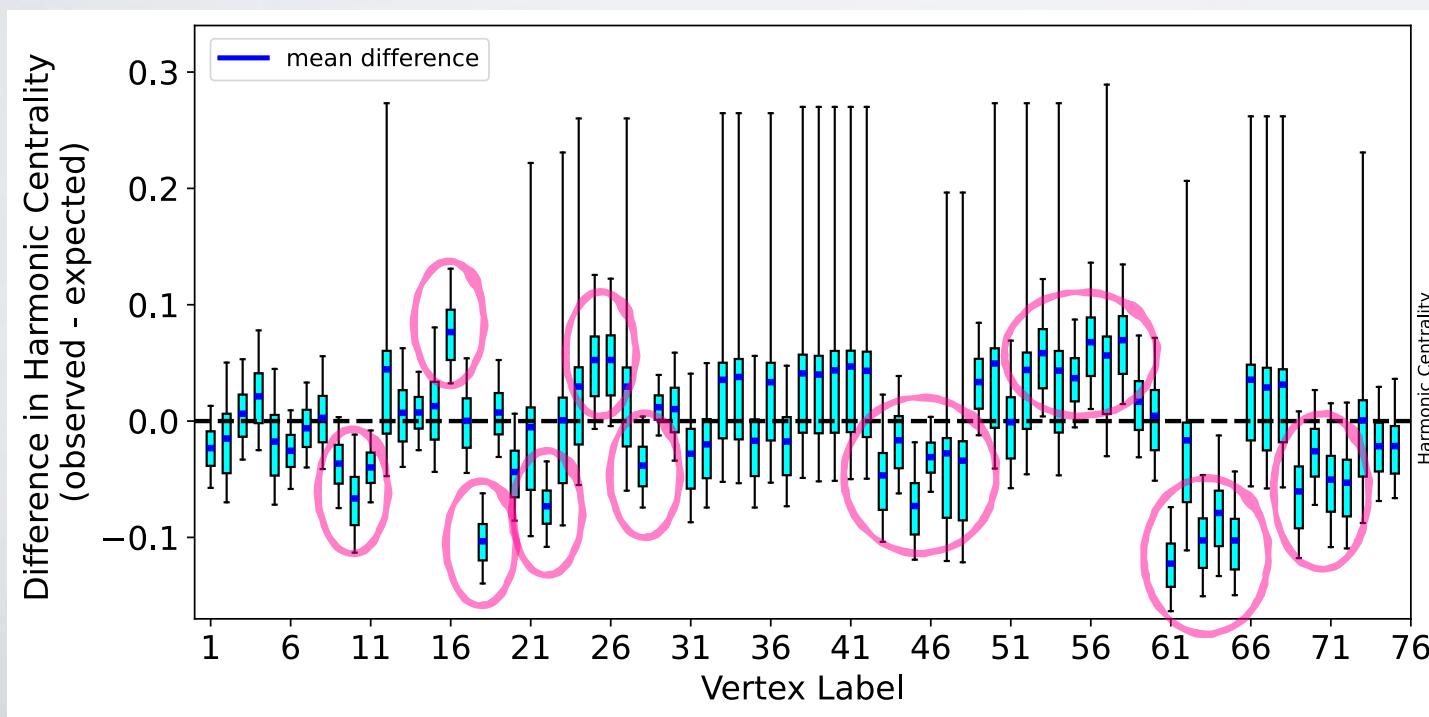
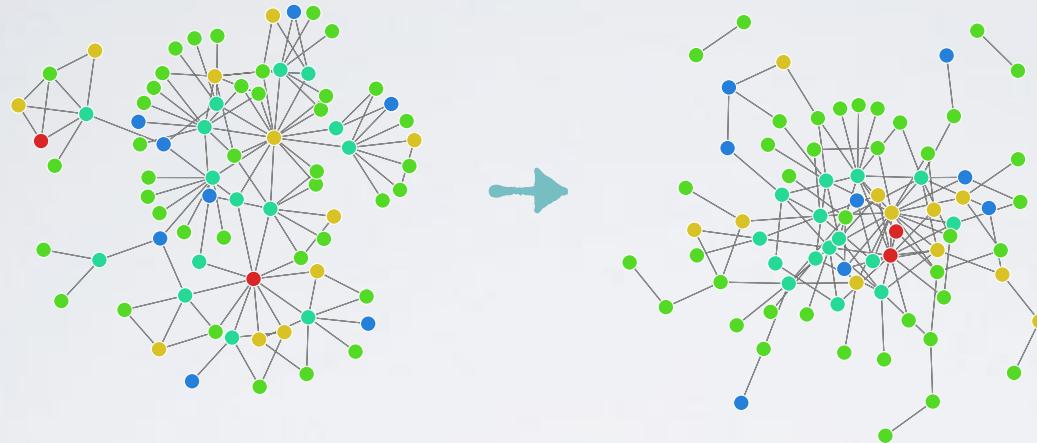
the standard **null model** for empirical patterns



* null distribution from 10^6 configuration models. what the configuration model gets wrong is the community structure. most everything else is well-explained by the degree structure alone

degree-based random graphs

the standard **null model** for empirical patterns



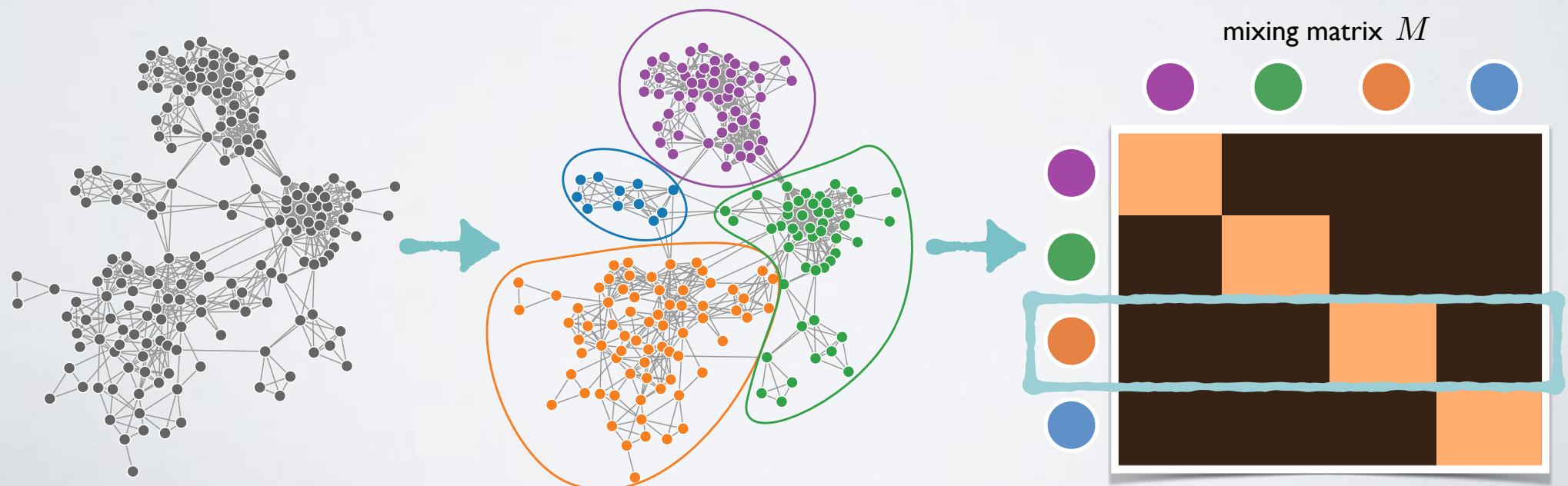
how are we doing?

feature			real networks
	$G(n, p)$	configuration	
degree distribution	Poisson	specified	heavy tailed
	$O(n^{-1})$	$O(n^{-1})$	social: higher non-social: lower
	$O(\ln n)$	$O(\ln n)$	small
	none	none	communities, dense core, hierarchies, etc.

stochastic block models

- each vertex i has type $z_i \in \{1, \dots, k\}$ (k vertex types or groups)
- stochastic block matrix M of group-level connection probabilities
- probability that i, j are connected = M_{z_i, z_j}

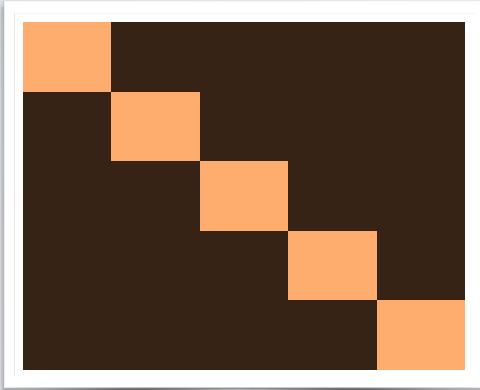
community = vertices with same pattern of inter-community connections



stochastic block models

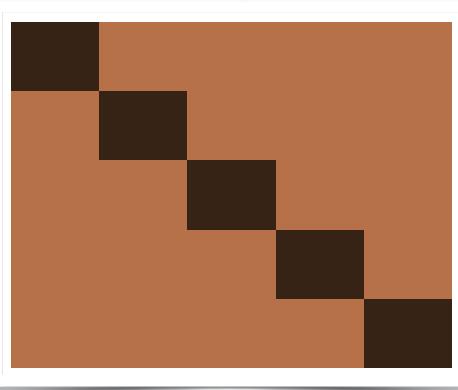
assortative

edges within groups



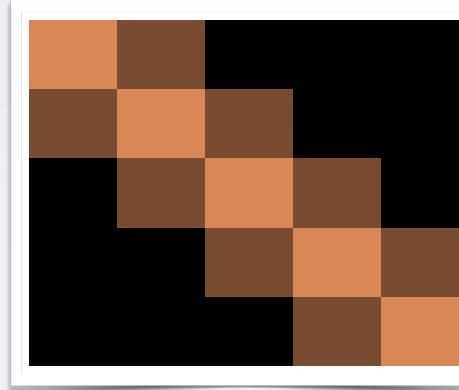
disassortative

edges between groups



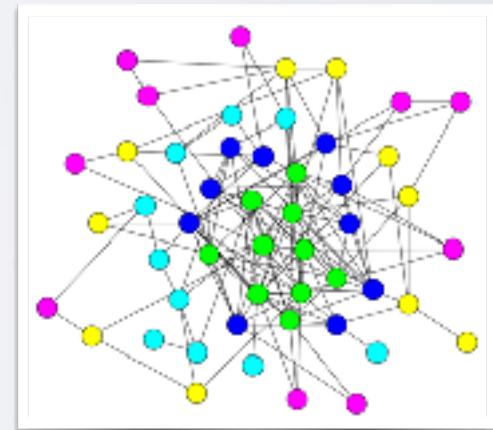
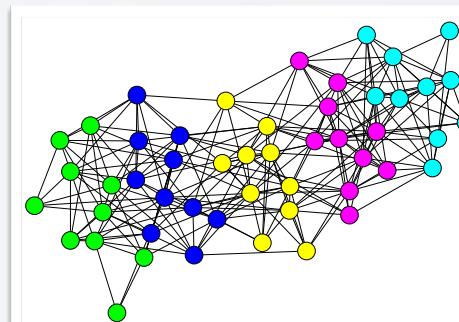
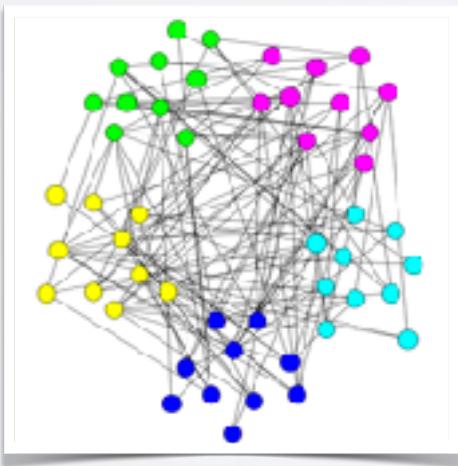
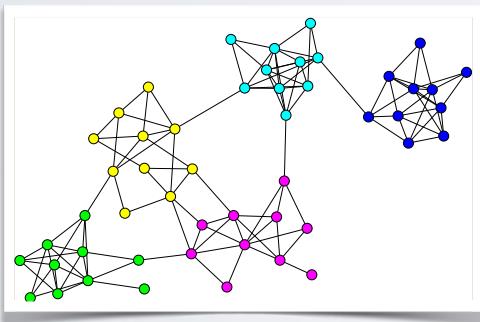
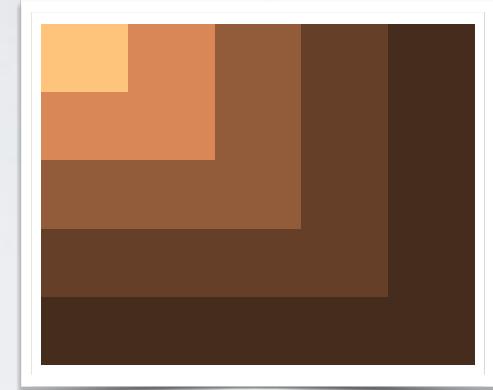
ordered

linear group hierarchy



core-periphery

dense core, sparse periphery



stochastic block models

likelihood function

the probability of G given labeling z and block matrix M

$$\Pr(G \mid z, M) = \underbrace{\prod_{(i,j) \in E} M_{z_i, z_j}}_{\text{edge}} / \underbrace{\prod_{(i,j) \notin E} (1 - M_{z_i, z_j})}_{\text{non-edge probability}}$$

stochastic block models

likelihood function

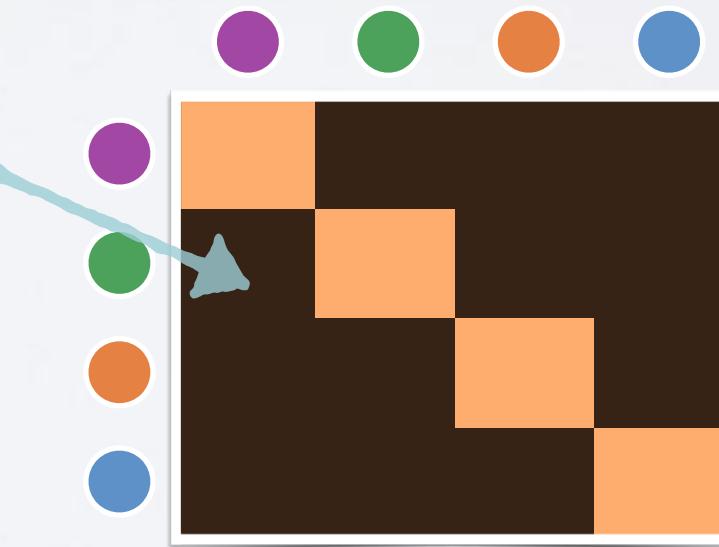
the probability of G given labeling z and block matrix M

$$\Pr(G \mid z, M) = \prod_{(i,j) \in E} M_{z_i, z_j} \prod_{(i,j) \notin E} (1 - M_{z_i, z_j})$$

$$= \prod_{rs} M_{r,s}^{e_{r,s}} (1 - M_{r,s})^{n_s n_r - e_{r,s}}$$

(Bernoulli edges)

Bernoulli random graph
with parameter $M_{r,s}$



stochastic block models

the most general SBM

$$\Pr(A \mid z, \theta) = \prod_{i,j} f(A_{ij} \mid \theta_{\mathcal{R}(z_i, z_j)})$$

A_{ij} : value of adjacency

\mathcal{R} : partition of adjacencies

f : probability function

$\theta_{a,*}$: pattern for a -type adjacencies

Binomial = simple graphs
Poisson = multi-graphs
Normal = weighted graphs
etc.

θ_{11}	θ_{12}	θ_{13}	θ_{14}
θ_{21}	θ_{22}	θ_{23}	θ_{24}
θ_{31}	θ_{32}	θ_{33}	θ_{34}
θ_{41}	θ_{42}	θ_{43}	θ_{44}

many stochastic block models

stochastic block models

k types of vertices, $\Pr(A_{ij} \mid M, z)$ depends only on node types z_i, z_j
originally invented by sociologists [Holland, Laskey, Leinhardt 1983]

many, many flavors, including

binomial SBM [Holland et al. 1983, Wang & Wong 1987]

simple assortative SBM [Hofman & Wiggins 2008]

mixed-membership SBM [Airoldi et al. 2008]

hierarchical SBM [Clauset et al. 2006,2008, Peixoto 2014]

fractal SBM [Leskovec et al. 2005]

infinite relational model [Kemp et al. 2006]

degree-corrected SBM [Karrer & Newman 2011]

SBM + topic models [Ball et al. 2011]

SBM + vertex covariates [Mariadassou et al. 2010, Newman & Clauset 2016]

SBM + edge weights [Aicher et al. 2013,2014, Peixoto 2015]

bipartite SBM [Larremore et al. 2014]

multilayer SBM [Peixoto 2015, Valles-Catata et al. 2016]

and many others

one important stochastic block model

degree-corrected SBM ($f = \text{Poisson}$)

one important stochastic block model

degree-corrected SBM ($f = \text{Poisson}$)

key assumption $\Pr(i \rightarrow j) = \theta_i \theta_j \omega_{z_i, z_j}$

stochastic block matrix $\omega_{r,s}$

(degree) propensity of node θ_i

likelihood:

$$\Pr(A \mid z, \theta, \omega) = \prod_{i < j} \frac{(\theta_i \theta_j \omega_{z_r, z_j})^{A_{ij}}}{A_{ij}!} \exp(-\theta_i \theta_j \omega_{z_r, z_j})$$

where $\hat{\theta}_i = \underbrace{\frac{k_i}{\sum_j k_j \delta_{z_i, z_j}}}_{\text{fraction of } i\text{'s group's stubs on } i}$

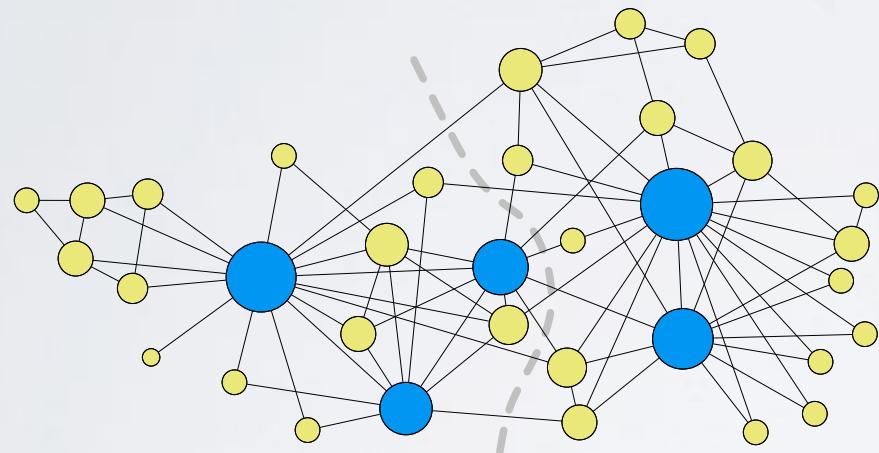
$$\hat{\omega}_{rs} = m_{rs} = \underbrace{\sum_{ij} A_{ij} \delta_{z_i, r} \delta_{z_j, s}}_{\text{total number of edges between } r \text{ and } s}$$

one important stochastic block model

comparing SBM vs. DC-SBM : Zachary karate club

one important stochastic block model

comparing SBM vs. DC-SBM : Zachary karate club



SBM

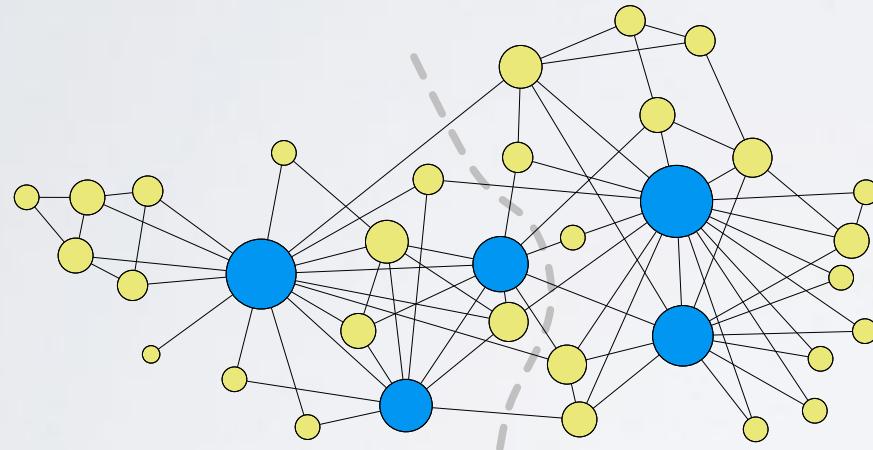
leader/follower division

DC-SBM

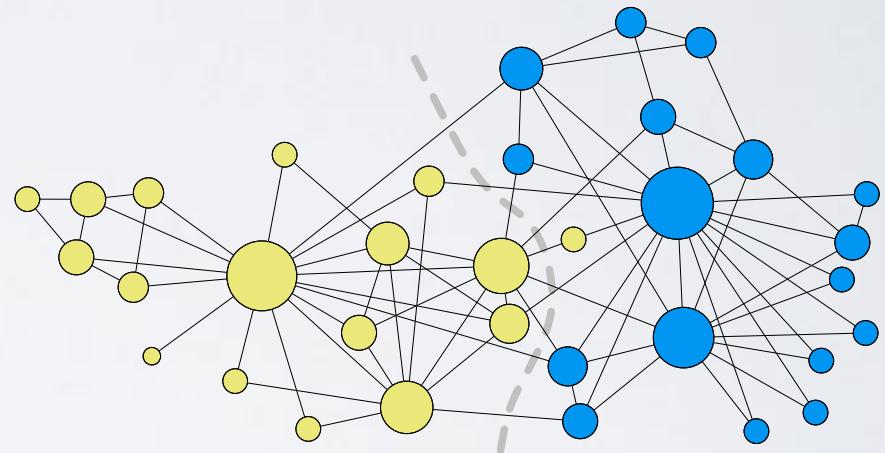
assortative group division

one important stochastic block model

comparing SBM vs. DC-SBM : Zachary karate club



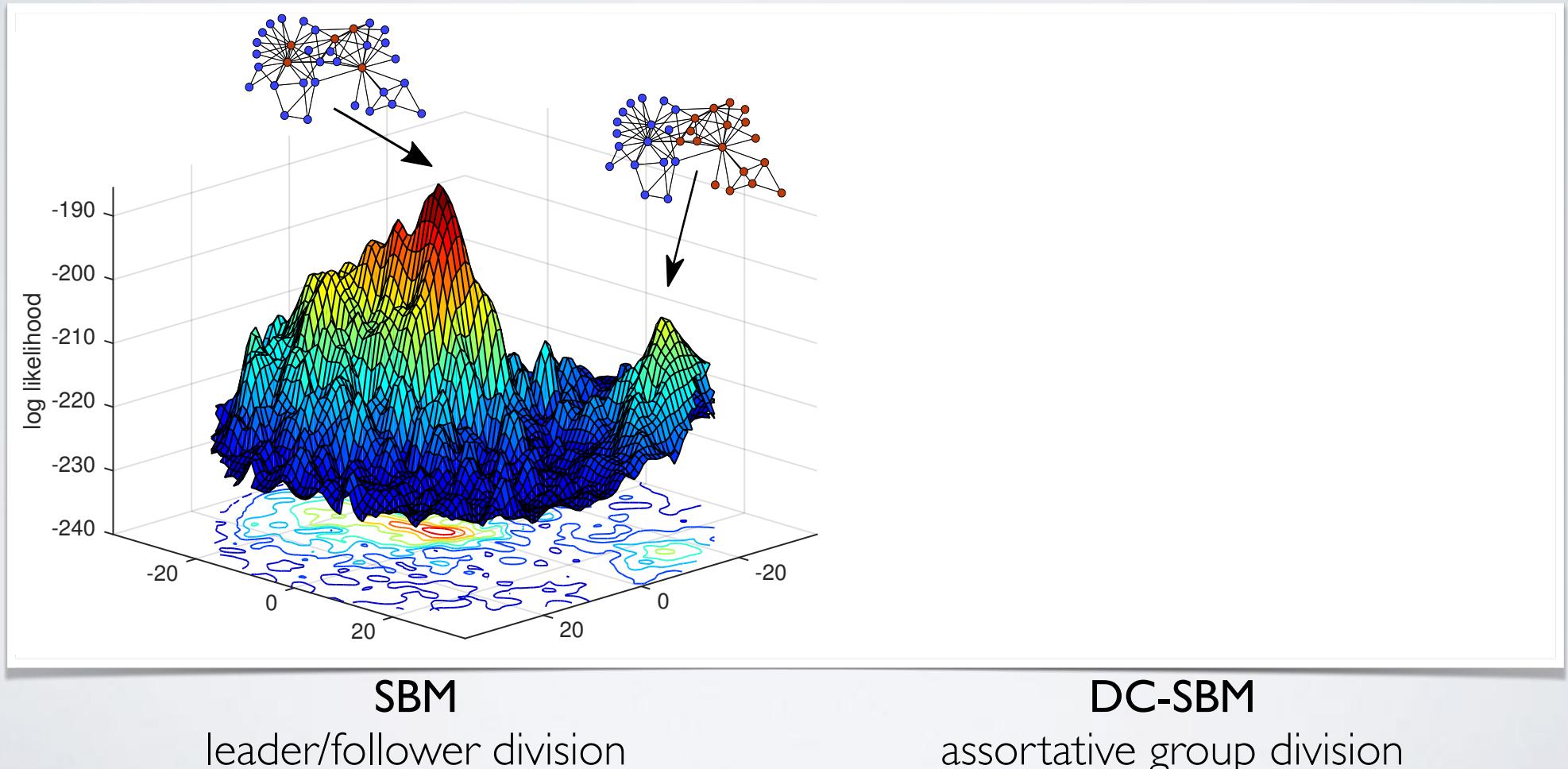
SBM
leader/follower division



DC-SBM
assortative group division

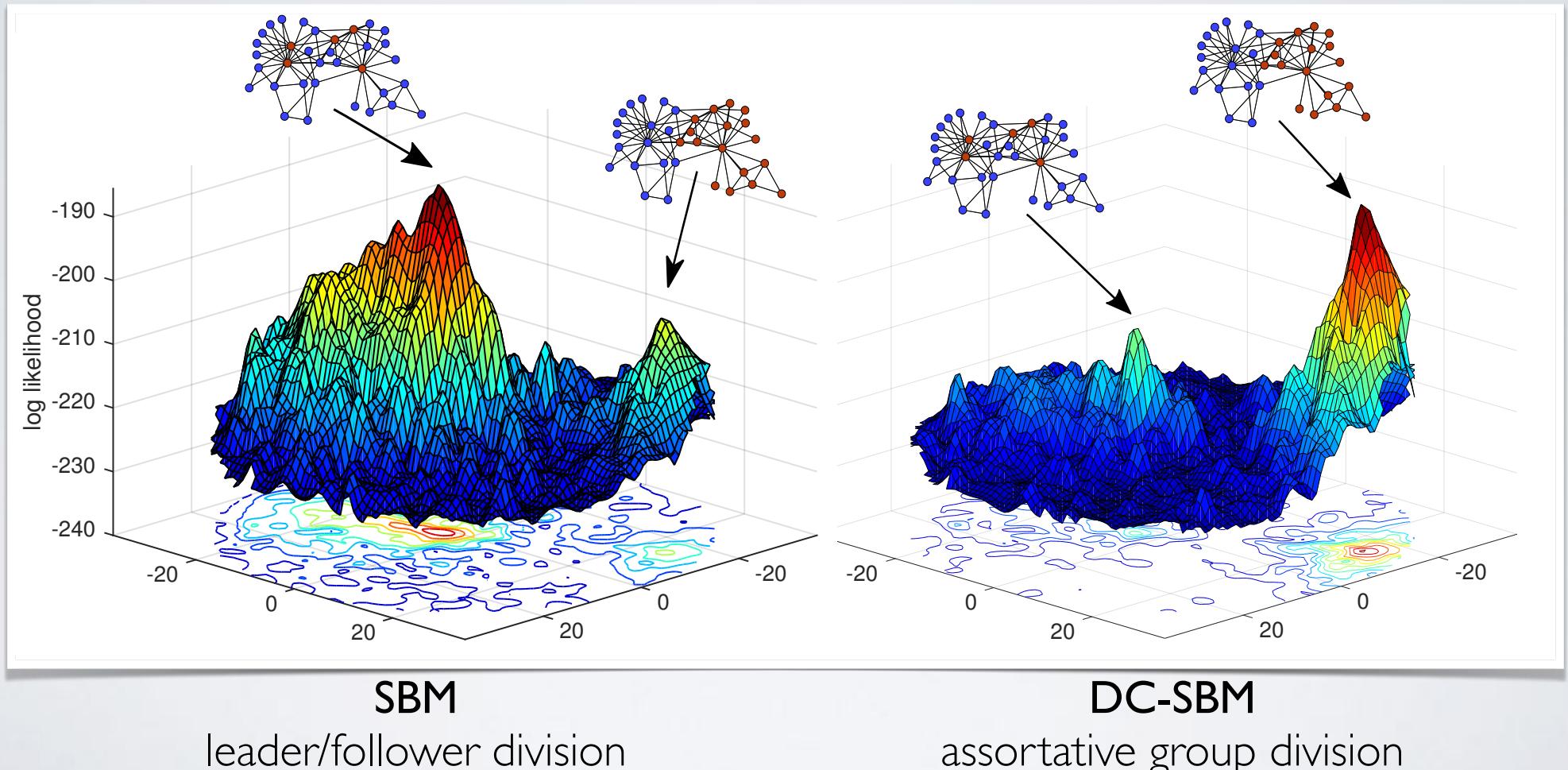
different models, different insights

comparing SBM vs. DC-SBM : Zachary karate club



different models, different insights

comparing SBM vs. DC-SBM : Zachary karate club



stochastic block models

SBM properties

k Erdos-Renyi random graphs

each with size n_r and internal density $M_{r,r}$

joined pairwise as random bipartite graph with density $M_{r,s}$

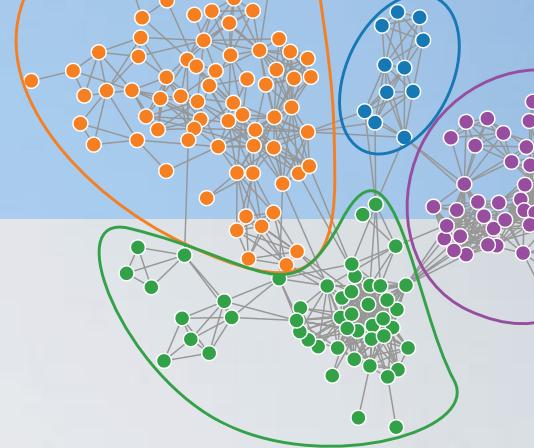
degree distribution: mixture of Poissons

diameter: $O(\ln n)$ or $O(\ln(kn))$

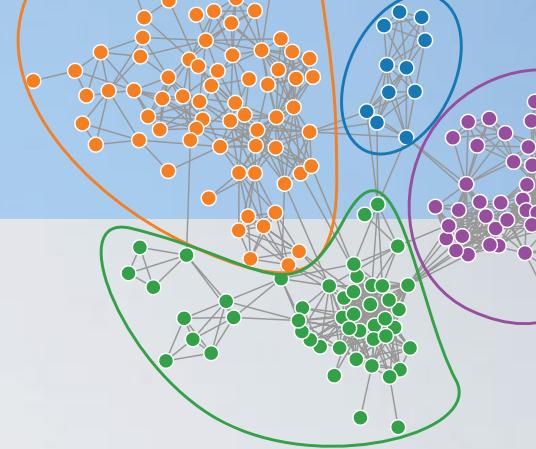
triangle density: low, except when $M_{r,s} \gg 0$

local structure: like a random graph

large-scale: mixtures of assortative & disassortative structure



stochastic block models



DC-SBM properties

k 'configuration model' random multi-graphs

each with size n_r , internal density $M_{r,r}$ and propensities $\{\theta_i\}_r$

joined pairwise as random bipartite graph with parameters $M_{r,s}$ and $\{\theta_i\}_{r,s}$

degree distribution: arbitrary $(\{\theta_i\})$

diameter: $O(\ln n)$ or $O(\ln(kn))$

triangle density: low, except when $M_{r,s} \gg 0$

local structure: like a random multi-graph

large-scale: mixtures of assortative & disassortative structure

how are we doing?

feature	$G(n, p)$	configuration	DC SBM	real networks
	Poisson	specified	specified	heavy tailed
	$O(n^{-1})$	$O(n^{-1})$	$O(n^{-1})$	social: high non-social: low
	$O(\ln n)$	$O(\ln n)$	$O(\ln n)$	small
	none	none	specified: communities, hierarchies, etc.	communities, dense core, hierarchies, etc.

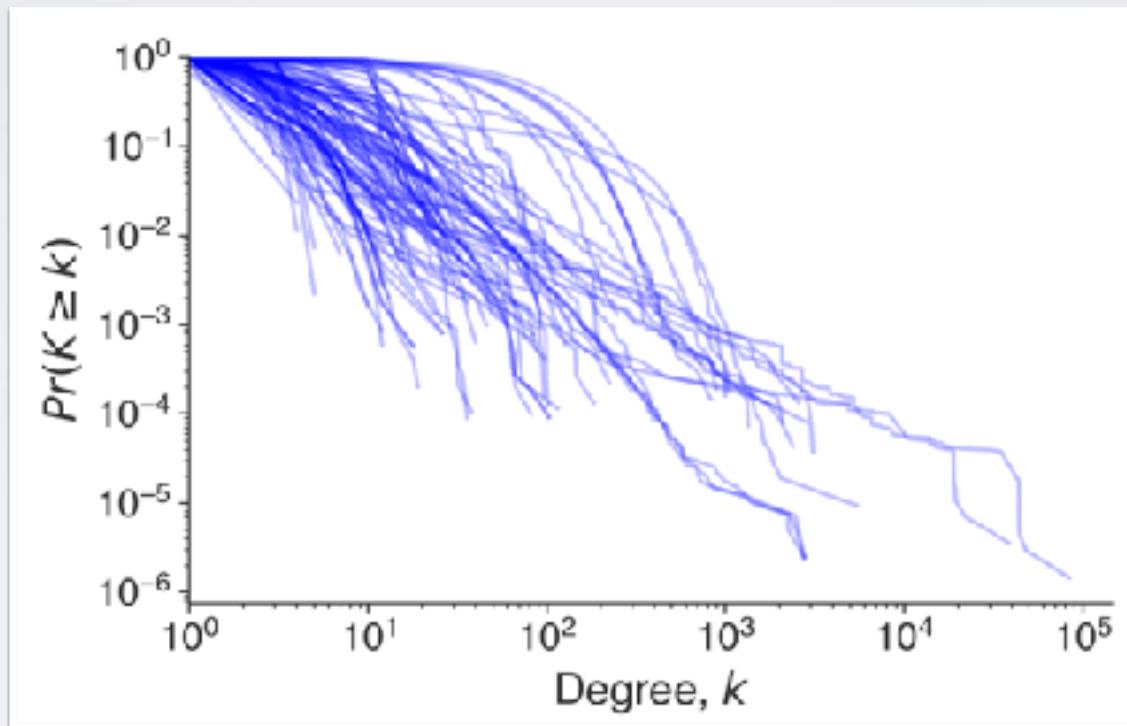
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what patterns do real networks exhibit?

degree distributions:

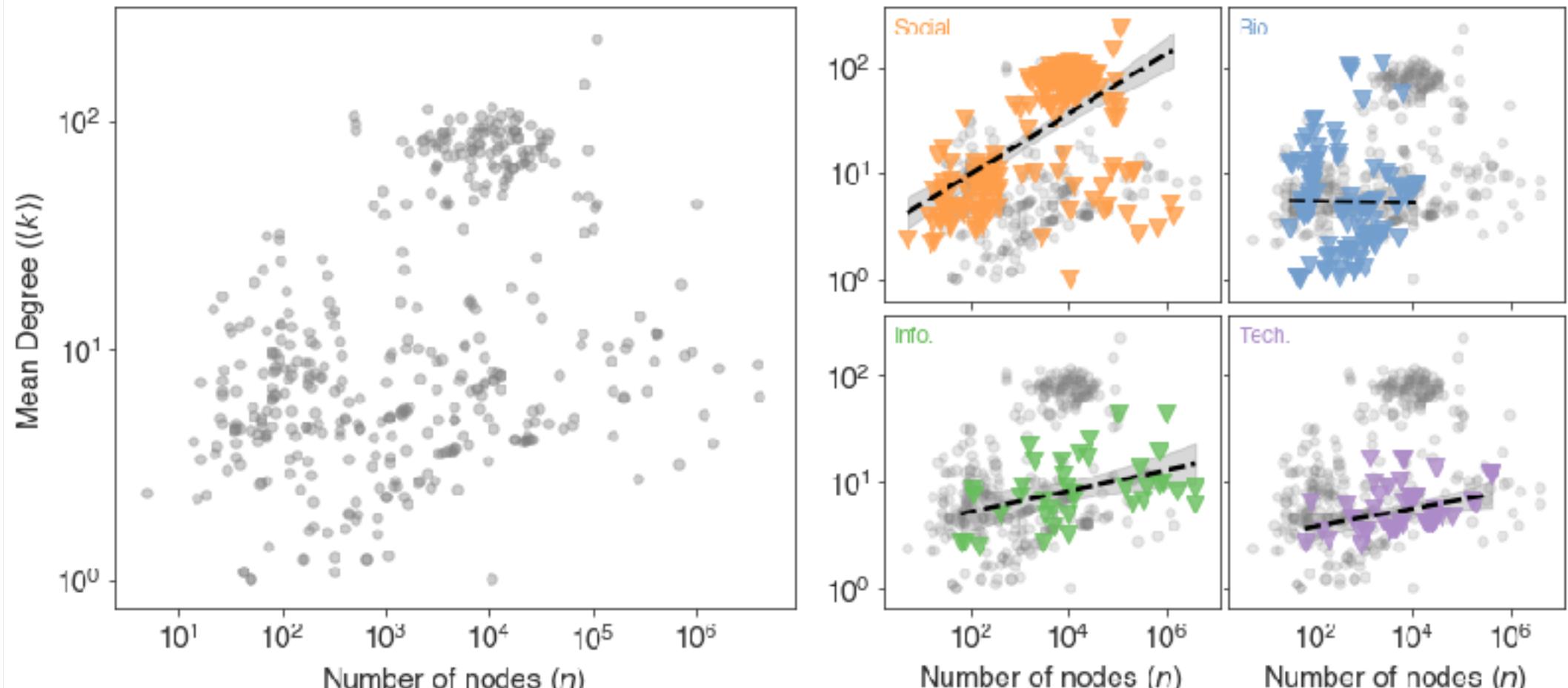
- ✓ heavy-tailed, with enormous diversity across networks and domains



what patterns do real networks exhibit?

mean degree (are networks sparse?):

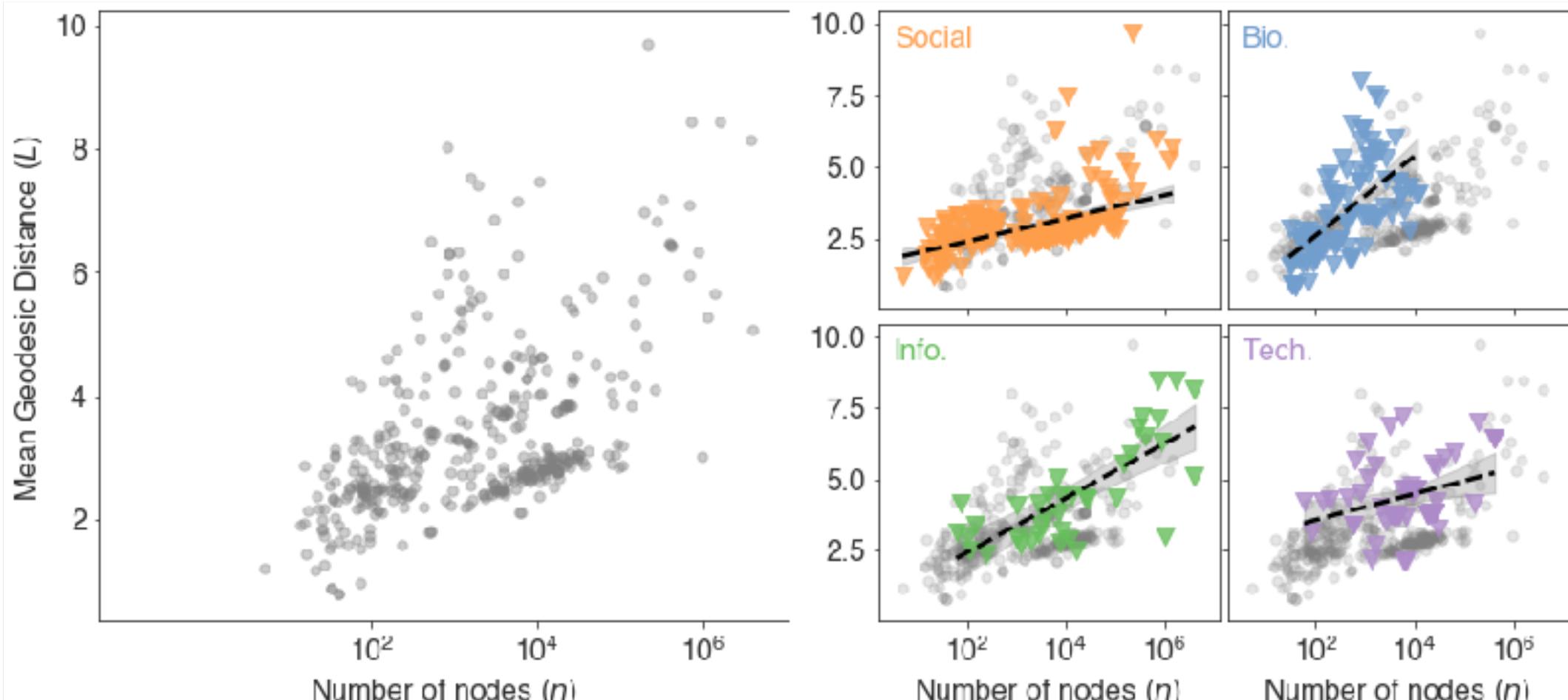
🤔 $O(n^\alpha)$, social networks generally far more dense than other types



what patterns do real networks exhibit?

mean geodesic distance (also, diameter):

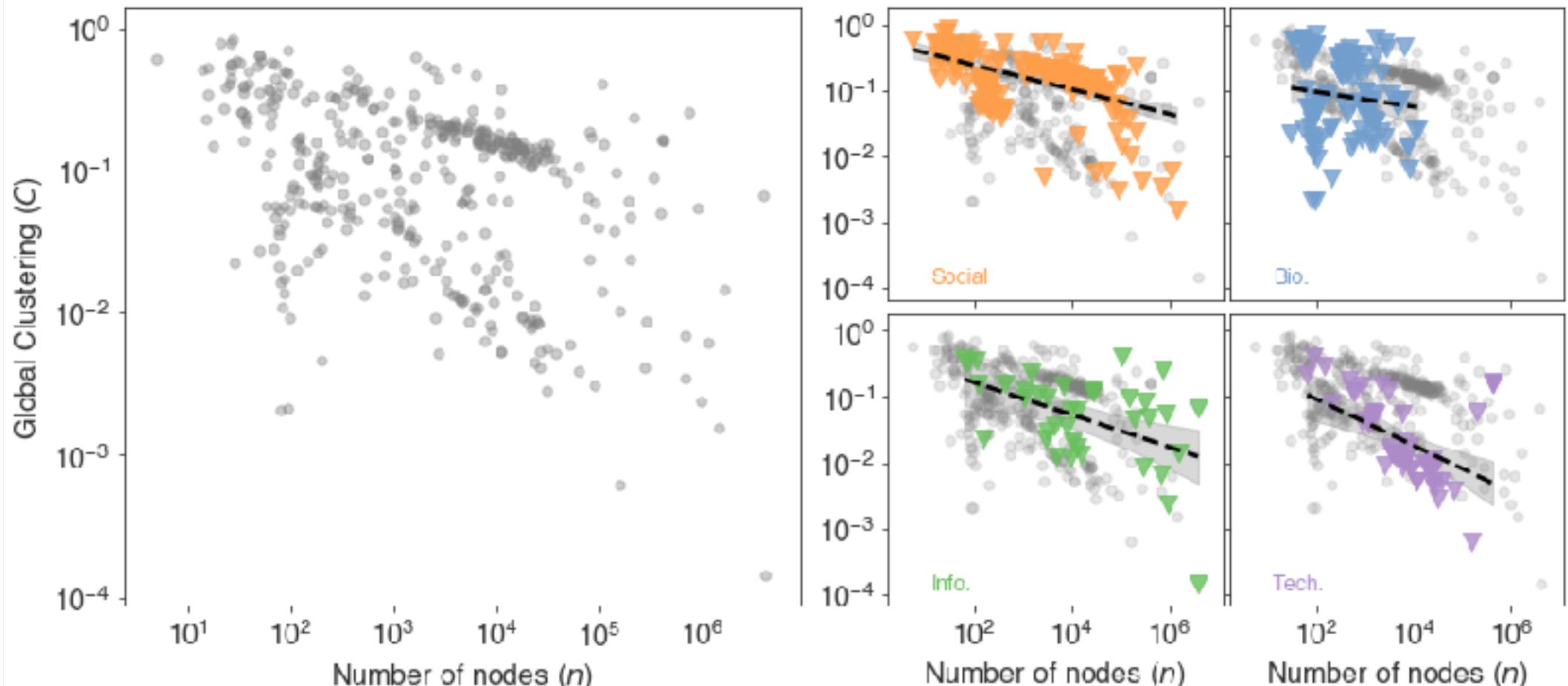
🤔 $O(\ln n)$, but with *different coefficients* for different domains



what patterns do real networks exhibit?

clustering coefficient:

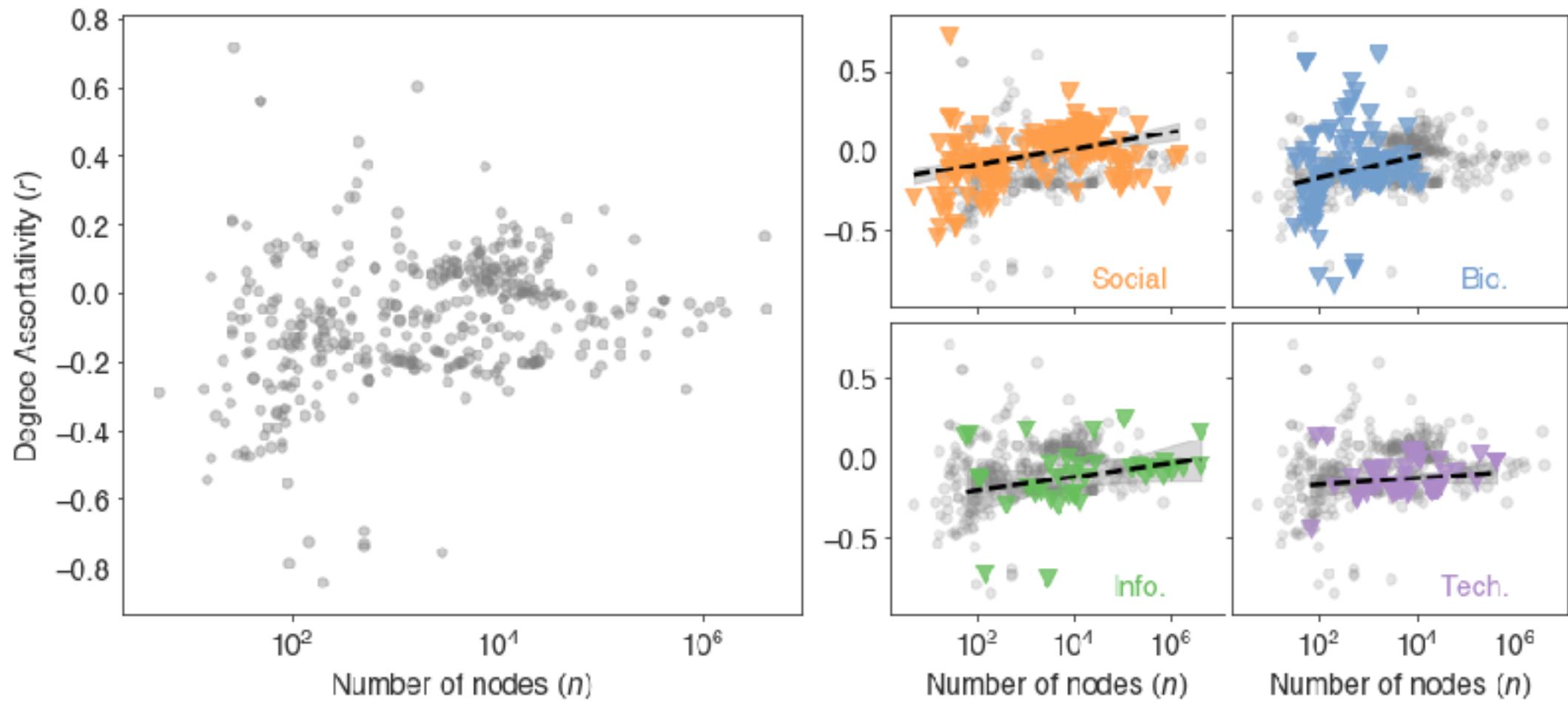
😱 $O(n^{-1})$, social networks have $5 - 10 \times$ more triangles at a given scale n , but all networks scale down



what patterns do real networks exhibit?

degree assortativity

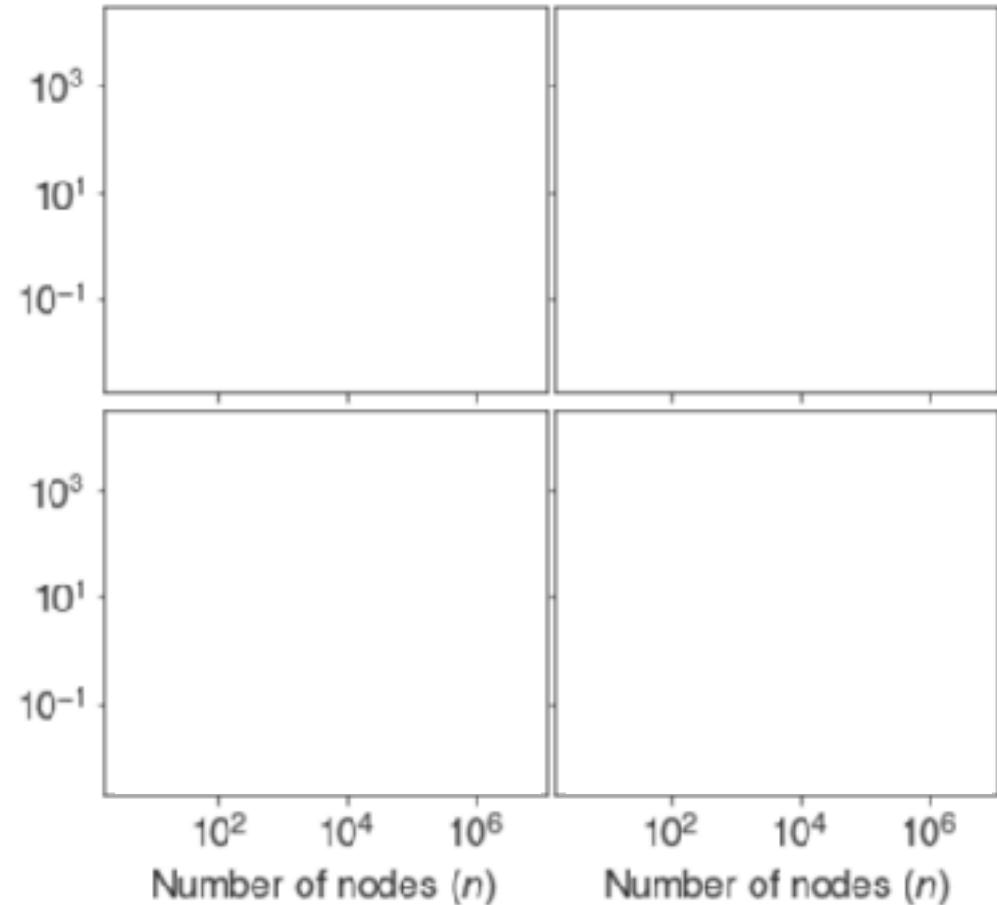
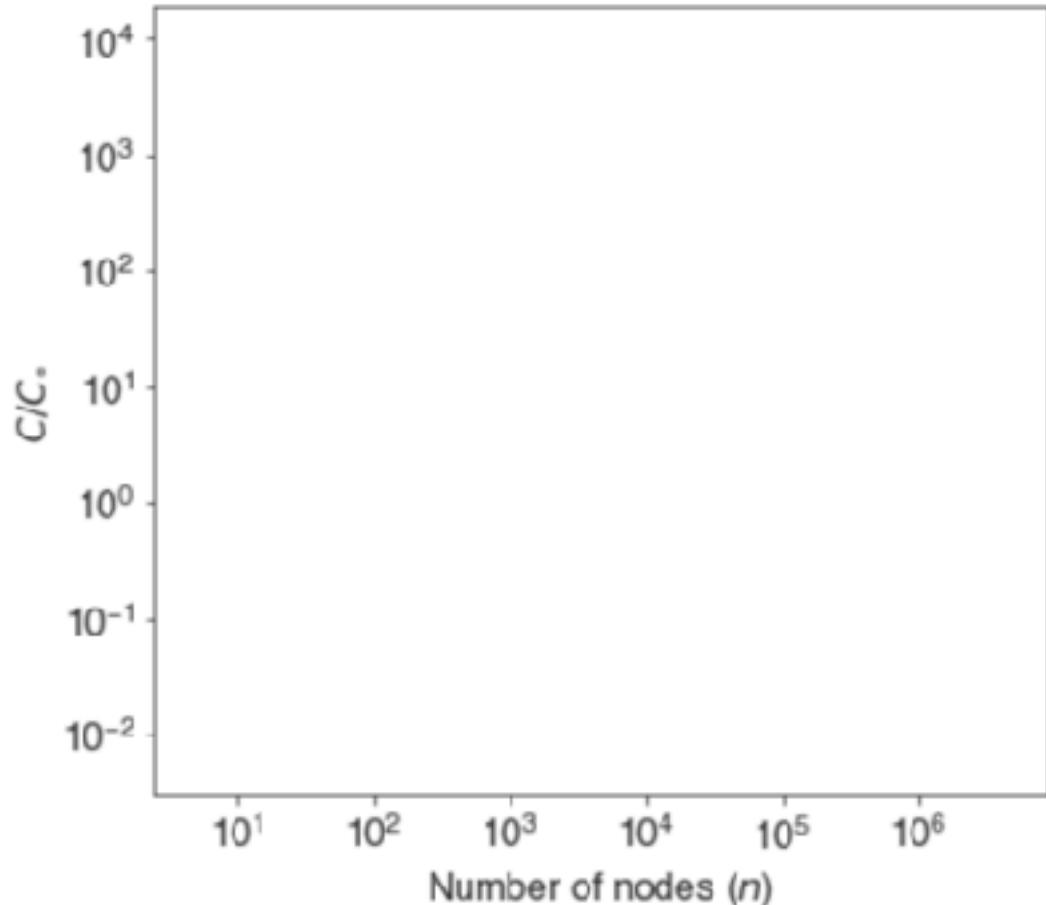
increases with scale — esp. in social networks



what patterns do real networks exhibit?

how much of clustering coefficient is due to degree structure?

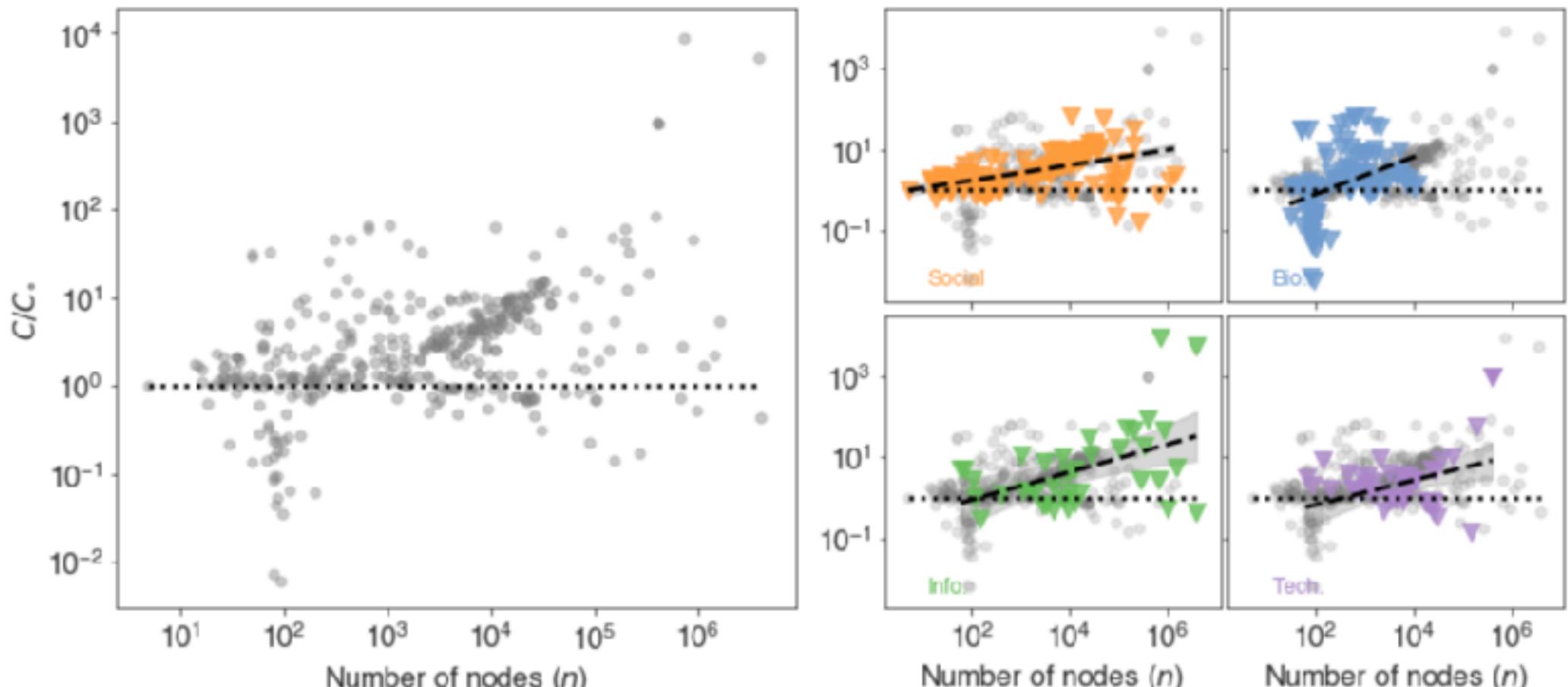
null models! compare empirical vs. configuration model: C/C_o



what patterns do real networks exhibit?

how much of clustering coefficient is due to degree structure?

- 的社会 networks' higher C is partly *explained* by their degree distributions
all domains exhibit similar triangle-enrichment across scales (a bit more for bio)

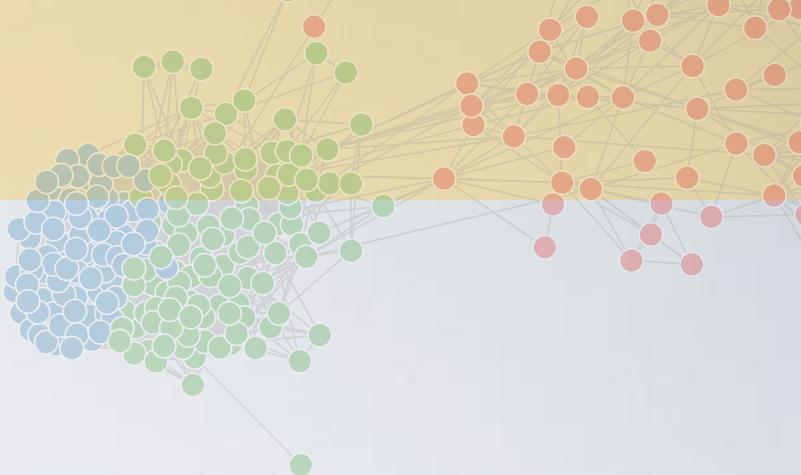


how are we doing?

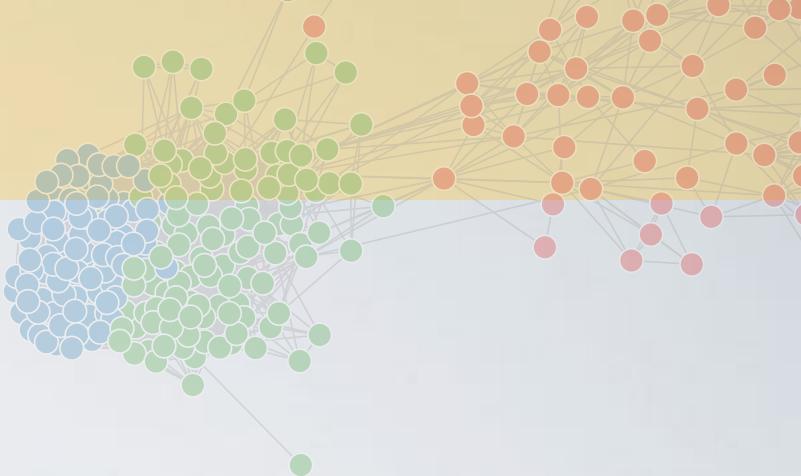
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parting thoughts on networks

- networks are cool!

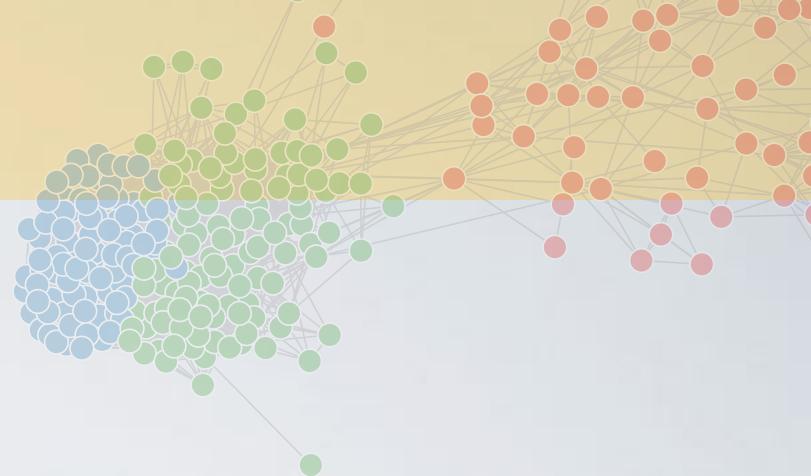


parting thoughts on networks



- **networks are cool!**
 - but also complicated objects = enormous structural diversity
 - many ways to describe a network's structure
- **null models & statistical inference**
 - among most powerful tools for describing network structure
 - highly flexible, scalable, useful
 - auxiliary data (weights, attributes, time)
 - applications abound [new ideas often come from these]
- **structure + dynamics = function**
 - how does structure constrain dynamics, robustness, etc.
 - to what degree does structure = function?

analyzing networks



6 major approaches

- ★ 1. **exploratory data analysis:** count & compare all the things (degree distributions, centrality scores, community detection, etc.)
- 2. **simple regressions:** convert network structure into node-level features, and do traditional explanatory modeling
- ★ 3. **null models:** use some kind of random graph to identify non-random patterns as deviations from the null
- 4. **mechanisms / simulations:** explain structural or dynamical patterns as caused by specific process
- ★ 5. **predictive models:** fit parametric model of network structure & use it to predict missing or future data (edges, labels, etc.)
- 6. **network experiments:** manipulate structure and measure node-level or graph-level behavior as function of changes

end of lecture 3

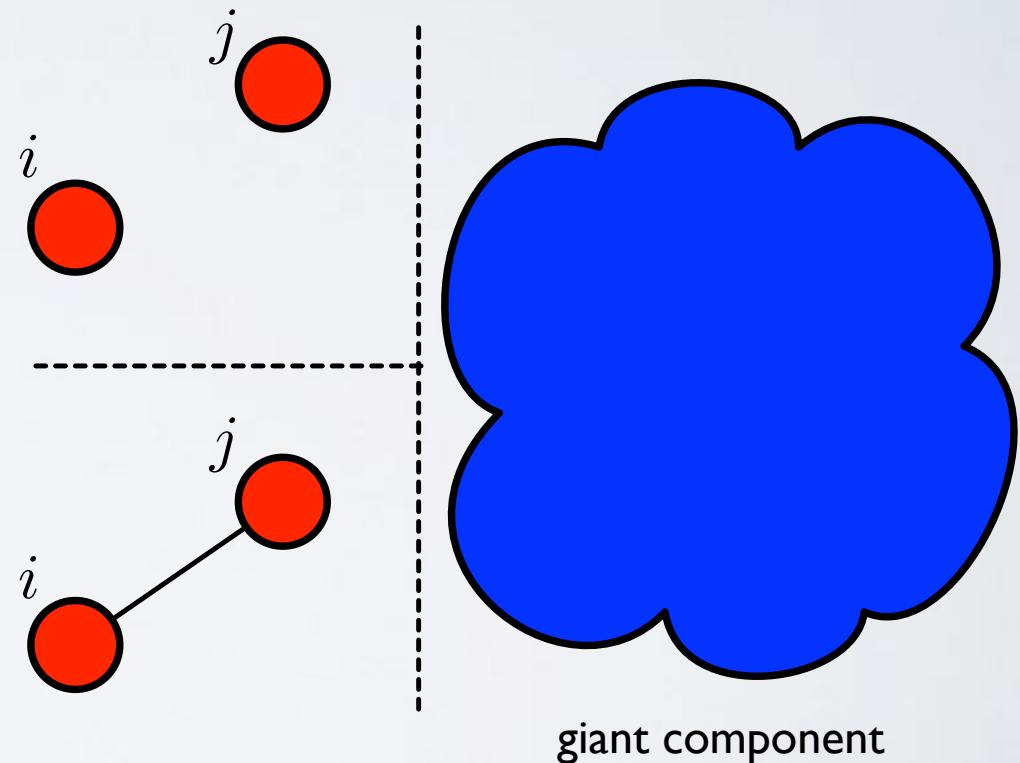


giant component (Erdos-Renyi graph)

let u be fraction of vertices *not* in **giant component**

for i not to be in the giant component, then for every j

1. i is not connected to j ,
or
2. i connects to j , and j is
not part of the giant
component



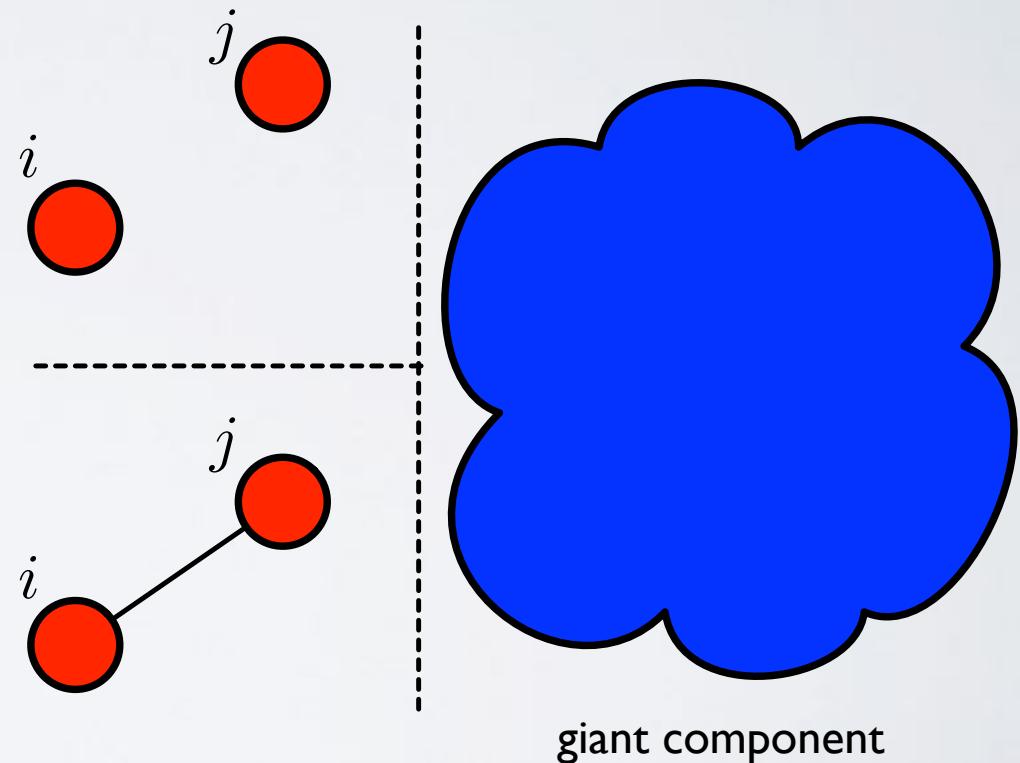
giant component (Erdos-Renyi graph)

let u be fraction of vertices *not* in **giant component**

for i not to be in the giant component, then for every j

1. with probability $1 - p$

2. with probability pu



giant component (Erdos-Renyi graph)

total probability that i **not** in giant component via
any of the $n - 1$ choices of j :

$$u = (1 - p + pu)^{n-1} = \left[1 - \frac{c}{n-1}(1-u) \right]^{n-1}$$

giant component (Erdos-Renyi graph)

total probability that i **not** in giant component via any of the $n - 1$ choices of j :

$$u = (1 - p + pu)^{n-1} = \left[1 - \frac{c}{n-1}(1-u) \right]^{n-1}$$

taking logs of both sides, and approximating:

$$\begin{aligned}\ln u &= (n-1) \ln \left[1 - \frac{c}{n-1}(1-u) \right] \\ &\approx -(n-1) \frac{c}{n-1}(1-u) \\ &= -c(1-u)\end{aligned}$$

giant component (Erdos-Renyi graph)

total probability that i **not** in giant component via any of the $n - 1$ choices of j :

$$u = e^{-c(1-u)}$$

and the fraction of vertices **in** the giant component is

$$S = 1 - u$$

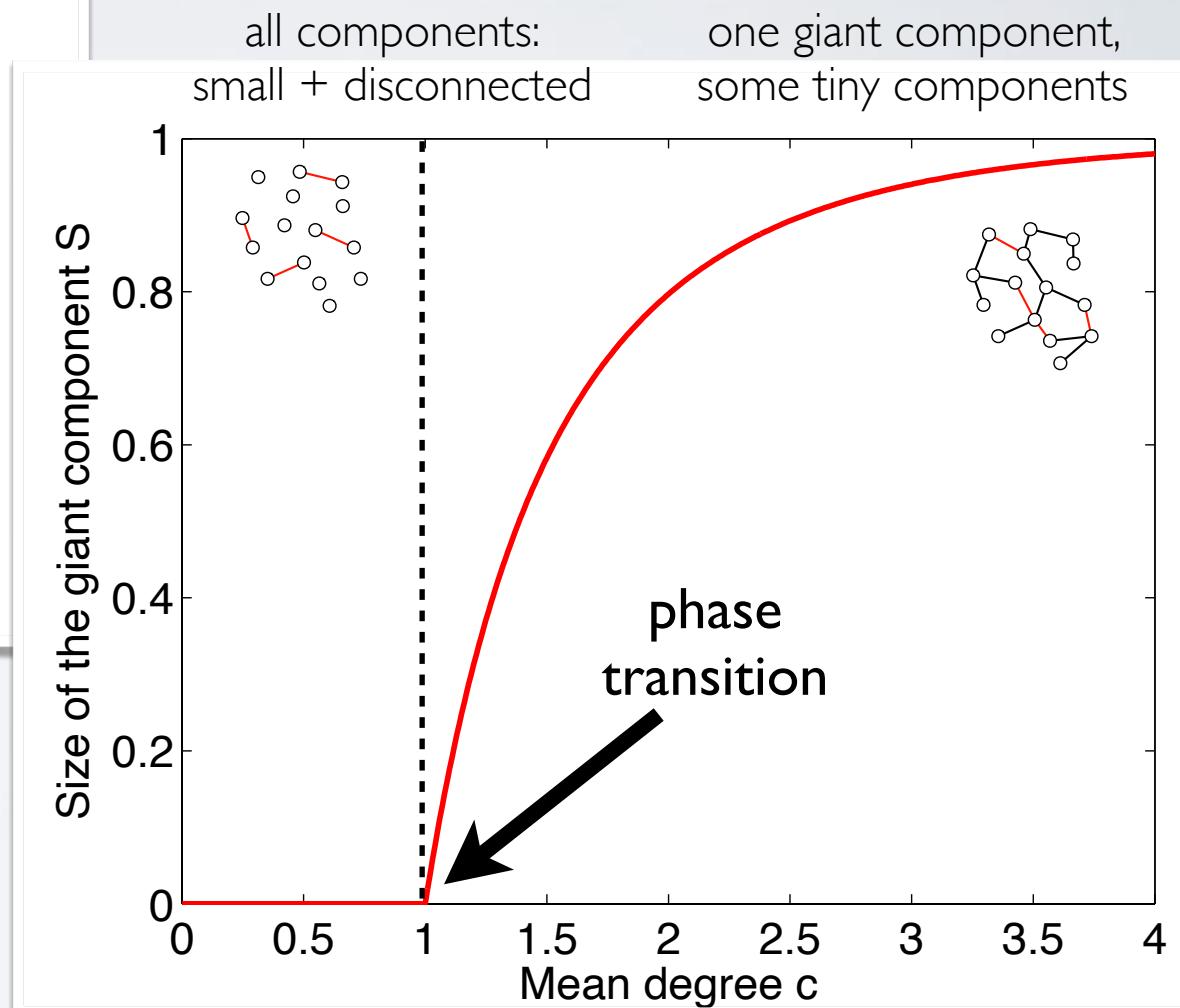
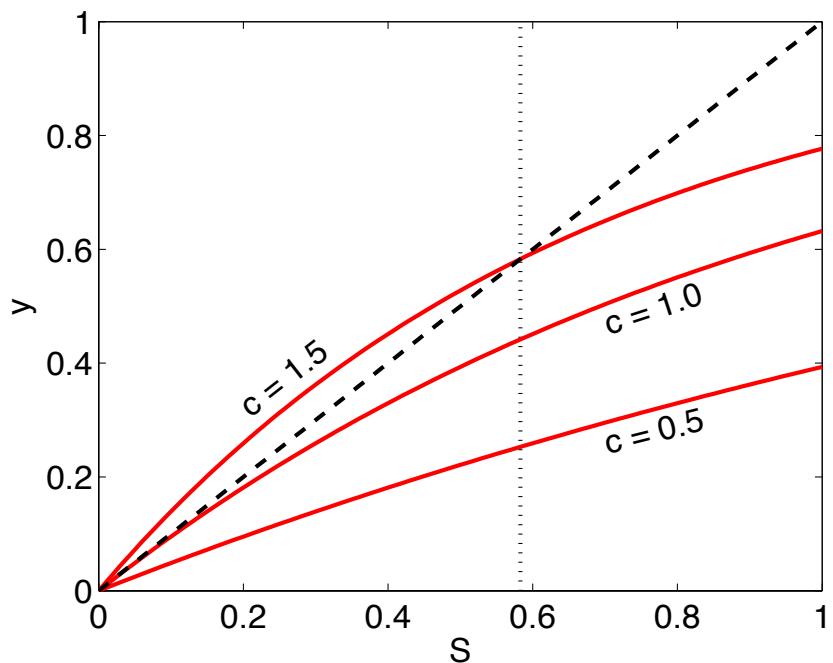
eliminating u for S yields the transcendental equation

$$S = 1 - e^{-cS}$$

[first given by Erdos and Renyi in 1959]

giant component

size of the giant component: $S = 1 - e^{-c} S$



citation networks

example of a **dynamic, growing network** model

example of a network **mechanistic** model

ample data

pleasing narcissistic qualities

long history of study

generally well understood

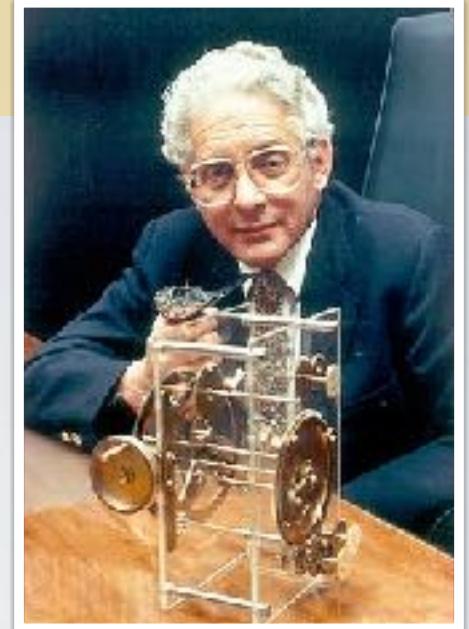
citation networks

Networks of Scientific Papers

The pattern of bibliographic references indicates the nature of the scientific research front.

1965

Derek J. de Solla Price



Price's model:

- papers are published continually [growing network]
- each paper has bibliography of length c [mean out degree]
- new papers cite previously published only [directed acyclic graph]
- attachment mechanism:

so long as $\Pr(c)$ has finite variance, the model is well defined

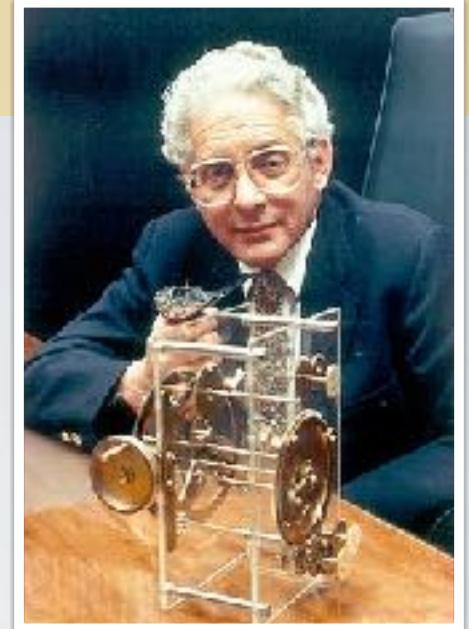
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- attachment mechanism:

$$p(j \text{ cites some paper } i) \propto k_i + a$$

preferential
attachment

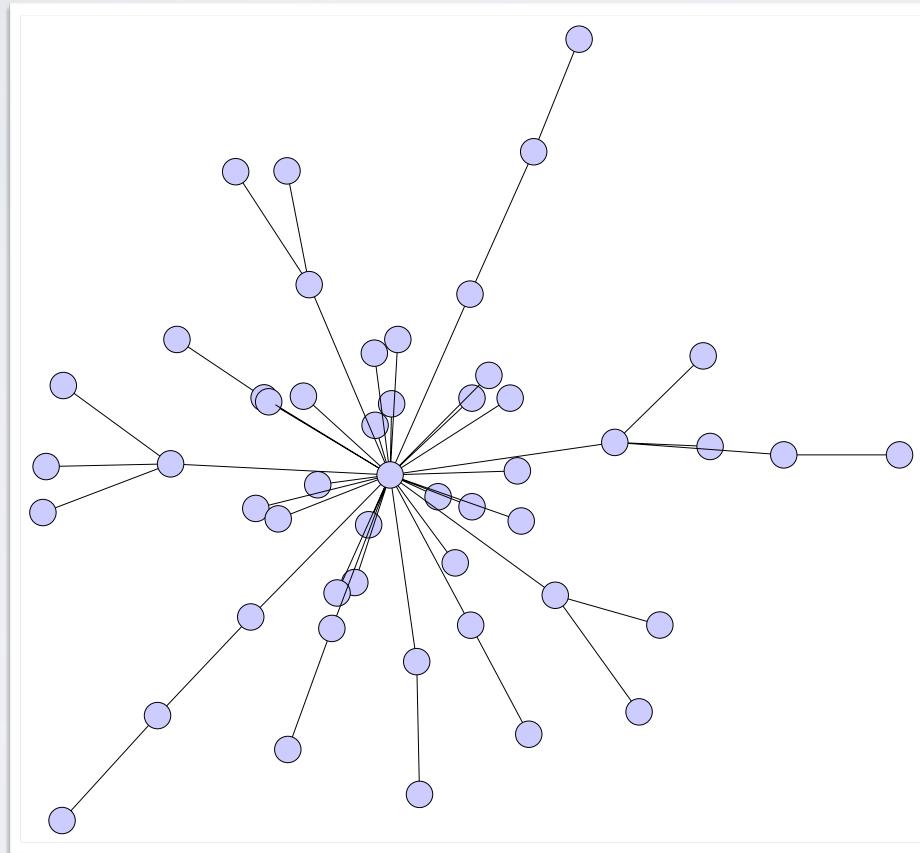
uniform
attachment

preferential attachment networks

$n = 50$

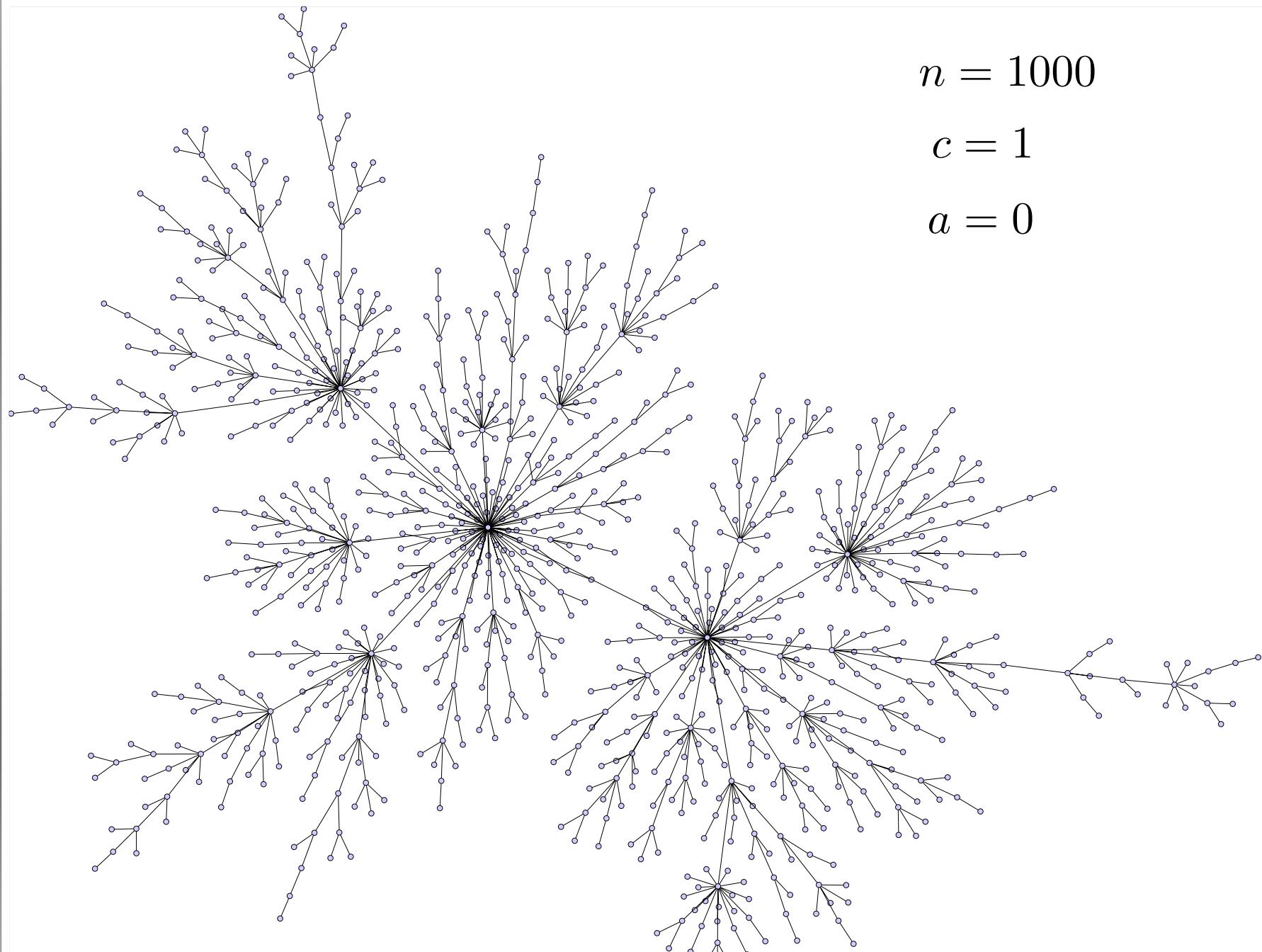
$c = 1$

$a = 0$



* this is a scale-free network, because the term "scale free" refers specifically to a graph with a power-law degree distribution (or tail), which this model produces

preferential attachment networks



* this is a scale-free network, because the term "scale free" refers specifically to a graph with a power-law degree distribution (or tail), which this model produces

degree distribution

exactly solvable in the limit [originally by Simon 1955]

$$p_k = \frac{B(k + a, \alpha)}{B(a, \alpha - 1)} \quad \alpha = 2 + a/c$$

degree distribution

exactly solvable in the limit

[originally by Simon 1955]

$$p_k = \frac{B(k+a, \alpha)}{B(a, \alpha-1)} \quad \alpha = 2 + a/c$$

recall that

$$B(a, b) = \Gamma(a)\Gamma(b)/\Gamma(a+b)$$

$$B(a, b) \sim a^{-b} \quad (\text{in the tail})$$

degree distribution

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$$p_k = \frac{B(k + a, \alpha)}{B(a, \alpha - 1)} \quad \alpha = 2 + a/c$$

recall that

$$B(a, b) = \Gamma(a)\Gamma(b)/\Gamma(a + b)$$

$$B(a, b) \sim a^{-b} \quad (\text{in the tail})$$

thus, distribution of citations

$$p_k \approx (k + a)^{-\alpha}$$

degree distribution

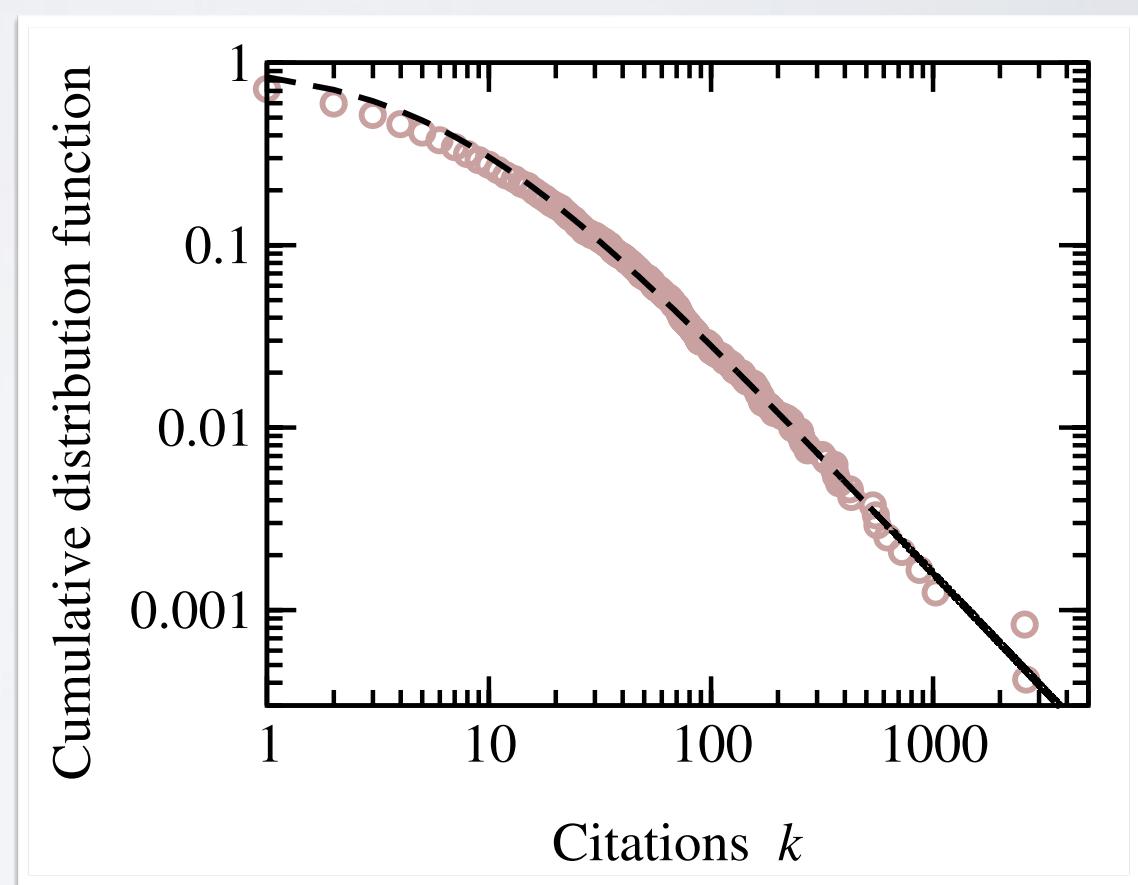
The first-mover advantage in scientific publication

M. E. J. NEWMAN^(a)

2009

$$p_k \approx (k + a)^{-\alpha}$$

- 2407 network science papers
- from 1998-2008
- fitted parameters
 $\alpha = 2.28$
 $a = 6.38$



the first-mover effect

The first-mover advantage in scientific publication

M. E. J. NEWMAN^(a)

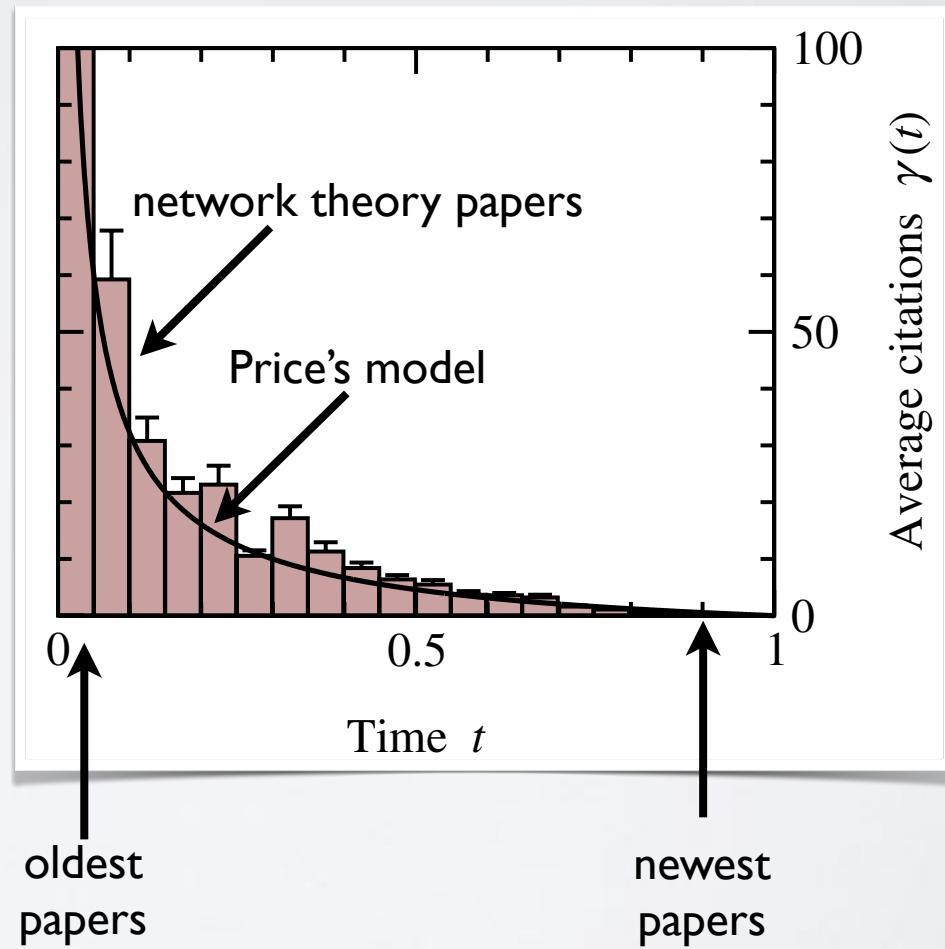
2009

- let t_i denote time that paper i was published
 - new papers only cite older papers
 - thus, first-mover effect: $k_i \propto 1/t_i$
-
- Price's model fully specified by α and a
 - idea:
 1. estimate them from total citation distribution
 2. derive predictions about citation counts vs. age of paper

the first-mover effect

average citations $\langle k \rangle$ vs. time of publication t

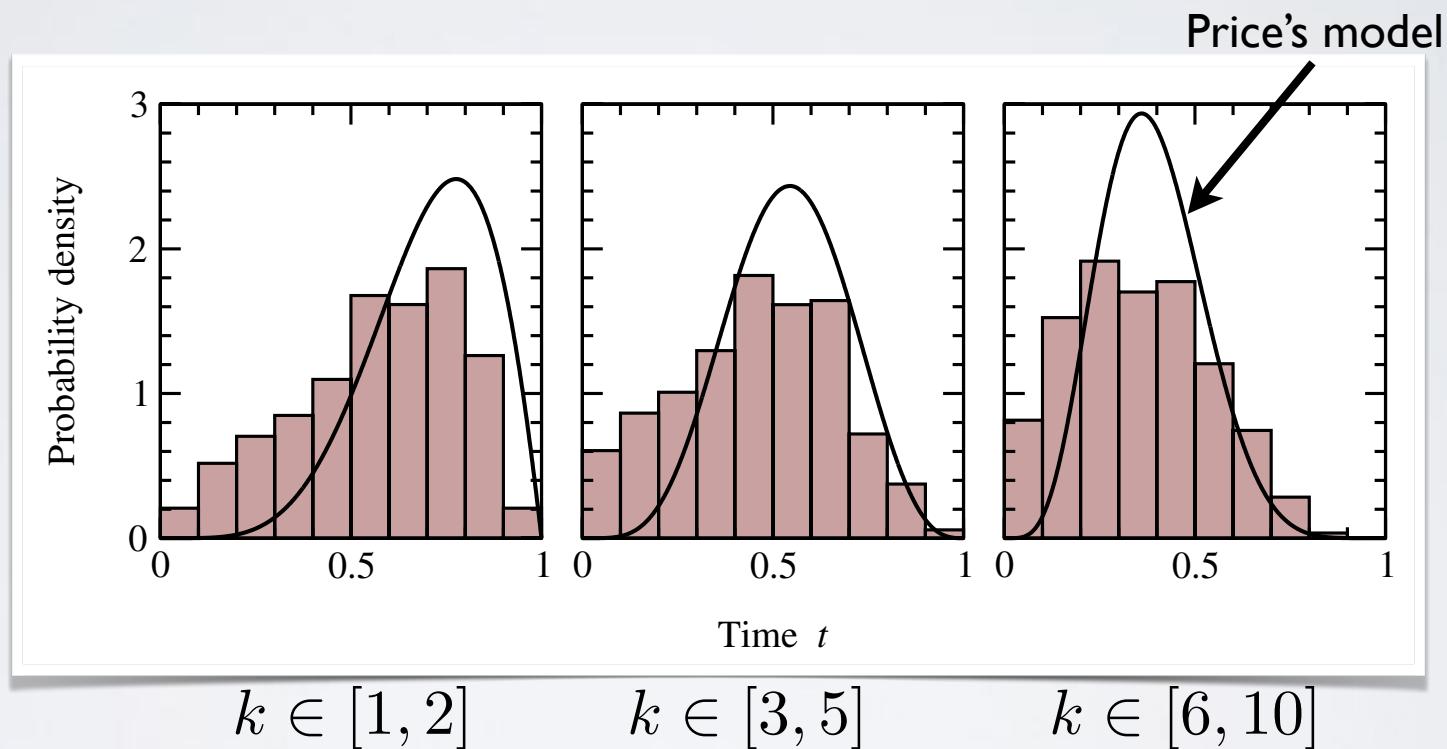
no free parameters



the first-mover effect

given k citations at time $t = 1$, probability of publication time t_i

no free parameters



checking the model



110 years of data (July 1893 - June 2003)

3.1 millions citations

330,000 papers with at least one citation

key question: is attachment function $\propto k_i$?

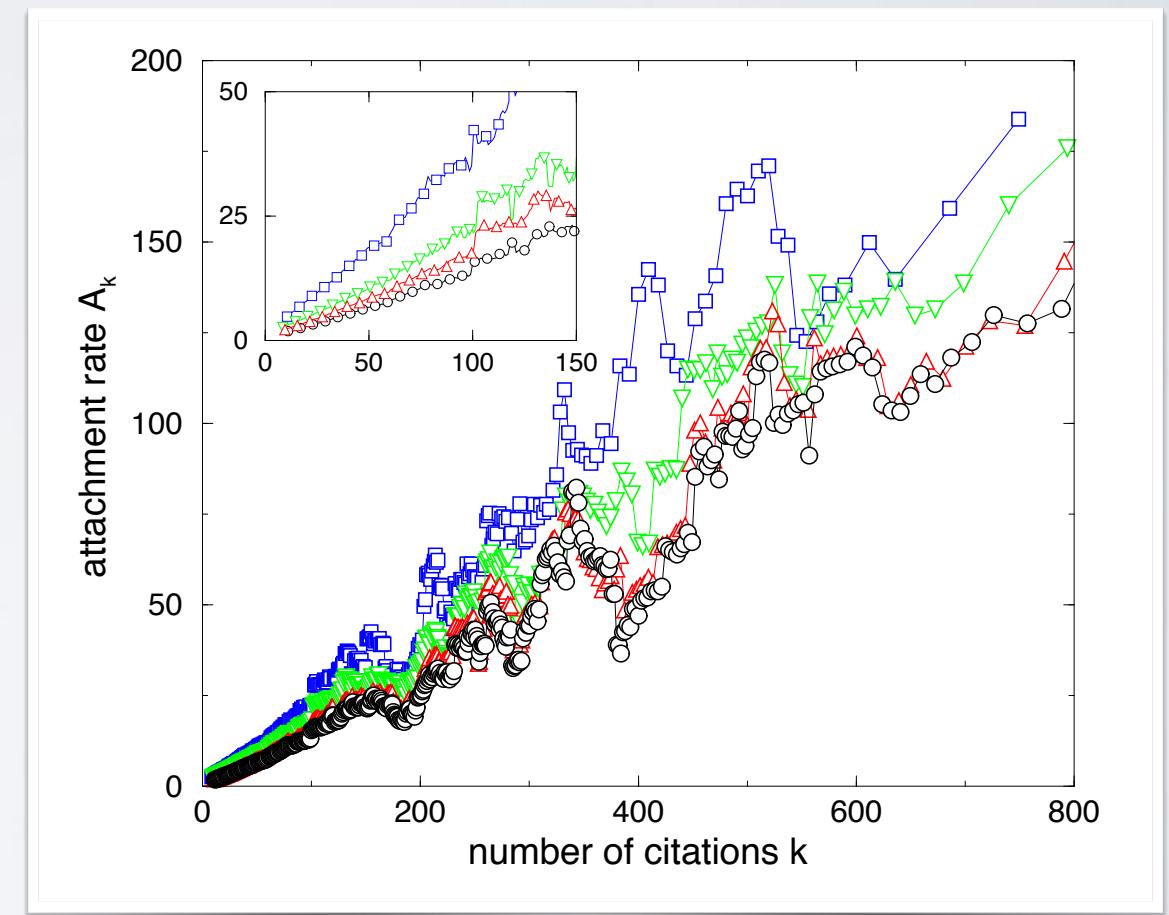
checking the model

key question: is attachment function $\propto k_i$?

pretty much.

caveat:

- ensemble only
(not individual papers)



citation networks

networks of scientific publications

summary of features

- Price's model: *preferential + uniform attachment*
 - excellent model of citation networks
 - also good model of WWW
 - a variation (duplication-mutation) good for gene networks
- not a great model of many other networks
 - especially social and spatial networks
 - ignores constraints (cost of edges)
- many additional mathematical, empirical results
 - see Redner's, Newman's, Fortunato's work