## Random Models

- Model $G(n, m)$ is a probability distribution $P(G)$ over all graphs with $n$ nodes and $m$ edges.
- Properties of model $=$ properties of ensemble
- Examples


- degTetal \{体 (nyder of gaph $n$

Q: what is the total number of graphs with no loops and multi-edges?
$2\binom{n}{2}$

## Random Models

- Model $G(n, p)$ - graphs with $n$ nodes and independent probability $p$ for placing an edge between two vertices (aka Erdös-Rényi model).
- Properties of model $=$ properties of ensemble where a particular graph $G$ with $m$ edges appears with probability

$$
P(G)=p^{m}(1-p)^{\binom{n}{2}-m}
$$

and probability of drawing a graph with $m$ edges from the ensemble is

$$
P(m)=\binom{\binom{n}{2}}{m} p^{m}(1-p)^{\binom{n}{2}-m} \text { and }\langle m\rangle=\sum_{m=0}^{\binom{n}{2}} m P(m)=\binom{n}{2} p
$$

- mean degree $\sum_{m=0}^{\binom{n}{2}} \frac{2 m}{n} P(m)=\frac{2}{n}\binom{n}{2} p=(n-1) p=c$
mean degree in a graph with exactly $m$ edges
- degree distribution
- node is connected to a particular $k$ others $q_{k}=p^{k}(1-p)^{n-1-k}$
- node is connected to exacly $k$ others $p_{k}=\binom{n-1}{k} q_{k}$
- mean degree $\sum_{m=0}^{\binom{n}{2}} \frac{2 m}{n_{<}} P(m)=\frac{2}{n}\binom{n}{2} p=(n-1) p=c$
mean degree in a graph with exactly $m$ edges
- degree distribution
- node is connected to a particular $k$ others $q_{k}=p^{k}(1-p)^{n-1-k}$
- node is connected to exacly $k$ others $p_{k}=\binom{n-1}{k} q_{k}$
- in large-scale networks $p=c /(n-1)$ can be very small, i.e.,

$$
\ln \left((1-p)^{n-1-k}\right)=(n-1-k) \ln (1-c /(n-1)) \approx-(n-1-k) \frac{c}{n-1} \approx-c
$$

Taylor series reminder: $\ln \left(1+\frac{1}{x}\right)=2\left(A+\frac{1}{3} A^{3}+\frac{1}{5} A^{5}+\ldots\right)$, where $A=\frac{1}{2 x+1}$ also if $\binom{n-1}{k}=\frac{(n-1)!}{(n-1-k)!k!} \approx \frac{(n-1)^{k}}{k!}$ then

$$
p_{k}=\frac{(n-1)^{k}}{k!} p^{k} e^{-c}=\frac{(n-1)^{k}}{k!}\left(\frac{c}{n-1}\right)^{k} e^{-c}=e^{-c} \frac{c^{k}}{k!}
$$

Poisson distribution in random models


In contrast to the degree distribution in random model, many real network degree distributions are different.


In contrast to the degree distribution in random model, many real network degree distributions are different.



- clustering coefficient $C=c /(n-1)=$ prob that any two nodes are neighbors

|  |  |  | clustering coefficient $C$ |  |
| :--- | ---: | ---: | :--- | :--- |
| network | $n$ | $z$ | measured | random graph |
| Internet (autonomous systems) $^{\mathrm{a}}$ | 6374 | 3.8 | 0.24 | 0.00060 |
| World-Wide Web (sites) $^{\mathrm{b}}$ | 153127 | 35.2 | 0.11 | 0.00023 |
| power grid $^{\mathrm{c}}$ | 4941 | 2.7 | 0.080 | 0.00054 |
| biology collaborations $^{\mathrm{d}}$ | 1520251 | 15.5 | 0.081 | 0.000010 |
| mathematics collaborations $^{\mathrm{e}}$ | 253339 | 3.9 | 0.15 | 0.000015 |
| film actor collaborations $^{\mathrm{f}}$ | 449913 | 113.4 | 0.20 | 0.00025 |
| company directors $^{\mathrm{f}}$ | 7673 | 14.4 | 0.59 | 0.0019 |
| word co-occurrence $^{\mathrm{g}}$ | 460902 | 70.1 | 0.44 | 0.00015 |
| neural network $^{\mathrm{c}}$ | 282 | 14.0 | 0.28 | 0.049 |
| metabolic network $^{\mathrm{h}}$ | 315 | 28.3 | 0.59 | 0.090 |
| food web $^{\mathrm{i}}$ | 134 | 8.7 | 0.22 | 0.065 |
|  | Newman, "Random graphs as models of networks" |  |  |  |

- giant component in $G(n, p)$

Giant component is a network component whose size grows in proportion to $n$.

Q : When $\mathrm{p}=0$ then $|\mathrm{gc}|=1$; when $\mathrm{p}=1$ then $|\mathrm{gc}|=\mathrm{n}$. What is the difference between them?


Co-authorship network
its largest connected component

- giant component in $G(n, p)$

Giant component is a network component whose size grows in proportion to $n$. $u=$ avg fraction of vertices that do not belong to the giant component.
$Q$ : When $p=0$ then $|g c|=1$; when $p=1$ then $|g c|=n$. Is this transition smooth? Is there a point of transition?
$i$ does not belong to gc if for every node $j$ either
a) there is no edge $i j$, prob $=1-p$
b) there is edge $i j$ but $j$ is not in giant component, prob = pu
$\operatorname{Pr}[i$ does not belong to gc via $j]=1-p+p u$, i.e., total probability of not being connected to gc via any of $n-1$ other vertices is

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

$\ln u \stackrel{n \rightarrow \infty}{\approx}_{\approx}-(n-1) \frac{c}{n-1}(1-u)=-c(1-u) \Rightarrow u=e^{-c(1-u)} \Rightarrow S=1-e^{-c S}$
vertices in giant component

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




Figure 12.1: Graphical solution for the size of the giant component. (a) The three curves in the left panel show $y=1-\mathrm{e}^{-c s}$ for values of $c$ as marked, the diagonal dashed line shows $y=S$, and the intersection gives the solution to Eq. (12.15), $S=1-\mathrm{e}^{-c S}$. For the bottom curve there is only one intersection, at $S=0$, so there is no giant component, while for the top curve there is a solution at $S=0.583 \ldots$ (vertical dashed line). The middle curve is precisely at the threshold between the regime where a non-trivial solution for $S$ exists and the regime where there is only the trivial solution $S=0$. (b) The resulting solution for the size of the giant component as a function of $c$.

|  | Medline | Physics E-print Archive |  |  |  | SPIRES | NCSTRL |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | complete | astro-ph | cond-mat | hep-th |  |  |
| total papers | 2163923 | 98502 | 22029 | 22016 | 19085 | 66652 | 13169 |
| total authors | 1520251 | 52909 | 16706 | 16726 | 8361 | 56627 | 11994 |
| first initial only | 1090584 | 45685 | 14303 | 15451 | 7676 | 47445 | 10998 |
| mean papers per author | 6.4(6) | 5.1(2) | 4.8(2) | 3.65(7) | 4.8(1) | 11.6(5) | 2.55(5) |
| mean authors per paper | 3.754(2) | $2.530(7)$ | 3.35 (2) | 2.66(1) | 1.99(1) | 8.96(18) | 2.22(1) |
| collaborators per author | 18.1(1.3) | 9.7(2) | 15.1(3) | 5.86(9) | 3.87 (5) | 173(6) | 3.59(5) |
| size of giant component | 1395693 | 44337 | 14845 | 13861 | 5835 | 49002 | 6396 |
| first initial only | 1019418 | 39709 | 12874 | 13324 | 5593 | 43089 | 6706 |
| as a percentage | 92.6(4)\% | 85.4(8)\% | 89.4(3) | 84.6(8)\% | 71.4(8)\% | 88.7(1.1)\% | 57.2(1.9)\% |
| 2nd largest component | 49 | 18 | 19 | 16 | 24 | 69 | 42 |
| clustering coefficient $C$ | 0.066(7) | 0.43(1) | 0.414(6) | 0.348(6) | 0.327(2) | 0.726(8) | 0.496(6) |
| mean distance | 4.6(2) | 5.9(2) | 4.66(7) | 6.4(1) | $6.91(6)$ | 4.0(1) | 9.7(4) |
| maximum distance | 24 | 20 | 14 | 18 | 19 | 19 | 31 |

Table 1: Summary of results of the analysis of seven scientific collaboration networks. Numbers in parentheses give an estimate of the error on the least significant figures.

|  | Network | Type | $\stackrel{1}{ }$ | 74 | ${ }_{6}$ | $S$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\stackrel{\bar{\pi}}{\mathscr{C}}$ | Film actors | Undirected | 449913 | 25516482 | 113.43 | 0.980 |
|  | Company directors | Undirected | 7673 | 55392 | 11.41 | 0.876 |
|  | Math coauthorghip | Undirected | 253339 | 4.96489 | 3.92 | 0.822 |
|  | Physics coauthorship | Undirected | 52909 | 245300 | 9.27 | 0.838 |
|  | Biology coauthorship | Undirected | 1520.251 | 11853064 | 15.53 | 0.518 |
|  | Telephone calt graph | Undirected | 4700000 | 80000000 | 3.46 |  |
|  | Emaiìmessages | Directed | 59812 | 86500 | 1.44 | 0.952 |
|  | Email address books | Directed | 15881 | 57029 | 3.38 | 0.590 |
|  | Stucent dating | Crdirected | 573 | 477 | 1.66 | 0.503 |
|  | Sexual contacts | Undirected | 2810 |  |  |  |
| $\begin{aligned} & \text { 煲 } \\ & \text { 总 } \\ & \text { C } \end{aligned}$ | WWW ad．odu | Directed． | ＇269504 | 1497135 | 5.55 | 1.000 |
|  | WWW Altarista | Directed | 203549046 | 1466000000 | 7.20 | 0.914 |
|  | Citation network | Directed | 783339 | 67：6198 | 8.57 |  |
|  | Roget＇s Thesaurus | Directed | 1022 | 5103 | 1.99 | 0.977 |
|  | Woric co－occumence | Lrdirected | 460902 | 16100000 | 66.96 | 1.000 |
|  | Internet | Undirected | 10697 | 31992 | 5.98 | 1.000 |
|  | Power gric | Undirected | 4941 | 6594 | 2.67 | 1.000 |
|  | Train roctes | Undirected | 587 | 19603 | 66.79 | 1.000 |
|  | Soitware packages | Directed | 1439 ． | 1723 | 1.20 | 0.998 |
|  | Software classes | Directed | 1376 | 2213 | 1.61 | 1.000 |
|  | Electronic circuits | Undirected | 24097 | 32248 | 4.34 | 1000 |
|  | Peer－to－peer network | Undirected | 880 | 1296 | 1.47 | 0.805 |
| $\begin{aligned} & \text { 哥 } \\ & \text { 药 } \\ & 0 \\ & 0 \\ & 0 \end{aligned}$ | Metabolic network | Undirected | 765 | 3686 | 9.64 | 0.996 |
|  | Protein interactions | Undirecteá |  | 2240 | 212 | 0.689 |
|  | Marine food web | Directed | 134 | 598 | 4.46 | 1.000 |
|  | Freshwater food web | Directed | 92 | 997 | 10.84 | 1.000 |
|  | Neural network | Directed | 307 | 2.359 | 7.68 | 0.967 |

Two giant components in $G(n, p)$ ?

- Generate $G$ with $p=c /(n-1)$
- Suppose that after adding edges with prob $p$, we have 2 giant components that cover fractions of nodes $S_{1}$, and $S_{2}$
- $S_{1}$ and $S_{2}$ remain separate with probability

$$
\begin{aligned}
& \qquad q=(1-p)^{S_{1} S_{2} n^{2}}=\left(1-\frac{c}{n-1}\right)^{S_{1} S_{2} n^{2}} \\
& \Rightarrow \ln q=S_{1} S_{2} \lim _{n \rightarrow \infty}\left(n^{2} \ln \left(1-\frac{c}{n-1}\right)\right)=S_{1} S_{2}\left(-c(n+1)+\frac{1}{2} c^{2}\right) \\
& q=q_{0} e^{-c S_{1} S_{2} n}, \\
& \text { where } q_{0} \text { is a constant, i.e., } q \xrightarrow{n \rightarrow \infty} 0 \\
& \begin{array}{l}
\text { Conclusion: In the limit of large } n \text {, the probability of existence } \\
\text { of two separate giant components goes to zero. }
\end{array}
\end{aligned}
$$

- Alright ... we have only one giant component. What about the sizes of small components?
$\pi_{s}$ is the probability that randomly chosen node belongs to a small component of size $s$.
- We cannot normalize $\pi_{s}$ to unity because some nodes may belong to the giant component, ie.,

$$
\sum_{s=0}^{\infty} \pi_{s}=1-S . \varlimsup_{\text {giant component }}^{\substack{\text { fraction of nodes in } \\ \text { gin }}}
$$

- Observation: small components are likely to be trees. Consider a small tree component of $s$ nodes. The total number of places we can add an extra edge to is $\binom{s}{2}-(s-1) \longleftarrow$ edges in tree

Average total number of added edges $\frac{1}{2}(s-1)(s-2) \cdot \frac{c}{n-1} \xrightarrow{n \rightarrow \infty} 0$
the component is still tree

Calculation of $\pi_{s}$ (the probability that randomly chosen node belongs to a small component of size $s$ ).

- Consider node $i$ in a small (tree) component
- ... and modified network with deleted $i$.

In the modified network, prob $p$ is the same and in the limit of $n$ the changes are negligible. Sizes of gc and sc will be indistinguishable for same $p$.

- Suppose $d(i)=k$ and $\operatorname{Pr}\left[n_{1} \in\right.$ sc of size $\left.s_{1}\right]=\pi_{s_{1}}$ and


$$
\operatorname{Pr}\left[\forall j \in N(i) n_{j} \in \text { sc of size } s_{j}\right]=\Pi_{j=1}^{k} \pi_{s_{j}}
$$

Since $\sum_{j \in N(i)} s_{j} s-1$ we have
Kronecker delta

$$
\begin{aligned}
& p_{k}=e^{-c} \frac{c^{k}}{k!} \operatorname{Pr}[s \mid k]=\sum_{s_{1}=1}^{\infty} \cdots \sum_{s_{k}=1}^{\infty}\left(\Pi_{j=1}^{k} \pi_{s_{j}}\right) \delta\left(s-1, \sum_{j} s_{j}\right) \\
& \pi_{s}=\sum_{k=0}^{\infty} p_{k} \operatorname{Pr}[s \mid k]=e^{-c} \sum_{k=0}^{\infty} \frac{c^{k}}{k!} \sum_{s_{1}=1}^{\infty} \cdots \sum_{s_{k}=1}^{\infty}\left(\Pi_{j=1}^{k} \pi_{s_{j}}\right) \delta\left(s-1, \sum_{j} s_{j}\right)
\end{aligned}
$$

$$
\pi_{s}=\sum_{k=0}^{\infty} p_{k} \operatorname{Pr}[s \mid k]=e^{-c} \sum_{k=0}^{\infty} \frac{c^{k}}{k!} \sum_{s_{1}=1}^{\infty} \cdots \sum_{s_{k}=1}^{\infty}\left(\prod_{j=1}^{k} \pi_{s_{j}}\right) \delta\left(s-1, \sum_{j} s_{j}\right)
$$

One way to evaluate $\pi_{s}$ is by using generating function

$$
h(z)=\sum_{s=1}^{\infty} \pi_{s} z^{s} \Rightarrow\langle s\rangle=\frac{\sum_{s} s \pi_{s}}{\sum_{s} \pi_{s}}=h^{\prime}(1) /(1-S)=1 /(1-c+c S) .
$$

see Newman's book, pp 412-413

## Average size of the small components in a random model does not grow with the number of vertices.

## Average component size

$$
R=\frac{2}{2-c+c S}
$$



Figure 12.4: Average size of the small components in a random graph. The upper curve shows the average size $\langle s\rangle$ of the component to which a randomly chosen vertex belongs, calculated from Eq. (12.34). The lower curve shows the overall average size $R$ of a component, calculated from Eq. (12.40). The dotted vertical line marks the point $c=1$ at which the giant component appears. Note that, as discussed in the text, the upper curve diverges at this point but the lower one does not.

## Distribution of component sizes


#### Abstract

Figure 12.5: Sizes of small components in the random graph. This plot shows the probability $\pi_{s}$ that a randomly chosen vertex belongs to a small component of size $s$ in a Poisson random graph with $c=0.75$ (top), which is in the regime where there is no giant component, and $c=1.5$ (bottom), where there is a giant component.

Component size $s$


- path lengths
- Intuition: avg number of nodes $s$ steps away from random $i$ is $c^{s}$. We reach all vertices when $c^{s} \approx n$, i.e., $s \approx \ln n / \ln c$.
- Problem: this argument doesn't work when $s$ is large.
- Consider two starting vertices $i$ and $j$ with their $s$ - and $t$-distance neighborhoods, respectively, when $s, t$ are small

1. if edge exists between surfaces then one can show that there are edges between larger surfaces


$$
\begin{aligned}
& \Longrightarrow \operatorname{Pr}\left[d_{i j}>s+t+1\right] \approx \text { prob } \nexists \text { edge between two surfaces } \\
& c^{s} \text {, and } c^{t} \text { when } t \text { is small }
\end{aligned}
$$

2. There are on avg $c^{s} \times c^{t}$ pairs of nodes, s.t. one lies on each surface and each pair is connected with prob $p=c /(n-1)$
i.e., $\operatorname{Pr}\left[d_{i j}>s+t+1\right]=(1-p)^{c^{s+t}}=(1-c / n)^{c^{l-1}}$ or $\ln \operatorname{Pr}\left[d_{i j}>l\right]=$ $c^{l-1} \ln (1-c / n) \approx-c^{l} / n$

$$
l=s+t+1
$$

| Networks | \# of nodes | Diameter |  |  |  |  | Modularity |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Observed | Expected | \% difference | $\begin{aligned} & z \\ & \text { score }{ }^{14} \end{aligned}$ | P-value | Observed | Expected | \% difference | $\begin{aligned} & z \\ & \text { score }{ }^{14} \end{aligned}$ | $P_{\text {value }}$ |
| Characters in "Les Miserables"1 | 77 | 2.64 | 2.50 | 5.6 | 3.58 | 0.0003 | 0.56 | 0.29 | 93.4 | 30.12 | $<10^{-4}$ |
| Words in "David Copperfield"2 | 112 | 2.54 | 2.48 | 2.3 | 1.81 | 0.0703 | 0.31 | 0.29 | 4.8 | 1.67 | 0.0949 |
| Dolphins ${ }^{3}$ | 62 | 3.36 | 2.70 | 24.3 | 14.40 | $<10^{-4}$ | 0.53 | 0.37 | 40.8 | 11.59 | $<10^{-4}$ |
| Political blogs ${ }^{4}$ | 1224 | 2.74 | 2.59 | 5.7 | 23.5 | $<10^{-4}$ | 0.43 | 0.14 | 206.9 | 189.27 | $<10^{-4}$ |
| Co-authorship ${ }^{5}$ | 7610 | 7.03 | 5.42 | 29.6 | 64.70 | $<10^{-4}$ | 0.81 | 0.49 | 64.9 | 12.50 | $<10^{-4}$ |
| Football ${ }^{6}$ | 115 | 2.51 | 2.23 | 12.5 | 54.30 | $<10^{-4}$ | 0.60 | 0.28 | 119.2 | 44.68 | $<10^{-4}$ |
| Power ${ }^{\text {² }}$ | 4941 | 18.99 | 8.32 | 128.3 | 14.30 | $<10^{-4}$ | 0.93 | 0.73 | 28.5 | 105.10 | $<10^{-4}$ |
| Airline ${ }^{\text {a }}$ | 810 | 3.06 | 2.61 | 17.4 | 3.53 | 0.0004 | 0.31 | 0.13 | 130.0 | 114.70 | $<10^{-4}$ |
| Electronic circuits* | 512 | 6.86 | 5.64 | 21.6 | 12.40 | $<10^{-4}$ | 0.81 | 0.63 | 28.6 | 35.96 | $<10^{-4}$ |
| Protein-protein interaction ${ }^{10}$ | 1870 | 6.81 | 5.78 | 17.8 | 9.19 | $<10^{-4}$ | 0.81 | 0.72 | 13.2 | 18.23 | $<10^{-4}$ |
| Neural ${ }^{11}$ | 297 | 2.46 | 2.35 | 4.5 | 3.38 | 0.0007 | 0.40 | 0.22 | 80.0 | 51.26 | $<10^{-4}$ |
| Transcriptional regulatory ${ }^{12}$ | 3459 | 3.72 | 3.39 | 9.7 | 3.60 | 0.0003 | 0.60 | 0.47 | 29.5 | 58.29 | $<10^{-4}$ |
| Metabolic ${ }^{13}$ | 563 | 8.78 | 6.54 | 34.3 | 18.67 | $<10^{-4}$ | 0.84 | 0.73 | 14.5 | 14.72 | $<10^{-4}$ |

"The network of coappearances of characters in Victor Hugo's novel "Les Miserables". Nodes represent characters and edges connect any pair of characters that appear in the same chapter,
${ }^{2}$ The network of common adjective and noun adjacencies for the novel "David Copperfield" by Charles Dickens. Nodes represent the most commonly occurring adjectives and nouns in the book.
'The network of frequent associations between 62 dolphins in a community living off Doubtful Sound, New Zealand.
${ }^{4}$ The network of political blogs. Nodes represent blogs and edges are the links between blogs.
The network of scientists posting preprints on the high-energy theory archive at wwwaroiv.org. 1995-1999. Nodes are authors and edges connect coauthors.
${ }^{6}$ The network of American football games between Division IA colleges during regular season Fall 2000. Nodes are teams and edges connect teams that contest in a game.
The network of the Western States Power Grid of the United States. Nodes are power plants, stations and households, and edges are powerlines.
"The network of scheduled air line connections in United States, 2005. Nodes are airports and edges are scheduled direct flights.
"Electronic circuits. Nodes are electronic elements and edges are electronic connections.
${ }^{10}$ The protein-protein interaction network of the budding yeast S . cerevisiae. Nodes are proteins and edges connect proteins that interact with each other.
"The neural network for the worm C. elegans. Nodes are neurons and edges link neurons that connect.
${ }^{12}$ The transcriptional regulatory network of the budding yeast $S$. cerevisioe. Nodes are genes and edges connect genes that regulate one another.
${ }^{13}$ The metabolic network of the bacterium $E$. coll. Nodes are metabolites and edges connect metabolites that can be converted by a blochemical reaction.
${ }^{14} Z$-score, number of standard deviations by which the observation deviates from the expectation.
docic.1371/journal.pone.0005686.t001

## Generating Functions and Degree Distributions

The generating function (gf) for the probability distribution $p_{k}$ is the polynomial

$$
g(z)=\sum_{k=0}^{\infty} p_{k} z^{k}
$$

If we know gf for $p_{k}$ then we can recover the values of $p_{k}$ by differentiating

$$
p_{k}=\left.\frac{1}{k!} \frac{d^{k} g}{d z^{k}}\right|_{z=0}
$$

Example: $k=0,1,2$ with the respective $p_{k}=\frac{1}{2}, \frac{7}{16}, \frac{1}{16}$ for all $k$ then

$$
g(z)=\frac{1}{2}+\frac{7}{16} z+\frac{1}{16} z^{2}
$$

Example: $k$ follows Poisson distribution, i.e., $p_{k}=e^{-c \frac{c^{k}}{k!}}$

$$
g(z)=e^{-c} \sum_{k=0}^{\infty} \frac{c^{k}}{k!} z^{k}=e^{c(z-1)}
$$

Power-law distributions $p_{k}=C k^{-\alpha}, \alpha>0, k>0$
Reminder: $C$ is calculated from normalization condition, i.e., $C=1 / \zeta(\alpha)$

$$
p_{k}=\left\{\begin{array}{ll}
0 & k=0 \\
k^{-\alpha} / \zeta(\alpha) & k>0
\end{array} \Longrightarrow g(z)=\frac{1}{\zeta(\alpha)} \sum_{k=1}^{\infty} k^{-\alpha} z^{k}=\frac{L i_{\alpha}(z)^{~}}{\zeta(\alpha)}\right.
$$

Since we are interested in differentiating $g(z)$ note that

$$
\frac{\partial L i_{\alpha}(z)}{\partial z}=\frac{L i_{\alpha-1}(z)}{z}
$$

Some properties of $g(z)$

- $g(1)=1$
- $\langle k\rangle=g^{\prime}(1),\left\langle k^{2}\right\rangle=\left[\left(z \frac{d}{d z}\right)^{2} g(z)\right]_{z=1}, \ldots,\left\langle k^{m}\right\rangle=\left[\left(z \frac{d}{d z}\right)^{m} g(z)\right]_{z=1}$
- Choose $m$ integers $k_{i}$ from $p_{k} \Rightarrow \operatorname{Pr}\left[\right.$ chosing particular set of values $\left.\left\{k_{i}\right\}\right]=$ $\prod_{i} p_{k_{i}}$


## Random Graphs and Configuration Model

Degrees: 1, 1, 2, 2, 3, 3


1. Add $n$ nodes
2. Add initial $d(i)$ stubs to each $i$
3. Connect stubs iteratively

Problems? Total degree is even; Can create self-loops, multi-edges

## Configuration Model

Multi-edges: Probability of adding an edge between $i$ and $j$ with degrees $k_{i}$, and $k_{j}$ is

$$
p_{i j}=\frac{k_{i} k_{j}}{2 m-1}
$$

Probability of second edge is $\left(k_{i}-1\right)\left(k_{j}-1\right) / 2 m$
Expected number of multiedges in conf model

$$
\frac{1}{2(2 m)^{2}} \sum_{i j} k_{i} k_{j}\left(k_{i}-1\right)\left(k_{j}-1\right)=\frac{1}{2\langle k\rangle^{2} n^{2}} \sum_{i} k_{i}\left(k_{i}-1\right) \sum_{j} k_{j}\left(k_{j}-1\right)=\frac{1}{2}\left[\frac{\left\langle k^{2}\right\rangle-\langle k\rangle}{\langle k\rangle}\right]^{2}
$$

Similar result for self-edges

$$
\sum_{i} p_{i i}=\sum_{i} \frac{k_{i}\left(k_{i}-1\right)}{4 m}=\frac{\left\langle k^{2}\right\rangle-\langle k\rangle}{2\langle k\rangle}
$$

## Conclusion? Expected number of multi-edges remains constant as network grows.

Expected number of common neighbors

$$
n_{i j}=\sum_{l} \frac{k_{i} k_{l}}{2 m} \frac{k_{j}\left(k_{l}-1\right)}{2 m}=\frac{k_{i} k_{j}}{2 m} \frac{\sum_{l} k_{l}\left(k_{l}-1\right)}{n\langle k\rangle}=p_{i j} \frac{\left\langle k^{2}\right\rangle-\langle k\rangle}{\langle k\rangle}
$$

$i$ is connected to $/$
$j$ is connected to / given il

## Random graphs with given expected degree

$\forall i \in V$ define parameter $c_{i}$. Then edge probability

$$
p_{i j}=\left\{\begin{array}{ll}
c_{i} c_{j} / 2 m & i \neq j \\
c_{i}^{2} / 4 m & i=j
\end{array}, \text { where } \sum_{i} c_{i}=2 m\right.
$$

average number of edges in network

$$
\sum_{i \leq j} p_{i j}=\sum_{i<j} \frac{c_{i} c_{j}}{2 m}+\sum_{i} \frac{c_{i}^{2}}{4 m}=m
$$

average degree

$$
\left\langle k_{i}\right\rangle=2 p_{i i}+\sum_{j \neq i} p_{i j}=\frac{c_{i}^{2}}{2 m}+\sum_{j \neq i} \frac{c_{i} c_{j}}{2 m}=\sum_{j} \frac{c_{i} c_{j}}{2 m}=c_{i}
$$

## More properties of random model

Excess degree distribution is the probability distribution, for a vertex reached by following an edge, of the number of other edges attached to that vertex.

$$
q_{k}=\frac{(k+1) p_{k+1}}{\langle k\rangle}
$$

Two academic collaboration networks, in which scientists are connected together by edges if they have coauthored scientific papers, and a snapshot of the structure of the Internet at the autonomous system level.

| Network | $n$ | Average <br> degree | Average <br> neighbor degree | $\frac{\left\langle k^{2}\right\rangle}{\langle k\rangle}$ |
| :--- | ---: | :---: | :---: | ---: |
| Biologists | 1520252 | 15.5 | 68.4 | 130.2 |
| Mathematicians | 253339 | 3.9 | 9.5 | 13.2 |
| Internet | 22963 | 4.2 | 224.3 | 261.5 |

According to these results a biologist's collaborators have, on average, more than four times as many collaborators as they do themselves. On the Internet, a node's neighbors have more than 50 times the average degree! Note that in each of the cases in the table the configuration model value of $\left\langle k^{2}\right\rangle /\langle k\rangle$ overestimates the real average neighbor degree.
M. Newman "Networks"

## POLL

## CHOICE FOR PRESIDENT IF VOTING TODAY

## BIDEN-HARRIS TRUMP-PENCE



## POLL

WHO DO YOU THINK YOUR NEIGHBORS ARE SUPPORTING FOR PRESIDENT?

## BIDEN <br> TRUMP <br> DEPENDS UNSURE

NOW
\%
\%
\%
$\%$

## More properties of random model

Excess degree distribution is the probability distribution, for a vertex reached by following an edge, of the number of other edges attached to that vertex.

$$
q_{k}=\frac{(k+1) p_{k+1}}{\langle k\rangle}
$$

Using the excess degree distribution it is easy to compute the clustering coefficient for configuration model

$$
C=\sum_{k_{i}, k_{j}=0}^{\infty} q_{k_{i}} q_{k_{j}} \frac{k_{i} k_{j}}{2 m}=\frac{1}{2 m}\left(\sum_{k=0}^{\infty} k q_{k}\right)^{2}=\cdots=\frac{1}{n} \frac{\left(\langle k\rangle^{2}-\langle k\rangle\right)^{2}}{\langle k\rangle^{3}}
$$

## Generating Functions and Degree Distributions

For degree and excess degree distributions we define generating functions

$$
g_{0}(z)=\sum_{k=0}^{\infty} p_{k} z^{k} \text { and } g_{1}(z)=\sum_{k=0}^{\infty} q_{k} z^{k}, \text { respectively }
$$

They are not independent

$$
g_{1}(z)=\frac{1}{\langle k\rangle} \sum_{k=0}^{\infty}(k+1) p_{k+1} z^{k}=\frac{1}{\langle k\rangle} \sum_{k=0}^{\infty} k p_{k} z^{k-1}=\frac{1}{\langle k\rangle} \frac{\mathrm{d} g_{0}}{\mathrm{~d} z}=\frac{g_{0}^{\prime}(z)}{g_{0}^{\prime}(1)}
$$

Example (Poisson): $p_{k}=e^{-c \frac{c^{k}}{k!}} \Rightarrow g_{0}(z)=e^{c(z-1)}$ and $g_{1}(z)=e^{c(z-1)}$
Example (power-law): $p_{k}=C k^{-\alpha} \Rightarrow g_{0}(z)=\frac{L i_{\alpha}(z)}{\zeta(\alpha)}$. Thus,

$$
g_{1}(z)=\frac{L i_{\alpha-1}(z)}{z L i_{\alpha-1}(1)}=\frac{L i_{\alpha-1}(z)}{z \zeta(\alpha-1)}
$$

Polylogarithm function

$$
L i_{s}(z)=\sum_{k=1}^{\infty} \frac{z^{k}}{k^{s}}=z+\frac{z^{2}}{2^{s}}+\frac{z^{3}}{3^{s}}+\ldots
$$

## Number of second neighbors of a vertex

Probability that $i$ has exactly $k$ second neighbors


Conclusion: Once we know generating functions of $g_{0}$ and $g_{1}$ the generating function of second neighbor distribution is straightforward to calculate. Moreover, this can be extended to

$$
g^{(3)}(z)=\sum_{k=0}^{\infty} \sum_{m=0}^{\infty} p_{m}^{(2)} P^{(3)}(k \mid m) z^{k}=\sum_{n=0}^{\infty} p_{m}^{(2)}\left(g_{1}(z)\right)^{m}=g^{(2)}\left(g_{1}(z)\right)=g_{0}\left(g_{1}\left(g_{1}(z)\right)\right)
$$

$$
\Longrightarrow g^{(d)}(z)=g^{(d-1)}\left(g_{1}(z)\right)=g_{0}\left(g_{1}\left(\ldots g_{1}(z) \ldots\right)\right)
$$

Problem: Sometimes it is difficult to extract explicit probabilities for numbers of second neighbors and it is hard to evaluate $n$ derivatives (in order to recover the probabilities). Solution: calculate the average number of neighbors at distance $d$. At $z=1$ of the first derivative we can evaluate the average of a distribution (see Slide 16).

$$
\begin{aligned}
& \frac{\mathrm{d} g^{(2)}}{\mathrm{d} z}=g_{0}^{\prime}\left(g_{1}(z)\right) g_{1}^{\prime}(z) \stackrel{z=1, g_{1}(1)=1}{\Longrightarrow} c_{2}=g_{0}^{\prime}(1) g_{1}^{\prime}(1) \stackrel{\uparrow_{0}^{\prime}(1)=\langle k\rangle}{\Longrightarrow} g_{1}^{\prime}(k)=\sum_{k=0}^{\infty} k q_{k}= \\
& \begin{array}{c}
\text { mean number of } \\
\text { second neighbors }
\end{array} \\
&\langle k\rangle \frac{1}{k=0}<\infty \\
& k(k+1) p_{k+1}=\frac{1}{\langle k\rangle}\left(\left\langle k^{2}\right\rangle-\langle k\rangle\right)
\end{aligned}
$$

Conclusion: $c_{2}=\left\langle k^{2}\right\rangle-\langle k\rangle$ and more general

$$
c_{d}=\left(\frac{c_{2}}{c_{1}}\right)^{d-1} c_{1} \Longrightarrow
$$

Condition of giant component's existance in configuration model is $\left\langle k^{2}\right\rangle-2\langle k\rangle>0$
[MR] A critical point for random graphs with given degree sequence

## Let's use theory for practical results ...

Given a network with power-law distribution $p_{k}=C k^{-\alpha}, \alpha>0, k>0$ Reminder: $C$ is calculated from normalization condition, i.e., $C=1 / \zeta(\alpha)$

$$
p_{k}= \begin{cases}0 & k=0 \\ k^{-\alpha} / \zeta(\alpha) & k>0\end{cases}
$$

This network will have a giant component iff $\left\langle k^{2}\right\rangle-2\langle k\rangle>0$

$$
\begin{gathered}
\langle k\rangle=\sum_{k=0}^{\infty} k p_{k}=\frac{1}{\zeta(\alpha)} \sum_{k=1}^{\infty} k^{-\alpha+1}=\frac{\zeta(\alpha-1)}{\zeta(\alpha)} \\
\left\langle k^{2}\right\rangle=\sum_{k=0}^{\infty} k^{2} p_{k}=\frac{1}{\zeta(\alpha)} \sum_{k=1}^{\infty} k^{-\alpha+2}=\frac{\zeta(\alpha-2)}{\zeta(\alpha)} \\
\Longrightarrow \zeta(\alpha-2)>2 \zeta(\alpha-1)
\end{gathered}
$$



Figure 13.8: Graphical solution of Eq. (13.138). The configuration model with a pure power-law degree distribution (Eq. (13.133)) has a giant component if $\zeta(\alpha-2)>2 \zeta(\alpha-1)$. This happens for values of $\alpha$ below the crossing point of the two curves.

