

Aalto University School of Science and Technology

CS-E5745 Mathematical Methods for Network Science

Mikko Kivelä

Department of Computer Science Aalto University, School of Science mikko.kivela@aalto.fi

January 25, 2024

Generating functions and their use in networks

Learning goals this week:

- Recap of probability generating functions from last week
- Learn how to use PGFs to solve component size distributions in networks
- We will be following the Section 13 in Newman: Networks, An Introduction



PGFs from last week

Definition:

$$g(z) = p(0) + p(1)z + p(2)z^2 \dots = \sum_{k=0}^{\infty} p(k)z^k$$
 (1)

• p(k) can be extracted through derivation:

$$p(k) = \left[\frac{1}{k!}\frac{d^k}{dz^k}g(z)\right]_{z=0}$$
(2)

-

Moments can also be calculated through derivation:

$$\langle X^m \rangle = \left[z \frac{d}{dz} \dots z \frac{d}{dz} g(z) \right]_{z=1} = \left[(z \frac{d}{dz})^m g(z) \right]_{z=1}$$
 (3)



PGFs from last week

Sums of independent RVs

$$g_{X_1+X_2}(z) = g_{X_1}(z) * g_{X_2}(z)$$
(4)

$$g_{\sum_{i=1}^{N} X_{i}}(z) = [g_{X_{i}}(z)]^{N}$$
(5)

$$g_{X_1+c}(z) = g_{X_1}(z) * z^c$$
 (6)

• If N is also a RV in
$$S = \sum_{i=1}^{N} X_i$$
:

$$g_{\mathcal{S}}(z) = g_{\mathcal{N}}(g_{X_i}(z)) \tag{7}$$



Notation for networks (from Newman)

For the degree distribution p(k):

$$g_0(z) = \sum_{k=0}^{\infty} p(k) z^k$$

► For the excess degree distribution *q*(*k*):

$$g_1(z)=\sum_{k=0}^{\infty}q(k)z^k$$

These two are related:

$$g_1(z) = rac{1}{\langle k
angle} rac{d}{dz} g_0(z)$$



Solving the Galton-Watson process for networks

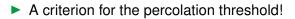
 Last week we derived the equations for Galton-Watson processes

$$g_{K_1}(z) = g_0(z)$$

 $g_{K_d}(z) = g_{K_{d-1}}(g_1(z))$

Solution for the expected value:

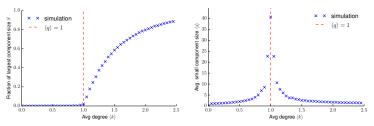
$$\langle \mathcal{K}_d \rangle = \langle q \rangle^{d-1} \langle k \rangle = \left(\frac{\langle k^2 \rangle - \langle k \rangle}{\langle k \rangle} \right)^{d-1} \langle k \rangle$$





Solving the Galton-Watson process for networks

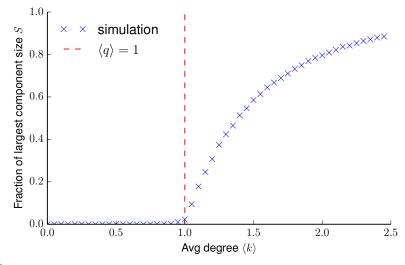
- We have the percolation threshold, but we are still missing
 - Shape of the relative giant size curve ($\sum_d K_d$ in the giant)
 - Size distributions of small components (S_d K_d when not in the giant)
- Note: GW process approximation for networks only works when d is small





Giant component size in ER networks

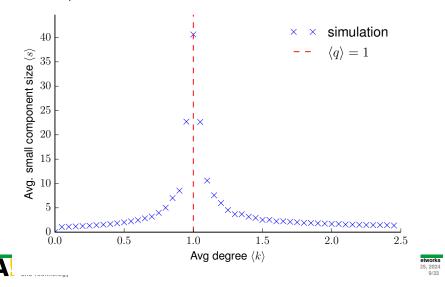
• Largest component size in ER networks with $N = 10^5$





Expected small component size in ER networks

Expected size of components other than the largest component in ER networks with $N = 10^5$



Uniqueness of the giant component

- Assume that there are two large components in ER networks with S₁, S₂ > 0 proportion of all nodes in them
- Number of nodes: $S_1 N$ and $S_2 N$
- Possible edges between the components: $S_1 N S_2 N = S_1 S_2 N^2$
- Probability that no two pairs are connected: $q = (1 - p)^{S_1 S_2 N^2} = (1 - \frac{\langle k \rangle}{N-1})^{S_1 S_2 N^2}$
- ► In "thermodynamic limit" ($N \to \infty$, s.t. $\langle k \rangle$ constant): $q = q_0 e^{-\langle k \rangle S_1 S_2 N} \to 0$



Solving for giant component size

- Idea: Instead of the GW process, we write down "self-consistency equations"
 - Write down the probability random node not being in giant u as a function of u
 - The fixed point where u = f(u) gives us u
 - The probability of random node in giant S = 1 u



Solving for giant component size for ER networks

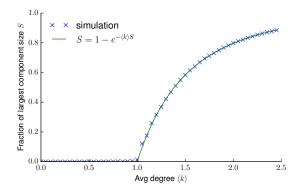
- Definition: u is the probability that (uniformly) randomly selected node doesn't belong to the giant component
 - Note that for Poisson degree distributions: p(k) = q(k)
- If node i doesn't belong to the giant then for every other node j:
 - ► There is no connection between *i* and *j* (probability 1 − *p*), or
 - There is connection, but j doesn't belong to the giant (probability pu)
- ► In total the following must hold $u = [(1 p) + pu]^{N-1}$

•
$$u = e^{-\langle k \rangle (1-u)}$$
, when $N \to \infty$



Solving for giant component size for ER networks

Shape of the relative giant size curve for ER network (with $N = 10^5$ nodes)





Solving for giant component size for configuration model (1/3)

- Definition: u is the probability that following an edge (and removing it) doesn't lead to the giant component
- If node *i* doesn't belong to the giant then none of the neighbors *j* belong to the giant
 - The number of new neighbors the node *i* has *k* is distributed as the excess degree q(k)
 - Probability that none of the k neighbors is in the giant is u^k
- ► In total the following must hold $u = \sum_{k=0}^{\infty} q(k) u^k$



Solving for giant component size for configuration model (2/3)

- ► In total the following must hold $u = \sum_{k=0}^{\infty} q(k)u^k$
 - Using the definition of the PGF [Eq. (1)]: $u = g_1(u)$
 - Probability of uniformly randomly selected node not being in the giant is ∑_{k=0}[∞] p(k)u^k = g₀(u)
- Returns the previous result for ER networks: $u = e^{-\langle k \rangle (1-u)}$



Solving for giant component size for configuration model (3/3)

- We can also solve the component size distributions of non-giant components
- Write equations for component size distributions using component size distributions
 - Leads to nice equations when done using PGF's



- Separate giant component size and small component size distribution
- S: Probability that uniformly randomly selected node belongs to the giant component
- RV s: size of non-giant component where uniformly randomly selected node belongs to
- Notation from Newman:

$$\blacktriangleright P(s=k)=\pi_k$$

$$\blacktriangleright h_0(z) = g_s(z)$$

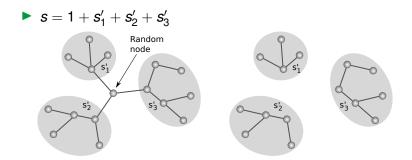
Sum of the above probabilities is one: $h_0(1) + S = 1$



- RV s': Follow a link and remove it. s' is the size of the non-giant component after that link is removed ("excess small component size")
- Notation from Newman:

$$\blacktriangleright P(s'=k) = \rho_k$$

$$h_1(z) = g_{s'}(z)$$

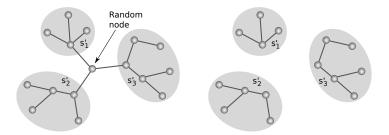




Mathematical Methods for Networks January 25, 2024 19/33

• RVs s, s', and the first neighborhood size k_1 are related



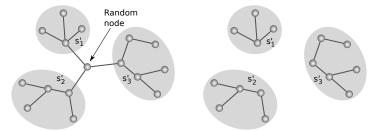




Mathematical Methods for Networks January 25, 2024 20/33

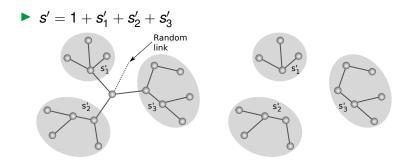
- ▶ Notation from Newman: $h_0(z) = g_s(z)$ and $h_1(z) = g_{s'}(z)$
- Using GF properties

$$egin{array}{rcl} m{s} &=& 1+\sum_{k=0}^{\kappa_1}m{s}' \iff \ h_0(z) &=& zg_0(h_1(z)) \end{array}$$





Mathematical Methods for Networks January 25, 2024 21/33

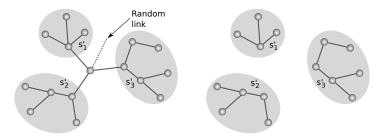




Mathematical Methods for Networks January 25, 2024 22/33

▶ $s'(h_1)$ and the excess degree $K_{1,i}(g_1)$ are related

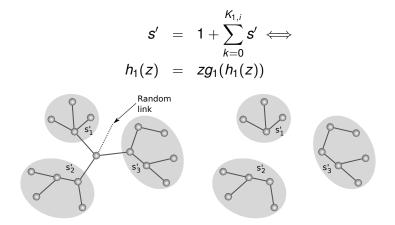
$$s'=1+\sum_{k=0}^{K_{1,i}}s'$$





Mathematical Methods for Networks January 25, 2024 23/33

Using GF properties





Mathematical Methods for Networks January 25, 2024 24/33

In total we have

$$h_0(z) = zg_0(h_1(z))$$
 (8)
 $h_1(z) = zg_1(h_1(z))$ (9)

- Solving h₀(z) possible but not easy
- ▶ Solving *h*₀(1) easier
 - Remember: $h_0(1) + S = 1$



• Using
$$h_0(1) + S = 1$$
 and Eq. (8):

$$S = 1 - h_0(1) = 1 - g_0(h_1(1))$$

• The value $h_0(1)$ can be solved from Eq. (9):

$$h_1(1) = g_1(h_1(1))$$

Using Newmans notation: u = h₁(1) (=probability that following a link does not lead to the giant component):

$$S = 1 - g_0(u)$$
 (10)
 $u = g_1(u)$ (11)



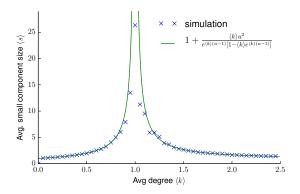
With some (home)work one can use previous equations to solve for the expected small component size

$$\langle s \rangle = 1 + \frac{g_0'(1)u^2}{g_0(u)[1 - g_1'(u)]}$$
 (12)



Solving for giant component size for ER networks

Shape of the expected size of the small components in ER network (with $N = 10^5$ nodes)





- With even more work one can find a formula for the whole small component size distribution!
 - See Newman pp. 468–469

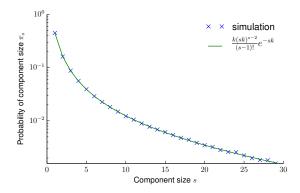
$$\pi_{s} = \frac{\langle k \rangle}{(s-1)!} \left[\frac{d^{s-2}}{dz^{s-2}} \left[g_{1}(z) \right]^{s} \right]_{z=0}$$
(13)

▶ Note that the formula works only for s > 1 and $\pi_1 = p_0$



Solving for component size distribution for ER networks

• Component sizes in ER network (with $N = 10^7$ nodes and $\langle k \rangle = 0.8$)





Summary on component sizes

We can now solve:

- Threshold for connectivity (percolation threshold)
- The (expected) number of nodes d steps away in BFS process
- The size of the giant component
- Component size distribution
- ... with some limitations:
 - Equations are derived for large configuration models (but might give good approximations for real networks)
 - We can write the equations but they might not have closed form solutions
 - Some equations do not work for the critical point (one can also solve for this separately)



Extensions

 Similar ideas (PGF's, self-consistency equations ...) can be used for many extensions

- Networks with degree correlations
- Networks with triangles
- Mutual connectivity
- ▶ ..

Some of these extensions are done in the possible projects



Percolation problems

- In percolation problems one removes nodes or edges can calculates the connected components
- Equivalent to finding component size distributions with altered degree distributions! (if done for configuration model or ER networks)
 - If K and K' are RV's of the original and the modified degree distribution, then percolation methods are given by P(K' = k|K = m)

•
$$P(K' = k) = \sum_{m=0}^{\infty} P(K' = k | K = m) * P(K = m)$$

- $g_{K'}(z) = \sum_{k=0}^{\infty} \sum_{m=0}^{\infty} P(K' = k | K = m) * P(K = m) z^k$
- Some projects might include percolation done in this way

