

# 6.207/14.15: Networks

## Lecture 4: Erdős-Renyi Graphs and Phase Transitions

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# Outline

- Phase transitions
- Connectivity threshold
- Emergence and size of a giant component
- An application: contagion and diffusion

## Reading:

- Jackson, Sections 4.2.2-4.2.5, and 4.3.

## Phase Transitions for Erdős-Renyi Model

- Erdős-Renyi model is completely specified by the link formation probability  $p(n)$ .
- For a given property  $A$  (e.g. connectivity), we define a **threshold function**  $t(n)$  as a function that satisfies:

$$\mathbb{P}(\text{property } A) \rightarrow 0 \quad \text{if} \quad \frac{p(n)}{t(n)} \rightarrow 0, \text{ and}$$

$$\mathbb{P}(\text{property } A) \rightarrow 1 \quad \text{if} \quad \frac{p(n)}{t(n)} \rightarrow \infty.$$

- This definition makes sense for “monotone or increasing properties,” i.e., properties such that if a given network satisfies it, any supernetwork (in the sense of set inclusion) satisfies it.
- When such a threshold function exists, we say that a **phase transition** occurs at that threshold.
- Exhibiting such phase transitions was one of the main contributions of the seminal work of Erdős and Renyi 1959.

# Threshold Function for Connectivity

## Theorem

(Erdős and Renyi 1961) A threshold function for the connectivity of the Erdős and Renyi model is  $t(n) = \frac{\log(n)}{n}$ .

- To prove this, it is sufficient to show that when  $p(n) = \lambda(n) \frac{\log(n)}{n}$  with  $\lambda(n) \rightarrow 0$ , we have  $\mathbb{P}(\text{connectivity}) \rightarrow 0$  (and the converse).
- However, we will show a stronger result: Let  $p(n) = \lambda \frac{\log(n)}{n}$ .

$$\text{If } \lambda < 1, \quad \mathbb{P}(\text{connectivity}) \rightarrow 0, \quad (1)$$

$$\text{If } \lambda > 1, \quad \mathbb{P}(\text{connectivity}) \rightarrow 1. \quad (2)$$

*Proof:*

- We first prove claim (1). To show disconnectedness, it is sufficient to show that the probability that **there exists at least one isolated node** goes to 1.

## Proof (Continued)

- Let  $I_i$  be a Bernoulli random variable defined as

$$I_i = \begin{cases} 1 & \text{if node } i \text{ is isolated,} \\ 0 & \text{otherwise.} \end{cases}$$

- We can write the probability that an individual node is isolated as

$$q = \mathbb{P}(I_i = 1) = (1 - p)^{n-1} \approx e^{-pn} = e^{-\lambda \log(n)} = n^{-\lambda}, \quad (3)$$

where we use  $\lim_{n \rightarrow \infty} \left(1 - \frac{a}{n}\right)^n = e^{-a}$  to get the approximation.

- Let  $X = \sum_{i=1}^n I_i$  denote the total number of isolated nodes. Then, we have

$$\mathbb{E}[X] = n \cdot n^{-\lambda}. \quad (4)$$

- For  $\lambda < 1$ , we have  $\mathbb{E}[X] \rightarrow \infty$ . We want to show that this implies  $\mathbb{P}(X = 0) \rightarrow 0$ .

- In general, this is not true.
- Can we use a Poisson approximation (as in the example from last lecture)? No, since the random variables  $I_i$  here are dependent.
- We show that the variance of  $X$  is of the same order as its mean.

## Proof (Continued)

- We compute the variance of  $X$ ,  $\text{var}(X)$ :

$$\begin{aligned}\text{var}(X) &= \sum_i \text{var}(I_i) + \sum_i \sum_{j \neq i} \text{cov}(I_i, I_j) \\ &= n\text{var}(I_1) + n(n-1)\text{cov}(I_1, I_2) \\ &= nq(1-q) + n(n-1)\left(\mathbb{E}[I_1 I_2] - \mathbb{E}[I_1]\mathbb{E}[I_2]\right),\end{aligned}$$

where the second and third equalities follow since the  $I_i$  are identically distributed Bernoulli random variables with parameter  $q$  (dependent).

- We have

$$\begin{aligned}\mathbb{E}[I_1 I_2] &= \mathbb{P}(I_1 = 1, I_2 = 1) = \mathbb{P}(\text{both 1 and 2 are isolated}) \\ &= (1-p)^{2n-3} = \frac{q^2}{(1-p)}.\end{aligned}$$

- Combining the preceding two relations, we obtain

$$\begin{aligned}\text{var}(X) &= nq(1-q) + n(n-1)\left[\frac{q^2}{(1-p)} - q^2\right] \\ &= nq(1-q) + n(n-1)\frac{q^2 p}{1-p}.\end{aligned}$$

## Proof (Continued)

- For large  $n$ , we have  $q \rightarrow 0$  [cf. Eq. (3)], or  $1 - q \rightarrow 1$ . Also  $p \rightarrow 0$ . Hence,

$$\begin{aligned} \text{var}(X) &\sim nq + n^2 q^2 \frac{p}{1-p} \sim nq + n^2 q^2 p \\ &= nn^{-\lambda} + \lambda n \log(n) n^{-2\lambda} \\ &\sim nn^{-\lambda} = \mathbb{E}[X], \end{aligned}$$

where  $a(n) \sim b(n)$  denotes  $\frac{a(n)}{b(n)} \rightarrow 1$  as  $n \rightarrow \infty$ .

- This implies that

$$\mathbb{E}[X] \sim \text{var}(X) \geq (0 - \mathbb{E}[X])^2 \mathbb{P}(X = 0),$$

and therefore,

$$\mathbb{P}(X = 0) \leq \frac{\mathbb{E}[X]}{\mathbb{E}[X]^2} = \frac{1}{\mathbb{E}[X]} \rightarrow 0.$$

- It follows that  $\mathbb{P}(\text{at least one isolated node}) \rightarrow 1$  and therefore,  $\mathbb{P}(\text{disconnected}) \rightarrow 1$  as  $n \rightarrow \infty$ , completing the proof.

## Converse

- We next show claim (2), i.e., if  $p(n) = \lambda \frac{\log(n)}{n}$  with  $\lambda > 1$ , then  $\mathbb{P}(\text{connectivity}) \rightarrow 1$ , or equivalently  $\mathbb{P}(\text{disconnectivity}) \rightarrow 0$ .
- From Eq. (4), we have  $\mathbb{E}[X] = n \cdot n^{-\lambda} \rightarrow 0$  for  $\lambda > 1$ .
- This implies probability of having isolated nodes goes to 0. However, we need more to establish connectivity.
- The event “graph is disconnected” is equivalent to the existence of  $k$  nodes without an edge to the remaining nodes, for some  $k \leq n/2$ .
- We have

$$\mathbb{P}(\{1, \dots, k\} \text{ not connected to the rest}) = (1 - p)^{k(n-k)},$$

and therefore,

$$\mathbb{P}(\exists k \text{ nodes not connected to the rest}) = \sum_{k=1}^{n/2} \binom{n}{k} (1 - p)^{k(n-k)}.$$



## Converse (Continued)

- Using the union bound [i.e.  $\mathbb{P}(\cup_i A_i) \leq \sum_i \mathbb{P}(A_i)$ ], we obtain

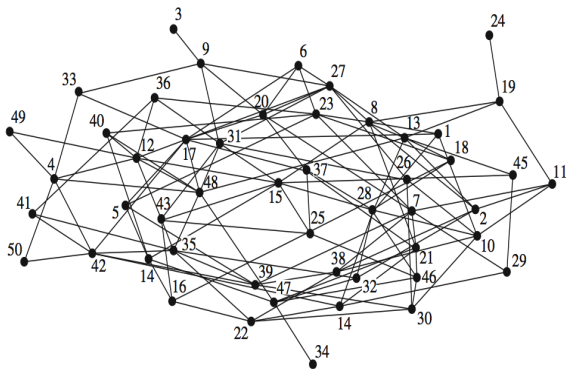
$$\mathbb{P}(\text{disconnected graph}) \leq \sum_{k=1}^{n/2} \binom{n}{k} (1-p)^{k(n-k)}.$$

- Using Stirling's formula  $k! \sim \left(\frac{k}{e}\right)^k$ , which implies  $\binom{n}{k} \leq \frac{n^k}{\left(\frac{k}{e}\right)^k}$  in the preceding relation and some (ugly) algebra, we obtain

$$\mathbb{P}(\text{disconnected graph}) \rightarrow 0,$$

completing the proof.

# Phase Transitions — Connectivity Threshold



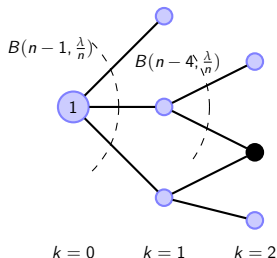
**Figure:** Emergence of connectedness: a random network on 50 nodes with  $p = 0.10$ .

# Giant Component

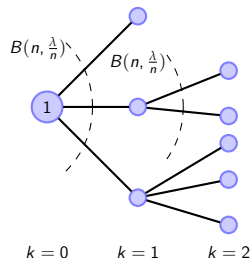
- We have shown that when  $p(n) \ll \frac{\log(n)}{n}$ , the Erdős-Renyi graph is disconnected with high probability.
- In cases for which the network is not connected, the component structure is of interest.
- We have argued that in this regime the expected number of isolated nodes goes to infinity. This suggests that the Erdős-Renyi graph should have an arbitrarily large number of components.
- We will next argue that the threshold  $p(n) = \frac{\lambda}{n}$  plays an important role in the component structure of the graph.
  - For  $\lambda < 1$ , all components of the graph are “small”.
  - For  $\lambda > 1$ , the graph has a **unique giant component**, i.e., a component that contains a constant fraction of the nodes.

# Emergence of the Giant Component—1

- We will analyze the component structure in the vicinity of  $p(n) = \frac{\lambda}{n}$  using a branching process approximation.
- We assume  $p(n) = \frac{\lambda}{n}$ .
- Let  $B(n, \frac{\lambda}{n})$  denote a binomial random variable with  $n$  trials and success probability  $\frac{\lambda}{n}$ .
- Consider starting from an arbitrary node (node 1 without loss of generality), and exploring the graph.



(a) Erdos-Renyi graph process.



(b) Branching Process Approx.

## Emergence of the Giant Component—2

- We first consider the case when  $\lambda < 1$ .
- Let  $Z_k^G$  and  $Z_k^B$  denote the number of individuals at stage  $k$  for the graph process and the branching process approximation, respectively.
- In view of the “overcounting” feature of the branching process, we have

$$Z_k^G \leq Z_k^B \quad \text{for all } k.$$

- From branching process analysis (see Lecture 3 notes), we have

$$\mathbb{E}[Z_k^B] = \lambda^k,$$

(since the expected number of children is given by  $n \times \frac{\lambda}{n} = \lambda$ ).

- Let  $S_1$  denote the number of nodes in the Erdős-Renyi graph connected to node 1, i.e., the size of the component which contains node 1.
- Then, we have

$$\mathbb{E}[S_1] = \sum_k \mathbb{E}[Z_k^G] \leq \sum_k \mathbb{E}[Z_k^B] = \sum_k \lambda^k = \frac{1}{1 - \lambda}.$$

## Emergence of the Giant Component—3

- The preceding result suggests that for  $\lambda < 1$ , the sizes of the components are “small”.

### Theorem

Let  $p(n) = \frac{\lambda}{n}$  and assume that  $\lambda < 1$ . For all (sufficiently large)  $a > 0$ , we have

$$\mathbb{P}\left(\max_{1 \leq i \leq n} |S_i| \geq a \log(n)\right) \rightarrow 0 \quad \text{as } n \rightarrow \infty.$$

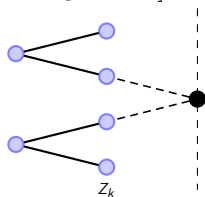
Here  $|S_i|$  is the size of the component that contains node  $i$ .

- This result states that for  $\lambda < 1$ , all components are small [in particular they are of size  $O(\log(n))$ ].
- Proof is beyond the scope of this course.

## Emergence of the Giant Component—4

- We next consider the case when  $\lambda > 1$ .
- We claim that  $Z_k^G \approx Z_k^B$  when  $\lambda^k \leq O(\sqrt{n})$ .
- The expected number of conflicts at stage  $k + 1$  satisfies

$$\mathbb{E}[\text{number of conflicts at stage } k + 1] \leq np^2 \mathbb{E}[Z_k^2] = n \frac{\lambda^2}{n^2} \mathbb{E}[Z_k^2].$$



- We assume for large  $n$  that  $Z_k$  is a Poisson random variable and therefore  $\text{var}(Z_k) = \lambda^k$ . This implies that

$$\mathbb{E}[Z_k^2] = \text{var}(Z_k) + \mathbb{E}[Z_k]^2 = \lambda^k + \lambda^{2k} \approx \lambda^{2k}.$$

- Combining the preceding two relations, we see that the conflicts become non-negligible only after  $\lambda^k \approx \sqrt{n}$ .

## Emergence of the Giant Component—5

- Hence, there exists some  $c > 0$  such that  $\mathbb{P}(\text{there exists a component with size } \geq c\sqrt{n} \text{ nodes}) \rightarrow 1$  as  $n \rightarrow \infty$ .
- Moreover, between any two components of size  $\sqrt{n}$ , the probability of having a link is given by

$$\mathbb{P}(\text{there exists at least one link}) = 1 - \left(1 - \frac{\lambda}{n}\right)^n \approx 1 - e^{-\lambda},$$

i.e., it is a positive constant independent of  $n$ .

- This argument can be used to see that components of size  $\leq \sqrt{n}$  connect to each other, forming a connected component of size  $qn$  for some  $q > 0$ , **a giant component**.



## Size of the Giant Component

- Form an Erdős-Renyi graph with  $n - 1$  nodes with link formation probability  $p(n) = \frac{\lambda}{n}$ ,  $\lambda > 1$ .
- Now add a last node, and connect this node to the rest of the graph with probability  $p(n)$ .
- Let  $q$  be the **fraction of nodes in the giant component** of the  $n - 1$  node network. We can assume that for large  $n$ ,  $q$  is also the fraction of nodes in the giant component of the  $n$ -node network.
- The probability that node  $n$  is not in the giant component is given by

$$\mathbb{P}(\text{node } n \text{ not in the giant component}) = 1 - q \equiv \rho.$$

- The probability that node  $n$  is not in the giant component is equal to the probability that none of its neighbors is in the giant component, yielding

$$\rho = \sum_d P_d \rho^d \equiv \Phi(\rho).$$

- Similar to the analysis of branching processes, we can show that this equation has a fixed point  $\rho^* \in (0, 1)$ .

## An Application: Contagion and Diffusion

- Consider a society of  $n$  individuals.
- A randomly chosen individual is infected with a contagious virus.
- Assume that the network of interactions in the society is described by an Erdős-Renyi graph with link probability  $p$ .
- Assume that any individual is immune with a probability  $\pi$ .
- We would like to find the expected size of the epidemic as a fraction of the whole society.
- The spread of disease can be modeled as:
  - Generate an Erdős-Renyi graph with  $n$  nodes and link probability  $p$ .
  - Delete  $\pi n$  of the nodes uniformly at random.
  - Identify the component that the initially infected individual lies in.
- We can equivalently examine a graph with  $(1 - \pi)n$  nodes with link probability  $p$ .

# An Application: Contagion and Diffusion

- We consider 3 cases:
- $\rho(1 - \pi)n < 1$ :

$$\mathbb{E}[\text{size of epidemic as a fraction of the society}] \leq \frac{\log(n)}{n} \approx 0.$$

- $1 < \rho(1 - \pi)n < \log((1 - \pi)n)$ :

$$\begin{aligned} \mathbb{E}[\text{size of epidemic as a fraction of the society}] \\ = \frac{q\rho(1 - \pi)n + (1 - q)\log((1 - \pi)n)}{n} \approx q^2(1 - \pi), \end{aligned}$$

where  $q$  denotes the fraction of nodes in the giant component of the graph with  $(1 - \pi)n$  nodes, i.e.,  $q = 1 - e^{-q(1 - \pi)\rho}$ .

- $\rho > \frac{\log((1 - \pi)n)}{(1 - \pi)n}$ :

$$\mathbb{E}[\text{size of epidemic as a fraction of the society}] = (1 - \pi).$$