



# Network theory Part III

**Complexity in Social Systems** AA 2023/2024 **Maxime Lucas** Lorenzo Dall'Amico





#### **Recap last lecture**

#### **Models**

Erdos-Renyi model Watts-Strogatz model **Configuration model** Chung-Lu model





#### **Concepts**

#### Importance of scale-freeness Giant component Phase transitions

# **Today's topics**

#### **Models**

Barabasi-Albert model Bianconi-Barabasi model Link/Copying model

Origins of scale-free distributions Growing networks Assortativity and correlations Bonus: Robustness





### Network heterogeneity

Where does it come from?



#### First ingredient: Growth/time

#### Second ingredient: Not all links are equally likely!





# **Network heterogeneity**

Microscopic mechanisms for macroscopic observables

In static ensemble models (last lecture) we defined network by constraints In evolving/growing network, we define growth rules and look for asymptotic stationary behaviour

(1) Networks continuously expand by the addition of new nodes WWW : addition of new documents

(2) New nodes prefer to link to highly connected nodes. (WWW : linking to well known sites)

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#### **Barabasi-Albert network model**

#### **GROWTH:**

At each timestep we add a new node with  $m (\leq m_0)$  links that connect the new node to *m* nodes already in the network.

#### **PREFERENTIAL ATTACHMENT:**

the probability that a node connects to a node with k links is proportional to k.

$$\Pi(k_i) = \frac{k_i}{\sum_j k_j}$$



Degree dynamics

$$\frac{dk_i}{dt} = m\Pi(k_i) = m\frac{k_i}{\sum_{j=1}^{N-1} k_j} \qquad \sum_{j=1}^{N-1} k_j = \frac{k_j}{\sum_{j=1}^{N-1} k_j}$$

$$\frac{\partial k_i}{k_i} = \frac{\partial t}{2t} \qquad \int_m^k \frac{\partial k_1}{k_i} = \int_{t_i}^t \frac{\partial t}{2t} \qquad \ln\left(\frac{k}{m}\right) = \frac{1}{2}$$

$$k_i(t) = m\left(\frac{t}{t_i}\right)^{\beta}$$
  $\beta = \frac{1}{2}$  dynamica

Barabási, Albert-László, Réka Albert, and Hawoong Jeong. "Mean-field theory for scale-free random networks." Physica A: Statistical Mechanics and its Applications 272.1-2 (1999): 173-187.





We call  $\beta$  the *dynamical exponent* and has the value

#### Degree distribution: simple derivation



$$\sum_{j=1}^{N-1} k_j = 2m\beta = \frac{1}{2}m$$

Equation (5.7) offers a number of predictions:

 $P(k_i(t) < k) = 1 - P(k_i(t) > k) = 1 - P(k_$ hetwork models with real data, we • The degree of each node increases following a power-law with the same A node *i* can come with equal probability any time between  $t_i = m_0$  and *t*, hence:  $\frac{1}{e}$  scales: the sa

#### • The growth in the degrees is sublinear (i.e. $\beta < 1$ ). This is a consequence World Wide Web of the growing nature of the Barab == Number of nodes with degree larger than k page was created in נפבו. נועפוו its trillion documents, more nodes to link to than the previous node. Hence, with time the exthe WWW added a node each milliistin There are overallinks with an indreasing pool of ther nodes. second (10<sup>-3</sup> sec).

• The configure added, the bighen is its degree *k*(t). Hence, but a random node has degree k or less is:

er advantage in marketing and business.

$$P(k) \sim 1 - \left(\frac{m}{k}\right)$$

• The rate at which the node kacquires new links is given by the derivative of (5.7) 1 I R

$$p_k \sim \frac{\partial P(k)}{\partial k} \frac{1}{dt} \frac{1}{dt} \frac{m^{1/p}}{\frac{1}{p}} \frac{1}{\sqrt{k_t}} \frac{1}{(\beta+1)}$$
(5.8)

indicating that in each time frame older nodes acquire more links (as they have smaller  $t_i$ ). Furthermore the rate at which a node acquires links decreases with time as  $2^{1/2}$ . Hence, few pand fewer links go to a node.  $P_k$ 

In summary, the Barabási-Albert model captures the fact that in real networks nodes arrive one after the other, offering a dynamical descrip-<sup>104</sup>tion of a netwissis ogreat but ethere are iterrors in the coefficient dition of each new node which the older nodes have an advantage over the younger ones, eventual-

#### TIME IN NETWORKS

 $k_i(t) = m$ 

# have to decide how to measure time

cell is the result of 4 billion years of evolution. With roughly 20,000 genes in a human cell, on average the cellular network added a node every 200,000 years (~10<sup>12</sup> sec).

Given these enormous time-scale differences it is impossible to use real time to compare the dynamics of different networks. Therefore, in network theory we use event time, advancing our time-step by one each time when there is a change in the network topology.

For example, in the Barabási-Albert corresponds to a new time step,





#### Rate-equation derivation

Number of degree k nodes at time t: < N(k,t) >= tP(k,t)

Number of links added to degree k nodes after the arrival of a new node:

$$(N+1)P(k,t+1) = NP(k,t)$$

k-nodes at time t+1

# k-nodes at time t

We do not have k=0,1,...,m-1 nodes in the network (each node arrives with degree m) Requires separate equation for degree m nodes.

$$(N+1)P(m,t+1) = NP(m,t) + 1 - \frac{m}{2}P(m,t)$$





**Rate-equation derivation**  $(N+1)P(k,t+1) = NP(k,t) + \frac{k-1}{2}P(k-1,t) - \frac{k}{2}P(k,t)$   $(N+1)P(m,t+1) = NP(m,t) + 1 - \frac{m}{2}P(m,t)$ 

Impose stationarity!  $P(k, \infty) = P(k)$  $(N+1)P(k,\infty) - NP(k,\infty) = P(k,\infty) = P(k)$ 

$$P(k) = \frac{k-1}{2}P(k-1)$$

 $P(m) = 1 - \frac{m}{2}P(m)$ 

 $-\frac{k}{2}P(k) \qquad P(k) = \frac{k-1}{k+2}P(k-1)$  $\rightarrow P(m) = \frac{2}{\sqrt{2}}$ m+2



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Rate-e

rabasi-Albert model  
quation derivation  

$$P(m) = \frac{2}{m+2}$$

$$P(m+1) = \frac{m}{m+3}P(m) = \frac{2m}{(m+2)(m+3)}$$

$$P(m+2) = \frac{m+1}{m+4}P(m+1) = \frac{2m(2m+1)}{(m+2)(m+3)(m+3)}$$

$$P(m+3) = \frac{m+2}{m+5}P(m+2) = \frac{2m(2m+1)}{(m+3)(m+4)(m+5)}$$
....= ...

$$P(k) = \frac{2m(2m+1)}{k(k+1)(k+2)} \sim$$

Krapivsky, Redner, Leyvraz, PRL 2000, Dorogovtsev, Mendes, Samukhin, PRL 2000, Bollobas et al, Random Struc. Alg. 2001

$$k^{-3}$$



#### γ 1 3 Barabasi-Albert model



(a) We generated networks with N=100,000and  $m_0=m=1$  (blue), 3 (green), 5 (grey), and 7 (orange). The fact that the curves are parallel to each other indicates that  $\gamma$  is independent of m and  $m_0$ . The slope of the purple line is -3, corresponding to the predicted degree exponent  $\gamma=3$ . Inset: (5.11) predicts  $p_k \sim 2m^2$ , hence  $p_k/2m^2$  should be independent of m. Indeed, by plotting  $p_k/2m^2$  vs. k, the data points shown in the main plot collapse into a single curve.

**(b)** The Barabási-Albert model predicts that  $p_k$  is independent of *N*. To test this we plot  $p_k$  for N = 50,000 (blue), 100,000 (green), and 200,000 (grey), with  $m_0 = m = 3$ . The obtained  $p_k$  are practically indistinguishable, indicating that the degree distribution is stationary, i.e. independent of time and system size.



Necessity of ingredients: growth









Necessity of ingredients: preferential attachment

$$\frac{\partial k_i}{\partial t} = \frac{N}{N-1} \frac{k_i}{2t} + k_i(t) = \frac{2(N-1)}{N(N-2)} t + Ct^{\frac{N}{2(N-1)}} \sim 0$$

p<sub>k</sub> : power law (initially) -> Gaussian -> Fully Connected





Necessity of ingredients?

#### Do we need both growth and preferential attachment? YEP.







Non-linear preferential attachments



Explicit derivation in Barabasi's book Sec 5.14





#### **Multiple questions:**

- why preferential attachment depends on k?
- Why linearly?
- Global (optimum) versus local (random) mechanisms?





Potential local mechanisms: link model

simplest example of a local mechanism that generates a scale-free network without preferential attachment



• *Growth*: At each time step we add a new node

• *Link Selection*: We select a link at random and connect the new node to one of the two nodes

lacks a built-in  $\Pi(k)$  function. Yet, it generates preferential attachment.

probability  $q_k$  that the node at the end of a randomly chosen link has degree k

 $q_k = 1$  $kp_k$ JU < k >

**Excess degree** distribution!!!!

Linear in k: pref attachment





Potential local mechanisms: copy model



- Random Connection: With probability p the new node links to u, which means that we link to the randomly selected web document.
- Copying: With probability 1-p we randomly choose an outgoing *link* of node *u* and link the new node to the link's target. In other words, the new webpage *copies* a link of node *u* and connects to its target, rather than connecting to node u directly.

$$\Pi(k) = \frac{p}{N} + \frac{1-p}{2L}k$$

probability of selecting a degree-k node





8

7

0

 $\langle d \rangle$ 

4

3

2

10<sup>1</sup>

Properties: diameter

# $\frac{\ln N}{\ln \ln N}$





Properties: clustering

Node I with degree k\_I has a number of triangles:

$$N_{r_l} = \int_{j=1}^{N} di \int_{i=1}^{N} dj P(i,j) P(i,l) P(j,l)$$

$$P(i,j) = m\Pi(k_i(j)) = m\frac{k_i(j)}{\sum_{l=1}^{j} k_l(j)} = m\frac{k_i(j)}{2}$$
  
using  $k_i(t) = m\left(\frac{t}{t_i}\right)^{\frac{1}{2}} = m\left(\frac{j}{i}\right)^{\frac{1}{2}} \to P(i,j) = \frac{m}{2}(ij)$ 

arrival time of node j is  $t_j = j$  and the arrival time of node i is  $t_i = i$ 

$$N_{r_l} = \frac{m^3}{8l} (\ln N)^2 \quad \to \quad C = \frac{2N_{r_l}}{k_l(N)(k_l(N) - 1)} \simeq \frac{m}{4} \frac{1}{4}$$

Konstantin Klemm, Victor M. Eguiluz,

Growing scale-free networks with small-world behavior, Phys. Rev. E 65, 057102 (2002),







#### Summary

- Power law with exp -3
- Ultrasmall world
- Undirected
- Vanishing clustering
- Does not capture:
  - variations in the shape of the degree distribution
  - variations in the degree exponent
  - size-independent clustering coefficient
- No other realistic methods:

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- Link deletion
- Internal links
- Ageing...
- Fitness

N = t

Number of Nodes N = t

Number of Links N = mt

Average Degree  $\langle k \rangle = 2m$ 

**Degree Dynamics**  $k_i(t) = m (t/t_i)^{\beta}$ 

**Dynamical Exponent**  $\beta = 1/2$ 

**Degree Distribution**  $p_k \sim k^{-\gamma}$ 

**Degree Exponent**  $\gamma = 3$ 

Average Distance  $\langle d \rangle \sim \log N / \log \log N$ 

Clustering Coefficient  $\langle C \rangle \sim (\ln N)^2 / N$ 



### **Beyond Barabasi-Albert model**

#### The Bianconi-Barabasi fitness model

**BA model:** first mover advantage! Can latecomers make it? **Fitness model!** 

$$\frac{\partial k_i}{\partial t} = m \frac{\eta_i k_i}{\sum_j \eta_j k_j} \quad \text{ansatz:} \quad k(t, t_i, \eta_i) = m \left(\frac{t}{t_i}\right)^{\beta}$$
$$\left\langle \sum_j \eta_j k_j \right\rangle = \int d\eta \rho(\eta) \eta \int_i^t dt_0 k_\eta(t, t_0) = \int d\eta \rho(\eta) \eta m \frac{t - t^{\beta(\eta)}}{1 - \beta(\eta)}$$
$$\left\langle \sum_j \eta_j k_j \right\rangle^{t \to \infty} Cmt[1 - O(t^{-\epsilon})]_{\epsilon = (1 - \max_\eta \beta(\eta)) > 0}$$
$$\frac{\partial k_\eta}{\partial k} = \frac{\eta k_\eta}{Ct} \qquad \beta(\eta) = \frac{\eta}{C} \quad 1 = \int_0^{\eta_{max}} d\eta \rho(\eta) \frac{1}{\frac{C}{\eta} - \frac{1}{2}}$$
$$Cmt = \sum_j k_j \le \eta_{max} \sum_j k_j = 2mt\eta_{max} \quad \rightarrow C \le 2\eta_{max}, \quad C > \eta_{max}$$





# **Beyond Barabasi-Albert model**

#### Derivation degree distribution

Number of nodes with degree larger than k

$$t_0 < t \left(\frac{m}{k}\right)^{C/r_k}$$

 $P(k_i < k) = 1 - P$ 

$$p(k) = \frac{\partial P(k)}{\partial k} = \int_0^{\eta_m}$$



$$\begin{aligned} (k_i > k) &= 1 - \frac{1}{m_0 + t} \int_0^{\eta_{max}} t\left(\frac{m}{k}\right)^{C/\eta} \rho(\eta) d\eta \\ &\sim 1 - \int_0^{\eta_{max}} \left(\frac{m}{k}\right)^{C/\eta} \rho(\eta) d\eta \end{aligned}$$

 $\sum_{\eta}^{\max} \frac{C}{\eta} m^{C/\eta} k^{-C/\eta+1)} \rho(\eta) d\eta$ 







# **Beyond Barabasi-Albert model**

Fitness examples: equal fitness

![](_page_25_Picture_2.jpeg)

![](_page_25_Figure_3.jpeg)

![](_page_25_Picture_4.jpeg)

![](_page_25_Picture_5.jpeg)

![](_page_25_Picture_6.jpeg)

# **Beyond Barabasi-Albert moc**

Limitations and extensions

![](_page_26_Figure_2.jpeg)

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MODEL CLASS	EXAMPLES	CHARACTERISTICS
Static Models	<ul> <li>Erdős–Rényi</li> <li>Watts-Strogatz</li> </ul>	<ul> <li><i>N</i> fixed</li> <li><i>p<sub>k</sub></i> exponentially bounded</li> <li>Static, time independent top</li> </ul>
Generative Models	<ul> <li>Configuration Model</li> <li>Hidden Parameter Model</li> </ul>	<ul> <li>Arbitrary pre-defined p<sub>k</sub></li> <li>Static, time independent top</li> </ul>
Evolving Network Models	<ul> <li>Barabási–Albert Model</li> <li>Bianconi-Barabási Model Initial Attractiveness Model Internal Links Model Node Deletion Model Accelerated Growth Model Aging Model</li> </ul>	<ul> <li><i>p<sub>k</sub></i> is determined by the proc that contribute to the networ evolution.</li> <li>Time-varying network topolo</li> </ul>

![](_page_26_Figure_4.jpeg)

![](_page_26_Figure_5.jpeg)

![](_page_26_Figure_6.jpeg)

![](_page_26_Picture_7.jpeg)

![](_page_26_Picture_8.jpeg)

#### Introduction to network correlations

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![](_page_27_Picture_1.jpeg)

**Figure 7.10: Friendship network at a US high school.** The vertices in this network represent 470 students at a US high school (ages 14 to 18 years). The vertices are color coded by race as indicated in the key. Data from the National Longitudinal Study of Adolescent Health [34, 314].

![](_page_27_Picture_3.jpeg)

### Introduction to network correlations

Homophily: This is not news to sociologists, who have long observed and discussed such divisions.

Assortative: like is associated with like

![](_page_28_Figure_3.jpeg)

Disassortative: like is associated with not-like

![](_page_28_Picture_5.jpeg)

![](_page_28_Picture_6.jpeg)

### **Enumerative assortativity**

Given ci class or type of vertex i  $(1,...,n_c = total number of classes)$ , then the total number of edges that run between vertices of the same type is:

$$\sum_{edges(i,j)} \delta(c_i, c_j) = \frac{1}{2} \sum_{ij} A_{ij} \delta(c_i, c_j)$$

However, we want to control for the random expectation of the mixing:

$$Q = \frac{1}{2m} \sum_{ij} \left( A \right)$$

**Modularity**: It is strictly less than 1, takes positive values if there are more edges between vertices of the same type than we would expect by chance, and negative ones if there are less.

![](_page_29_Picture_6.jpeg)

 $\frac{1}{2}\sum ij\frac{k_ik_j}{2m}\delta(c_i,c_j)$ 

 $A_{ij} - \frac{k_i k_j}{2m} \bigg) \, \delta(c_i, c_j)$ 

# **Enumerative assortativity**

How can we normalise it?

Obtained for full mixing:

Assortativity coefficient

![](_page_30_Picture_5.jpeg)

![](_page_30_Picture_6.jpeg)

 $Q_{max} = \frac{1}{2m} \left( 2m - \sum_{ij} \frac{k_i k_j}{2m} \delta(c_i, c_j) \right)$ 

 $\frac{Q}{Q_{max}} = \frac{\sum_{ij} \left( A_{ij} - \frac{k_i k_j}{2m} \right) \delta(c_i, c_j)}{\left( 2m - \sum_{ij} \frac{k_i k_j}{2m} \delta(c_i, c_j) \right)}$ 

![](_page_30_Picture_9.jpeg)

#### Scalar assortativity

![](_page_31_Figure_1.jpeg)

$$\mu = \frac{\sum_{ij} A_{ij} x_i}{\sum_{ij} A_{ij}} = \frac{\sum_i k_i x_i}{\sum_i k_i} = \frac{1}{2m} \sum_i k_i x_i$$

$$cov(x_i, x_j) = \frac{\sum_{ij} A_{ij} (x_i - \mu) (x_j - \mu)}{\sum_{ij} A_{ij}}$$

$$=\frac{1}{2m}\sum_{ij}(A_{ij}-\frac{k_ik_j}{2m})x_ix_j$$

$$r = \frac{\sum_{ij} (A_{ij} - k_i k_j/2m) x_i x_j}{\sum_{ij} (k_i \delta_{ij} - k_i k_j/2m) x_i x_j}$$

![](_page_31_Picture_6.jpeg)

#### Degree assortativity coefficient

It can be misleading when:

- complicated behavior of the correlation functions (non-monotonous behavior)
- Pearson coefficient gives a larger weight to the more abundant degree classes

#### Average nearest neighbour degree

$$k_{nn,i} = \frac{1}{k_i} \sum_{j \in \nu(i)} k_j \qquad k_{nn}$$

 $r = \frac{\sum_{ij} (A_{ij} - k_i k_j/2m) k_i k_j}{\sum_{ij} (k_i \delta_{ij} - k_i k_j/2m) k_i k_j}$ 

 $\overline{N}_k \not$  $\kappa_{nn,i}$  $-\sum_{K} K I (K | K)$ k'

![](_page_32_Picture_10.jpeg)

![](_page_32_Picture_11.jpeg)

What are the possible scenarios?

![](_page_33_Figure_2.jpeg)

![](_page_33_Picture_3.jpeg)

![](_page_33_Figure_4.jpeg)

![](_page_33_Figure_5.jpeg)

![](_page_33_Figure_6.jpeg)

![](_page_33_Figure_7.jpeg)

![](_page_33_Figure_8.jpeg)

e.

f.

![](_page_33_Figure_9.jpeg)

![](_page_33_Figure_10.jpeg)

![](_page_33_Picture_11.jpeg)

![](_page_34_Figure_3.jpeg)

![](_page_34_Figure_4.jpeg)

#### Origins: real or structural?

![](_page_35_Figure_2.jpeg)

#### **Degree Preserving Randomization with Simple Links (R-S)**

We apply degree-preserving randomization to the original network and at each step we make sure that we do not permit more than one link between a pair of nodes. On the algorithmic side this means that each rewiring that generates multi-links is discarded.

#### **Degree Preserving Randomization with Multiple Links (R-M)**

For a self-consistency check it is sometimes useful to perform degree-preserving randomization that allows for multiple links between the nodes. On the algorithmic side this means that we allow each random rewiring, even if it leads to multi-links

Origins: real or structural?

![](_page_36_Figure_2.jpeg)

![](_page_36_Figure_3.jpeg)

![](_page_36_Picture_4.jpeg)

# **Recap today's topics**

#### **Models**

Barabasi-Albert model Bianconi-Barabasi model Link/Copying model

Origins of scale-free distributions Growing networks Assortativity and correlations "Robustness"

![](_page_37_Picture_5.jpeg)

#### **Concepts**

Next time

Percolation! Spectral properties Random Walks

![](_page_37_Picture_9.jpeg)