

Kronecker graphs and the Structure of Large Networks

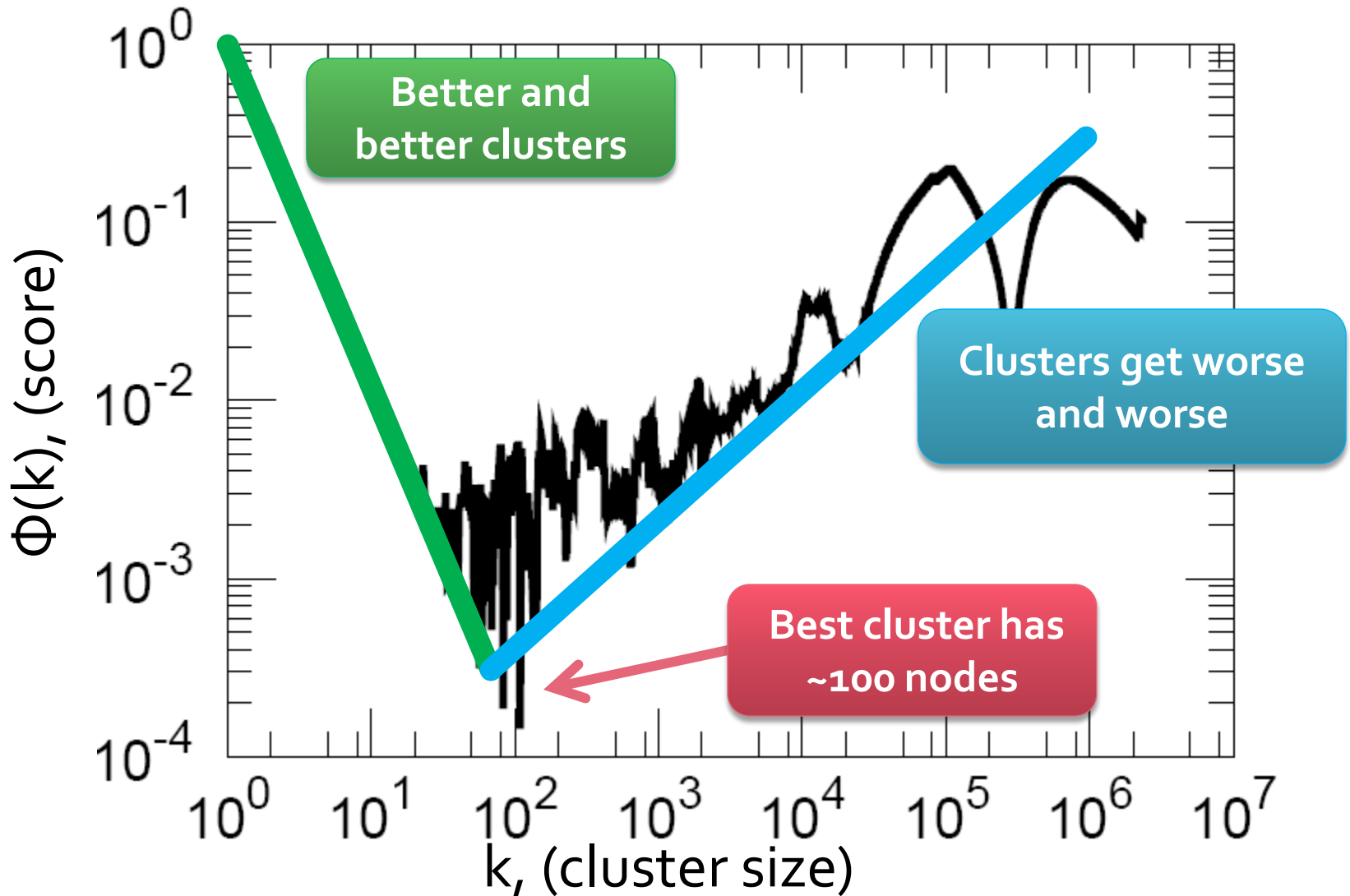
CS224W: Social and Information Network Analysis

Jure Leskovec, Stanford University

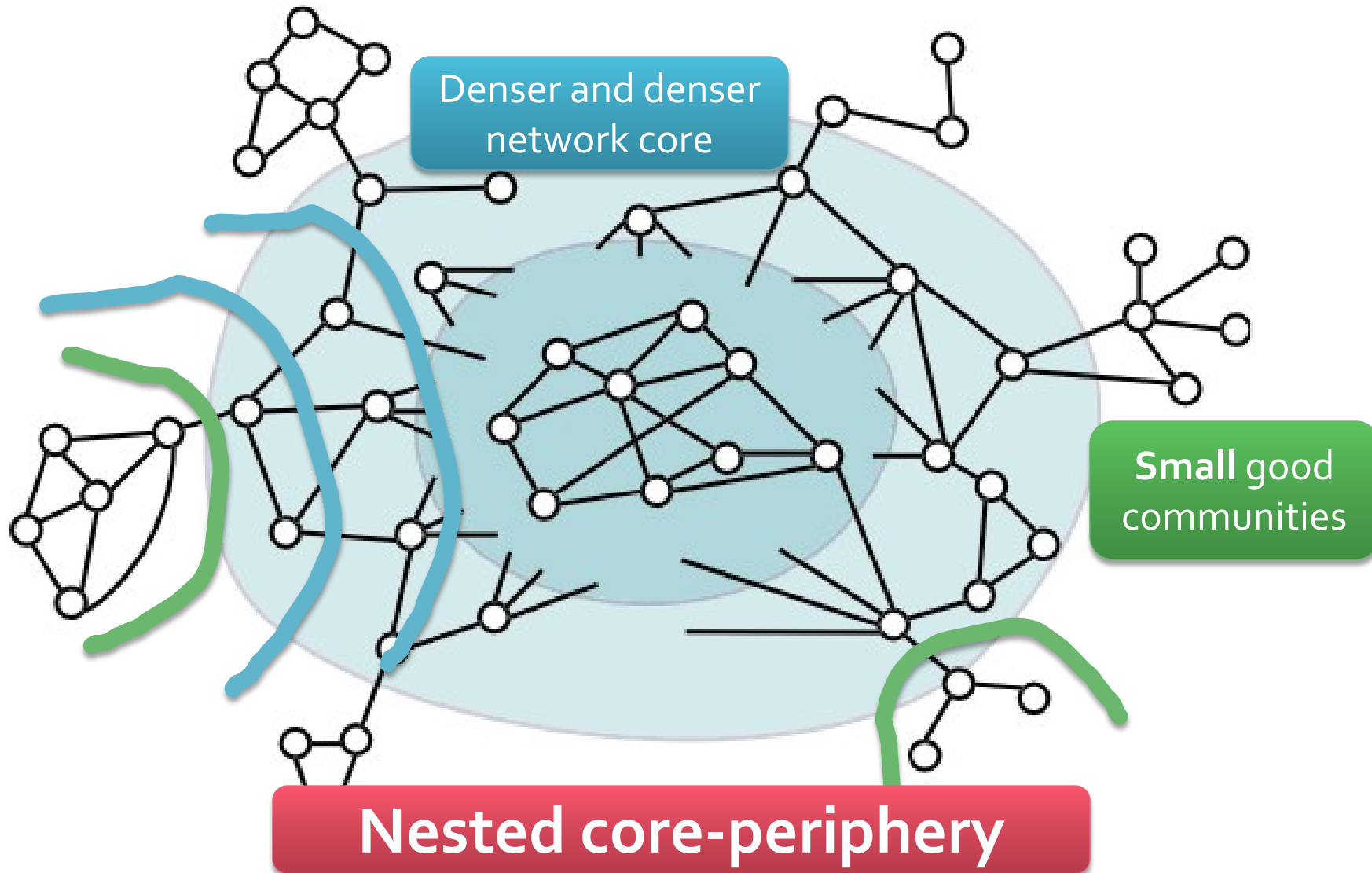
<http://cs224w.stanford.edu>



Recap: Network Community Profile

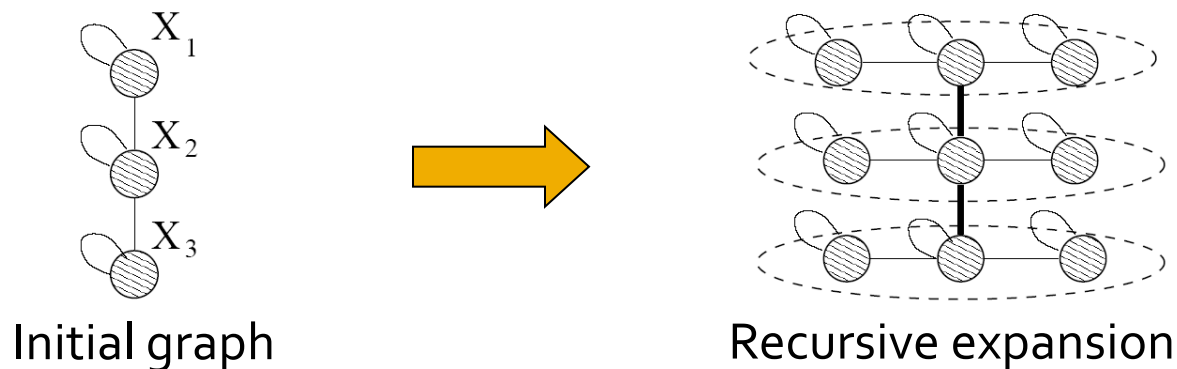


Explanation: Nested Core-Periphery



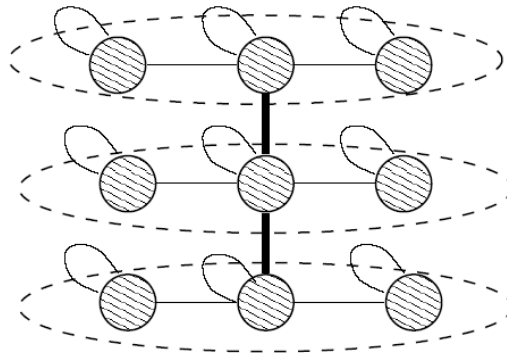
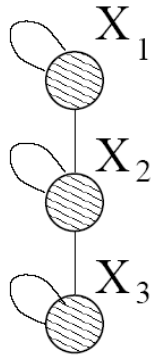
Idea: Recursive Graph Generation

- **Intuition: Self-similarity**
 - Object is similar to a part of itself (i.e. the whole has the same shape as one or more of the parts)
- Mimic recursive graph / community growth



- **Kronecker Product** is a way of generating self-similar matrices

Kronecker: Graph Growth



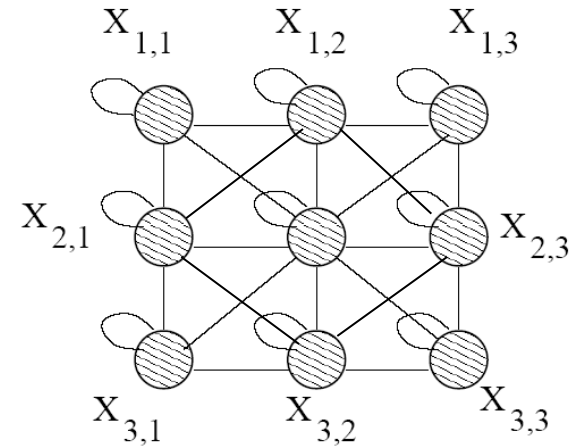
Intermediate stage

1	1	0
1	1	1
0	1	1

(3x3)

K_1

Initiator graph



K_1	K_1	0
K_1	K_1	K_1
0	K_1	K_1

(9x9)

$$K_2 = K_1 \otimes K_1$$

After the growth phase

Kronecker Product: Definition

- **Kronecker product** of matrices A and B is given by

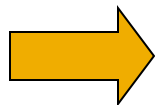
$$\underset{N \times M}{\mathbf{C}} = \underset{N \times M}{\mathbf{A}} \otimes \underset{K \times L}{\mathbf{B}} \doteq \underset{N * K \times M * L}{\begin{pmatrix} a_{1,1}\mathbf{B} & a_{1,2}\mathbf{B} & \dots & a_{1,m}\mathbf{B} \\ a_{2,1}\mathbf{B} & a_{2,2}\mathbf{B} & \dots & a_{2,m}\mathbf{B} \\ \vdots & \vdots & \ddots & \vdots \\ a_{n,1}\mathbf{B} & a_{n,2}\mathbf{B} & \dots & a_{n,m}\mathbf{B} \end{pmatrix}}$$

- Define a Kronecker product of two graphs as a Kronecker product of their **adjacency matrices**

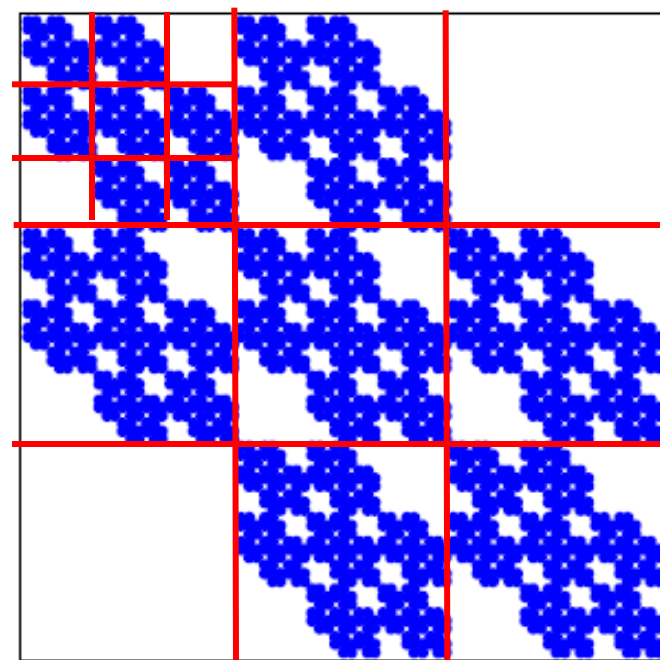
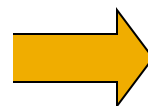
Kronecker Product: Graph

- Continuing multiplying with K_1 we obtain K_4 and so on ...

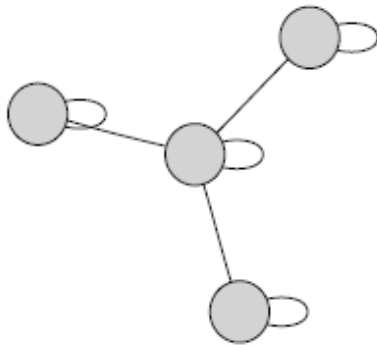
1	1	0
1	1	1
0	1	1

 K_1
 3×3


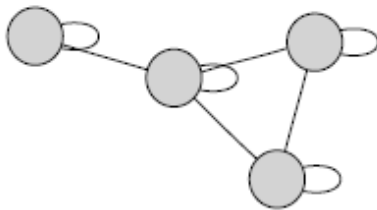
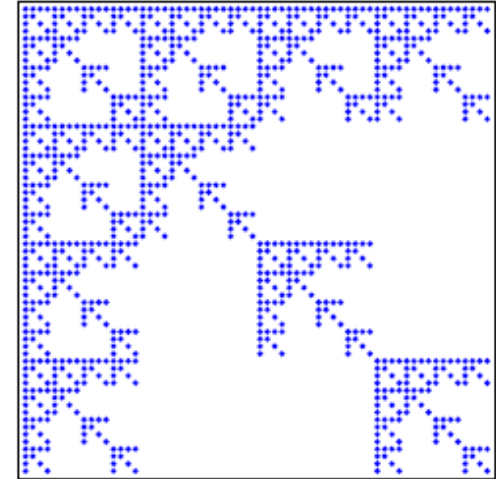
K_1	K_1	0
K_1	K_1	K_1
0	K_1	K_1

 $K_2 = K_1 \otimes K_1$
 9×9

 K_4 adjacency matrix

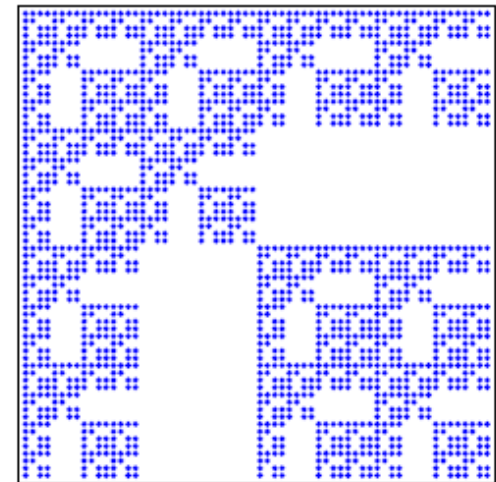
Kronecker Initiator Matrices



1	1	1	1
1	1	0	0
1	0	1	0
1	0	0	1



1	1	1	1
1	1	0	0
1	0	1	1
1	0	1	1

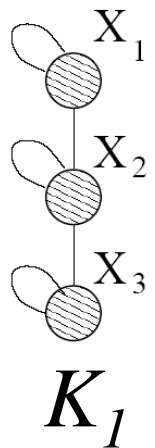
Initiator K_1 K_1 adjacency matrix K_3 adjacency matrix

Kronecker Graphs

- **Kronecker graph:** a growing sequence of graphs by iterating the Kronecker product

$$K_1^{[k]} = K_k = \underbrace{K_1 \otimes K_1 \otimes \dots \otimes K_1}_{k \text{ times}} = K_{k-1} \otimes K_1$$

1	1	0
1	1	1
0	1	1



- **Note:** One can easily use multiple initiator matrices (K_1', K_1'', K_1''') (even of different sizes)

Kronecker Graphs

$$K_1^{[k]} = K_k = \underbrace{K_1 \otimes K_1 \otimes \dots K_1}_{k \text{ times}} = K_{k-1} \otimes K_1$$

- For K_1 on N_1 nodes and E_1 edges K_k (k^{th} Kronecker power of K_1) has:
 - N_1^k nodes
 - E_1^k edges
- We get **densification power-law**:
 - $E(t) \propto N(t)^a$, What is a ?
 - $a = \frac{\log(E(t))}{\log(N(t))} = \frac{\log(E_1)}{\log(N_1)}$

1	1	0
1	1	1
0	1	1

K_1

Properties of Kronecker graphs

- **Kronecker graphs have many properties found in real networks:**
 - **Properties of static networks**
 - Power-Law like Degree Distribution
 - Power-Law eigenvalue and eigenvector distribution
 - Small Diameter
 - **Properties of dynamic networks**
 - Densification Power Law
 - Shrinking/Stabilizing Diameter

Constant Diameter

- Observation: Edges in Kronecker graphs:

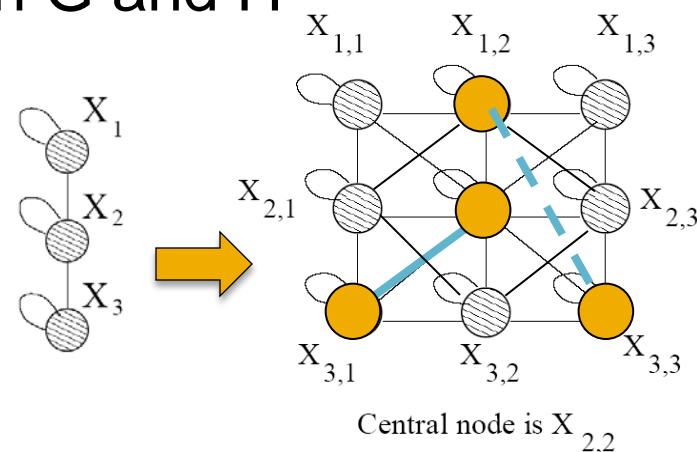
$$\text{Edge } (X_{ij}, X_{kl}) \in G \otimes H$$

$$\text{iff } (X_i, X_k) \in G \text{ and } (X_j, X_l) \in H$$

where X are appropriate nodes in G and H

- **Why?**

- An entry in matrix $G \otimes H$ is a multiplication of entries in G and H .



1	1	0
1	1	1
0	1	1

Central node is $X_{2,2}$

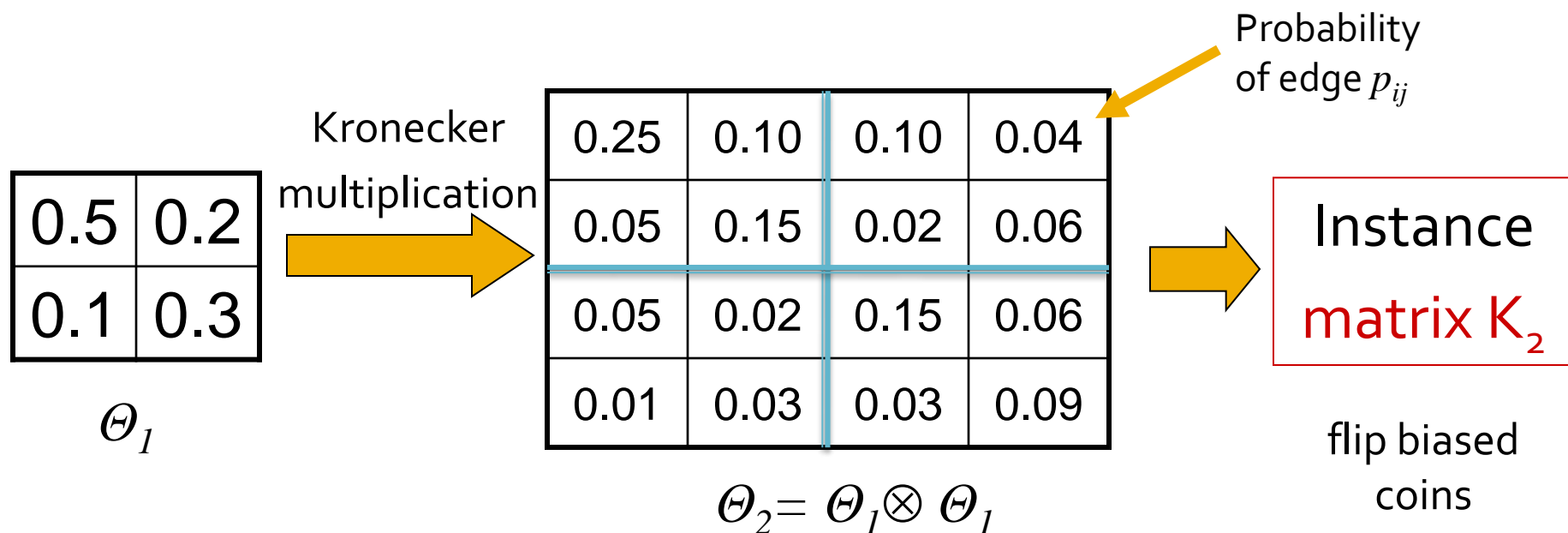
1	1	0	1	1	0	0	1	1	0
1	1	1	1	1	1	1	1	1	1
0	1	1	0	1	1	1	1	1	1
1	1	0	1	1	0	1	1	0	1
1	1	1	1	1	1	1	1	1	1
0	1	1	0	1	1	0	1	1	1
0	1	1	0	1	1	1	1	0	1
1	1	0	1	1	0	1	1	0	1
1	1	1	1	1	1	1	1	1	1
0	1	1	0	1	1	0	1	1	1

Constant Diameter

- Theorem: **Constant diameter**: If G, H have diameter d then $G \otimes H$ has diameter d
- **What is distance between nodes u, v in $G \otimes H$?**
 - Let $u=[a,b], v=[a',b']$ (using notation from last slide) **then** edge (u,v) in $G \otimes H$ iff $(a,a') \in G$ and $(b,b') \in H$
 - So, path a to a' in G is less d steps: $a_1, a_2, a_3, \dots, a_d$
 - And path b to b' in H is less d steps: $b_1, b_2, b_3, \dots, b_d$
 - **Then:** edge $([a_1, b_1], [a_2, b_2])$ is in $G \otimes H$
 - So it takes $< d$ steps to get from u to v in $G \otimes H$
- **Consequence:**
 - If K_l has diameter d then graph K_k also has diameter d

Stochastic Kronecker Graphs

- Create $N_I \times N_I$ probability matrix Θ_1
- Compute the k^{th} Kronecker power Θ_k
- For each entry p_{uv} of Θ_k include an edge (u,v) in K_k with probability p_{uv}



Stochastic Kronecker Graphs

What is known about Stochastic Kronecker?

■ **Undirected** Kronecker graph model with:

■ **Connected**, if:

- $b+c > 1$

$$\Theta_1 = \begin{array}{|c|c|} \hline a & b \\ \hline b & c \\ \hline \end{array}$$

■ **Connected component of size $\Theta(n)$** , if:

- $(a+b)(b+c) > 1$

$$a > b > c$$

■ **Constant diameter**, if:

- $b+c > 1$

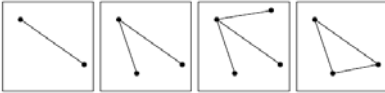
■ **Not searchable** by a decentralized algorithm

Estimating Kronecker graphs

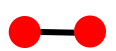
- Given a real network G

Want to estimate the initiator matrix: $\Theta_1 = \begin{bmatrix} a & b \\ b & d \end{bmatrix}$

- Method of moments [Gleich&Owen '09]

- Compare counts of  and solve the system of equations

- For every of the 4 subgraphs, we get an equation:

- $2 E[\# \text{ ] = (a+2b+c)^k - (a+c)^k$ where $k = \log_2(N)$

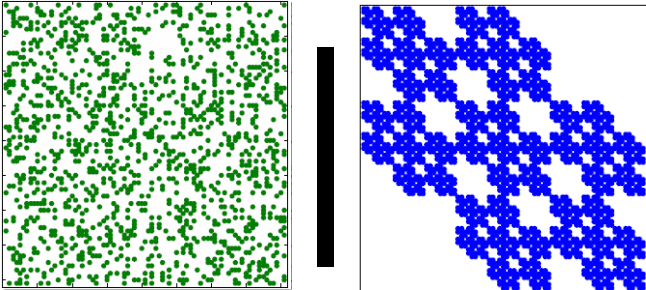
- $2 E[\# \text{ ] = \dots$

- ...

- Now solve the system of equations by trying all possible values (a,b,c)

Kronecker graphs: Estimation

■ Maximum Likelihood Estimation

$$\arg \max_{\Theta_1} P(\text{[Green Graph]} \mid \text{[Blue Graph]} \leftarrow \text{Kronecker}(\Theta_1))$$


■ Naïve estimation takes $O(N!N^2)$:

■ $N!$ for different node labelings:

■ **Solution:** Metropolis sampling: $N! \rightarrow$ *(big) const*

■ N^2 for traversing graph adjacency matrix

■ **Solution:** Kronecker product ($E \ll N^2$): $N^2 \rightarrow E$

■ Do gradient descent

$$\Theta_1 = \begin{bmatrix} a & b \\ c & d \end{bmatrix}$$

Parameter Estimation: Approach

KronFit: Maximum likelihood estimation

- Given real graph G
- Find Kronecker initiator graph Θ (i.e.,

a	b
c	d

) which

$$\arg \max_{\Theta} P(G | \Theta)$$

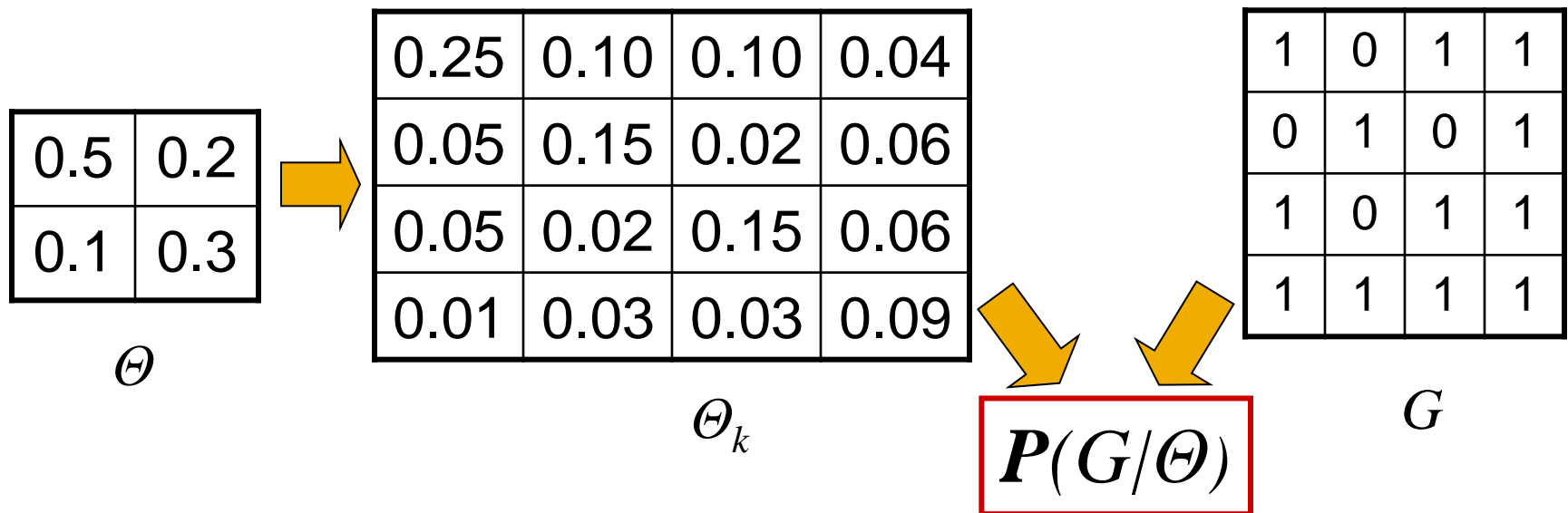
- We then need to (efficiently) calculate

$$P(G | \Theta)$$

- **And maximize over Θ**
(e.g., using gradient descent)

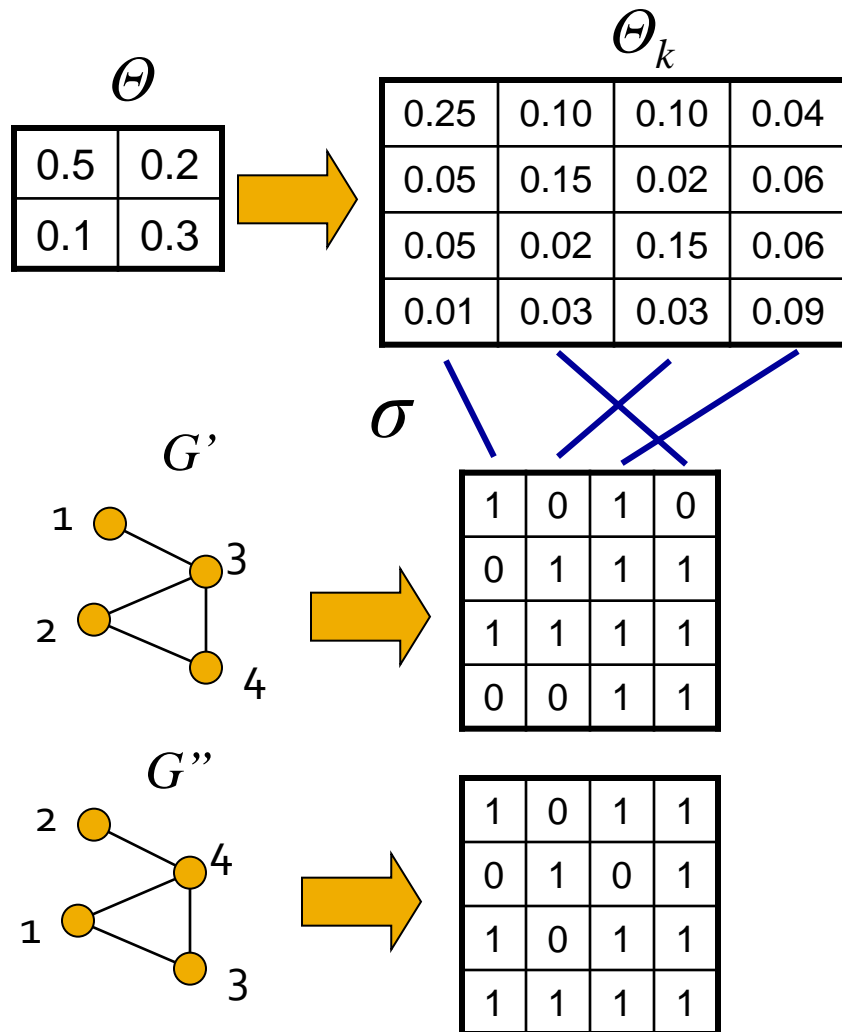
KronFit: Likelihood $P(G|\Theta)$

- Given a graph G and Kronecker matrix Θ we calculate probability that Θ generated G $P(G/\Theta)$



$$P(G | \Theta) = \prod_{(u,v) \in G} \Theta_k[u,v] \prod_{(u,v) \notin G} (1 - \Theta_k[u,v])$$

Challenge 1: Node Correspondence



$$P(G'|\Theta) = P(G''|\Theta)$$

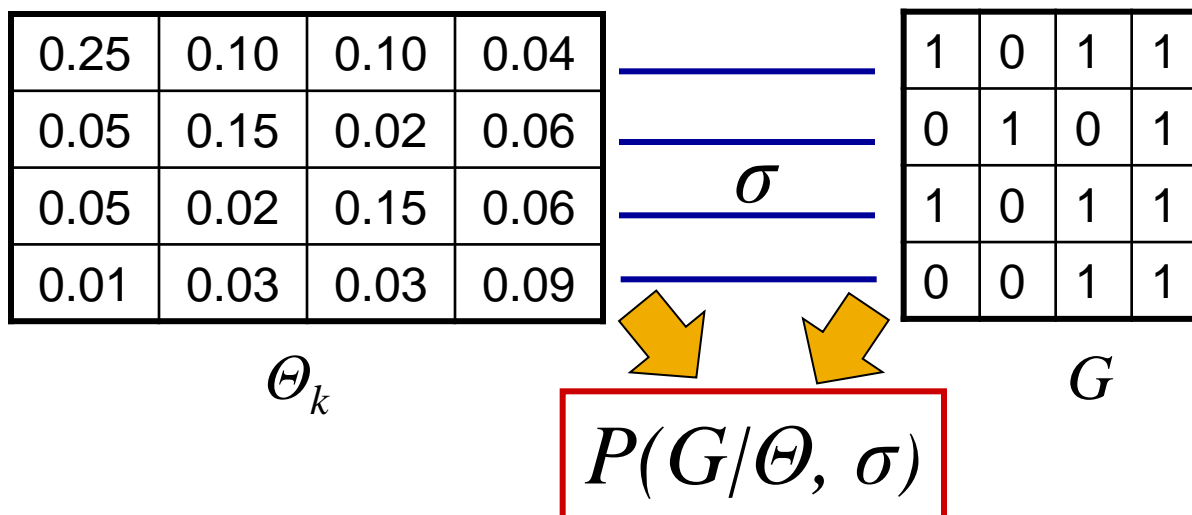
- Nodes are **unlabeled**
 - Graphs G' and G'' should have the same probability $P(G'|\Theta) = P(G''|\Theta)$
 - One needs to consider all node correspondences σ
- $$P(G|\Theta) = \sum_{\sigma} P(G|\Theta, \sigma)P(\sigma)$$
- All correspondences are a priori equally likely
 - There are **$O(N!)$** correspondences

Challenge 2: Calculating $P(G|\Theta, \sigma)$

- Assume that we solved the node correspondence problem
- Calculating

$$P(G | \Theta) = \prod_{(u,v) \in G} \Theta_k[u, v] \prod_{(u,v) \notin G} (1 - \Theta_k[u, v])$$

- Takes $O(N^2)$ time



Experiments: real networks

■ Experimental setup

- Given real graph G
- Gradient descent from random initial point
- Obtain estimated parameters Θ
- Generate synthetic graph K using Θ
- Compare properties of graphs G and K

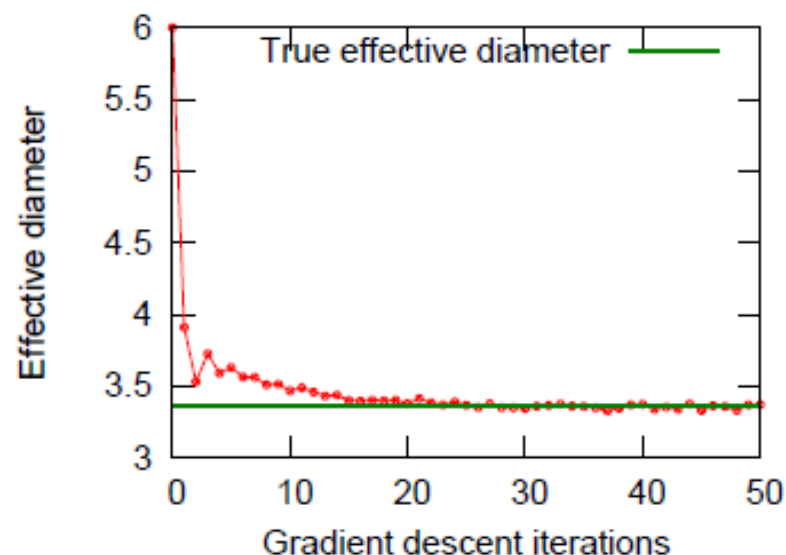
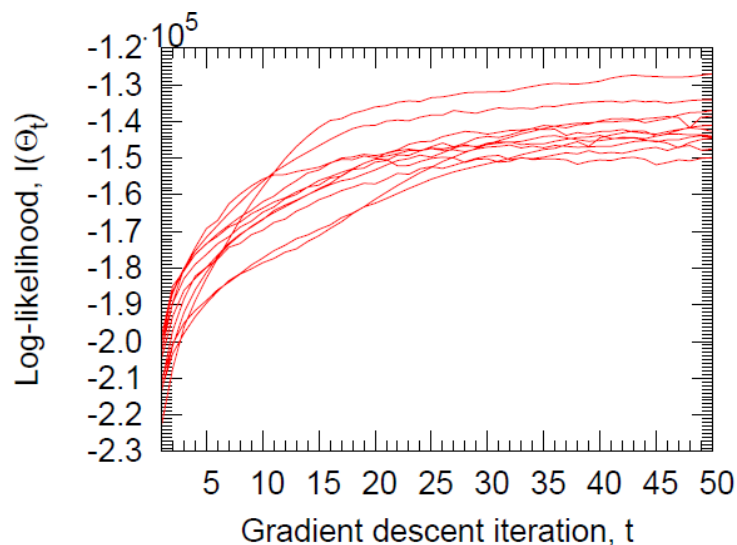
$$\Theta = \begin{array}{|c|c|} \hline a & b \\ \hline c & d \\ \hline \end{array}$$

■ Note:

- We do not fit the graph properties themselves
- We fit the likelihood and then compare the properties

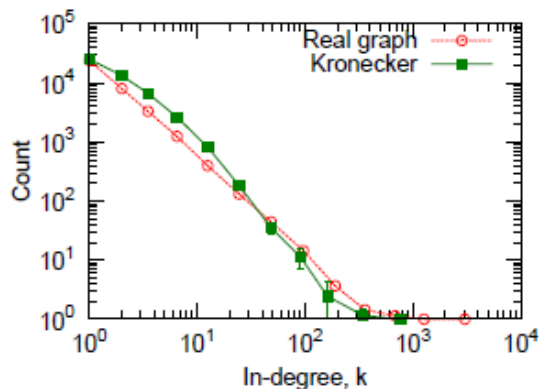
Convergence of fitting

- Can gradient descent recover true parameters?
 - Generate a graph from random parameters
 - Start at random point and use gradient descent
 - We recover true parameters 98% of the times

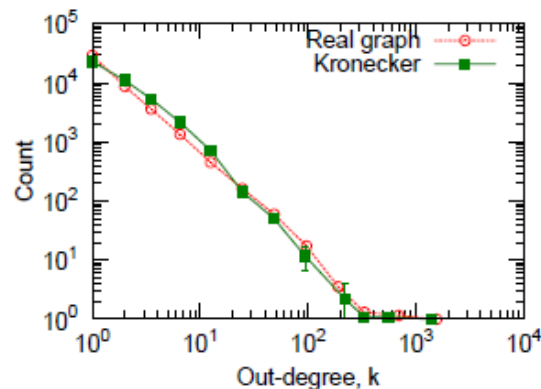


Estimation: Epinions (n=76k, m=510k)

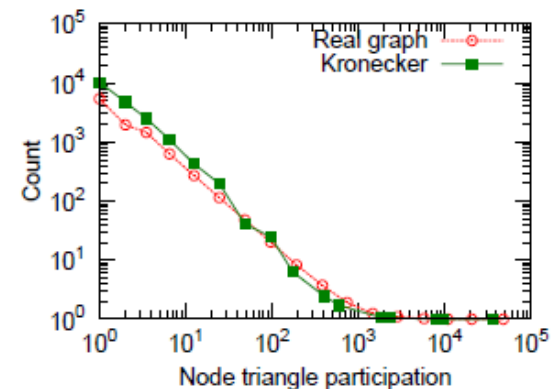
- Real and Kronecker are very close:

$$\Theta_1 = \begin{bmatrix} 0.99 & 0.54 \\ 0.49 & 0.13 \end{bmatrix}$$


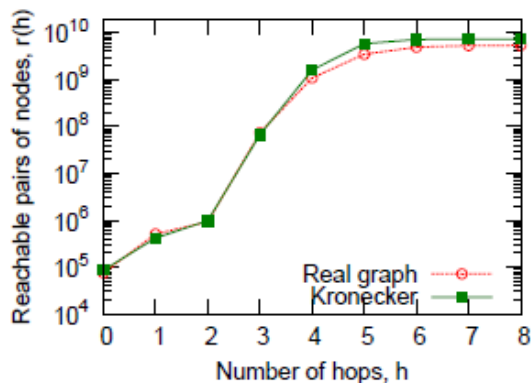
(a) In-Degree



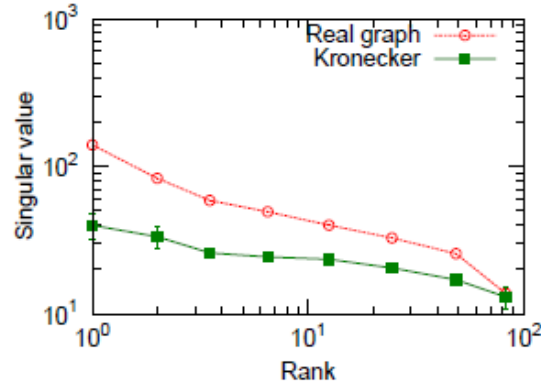
(b) Out-degree



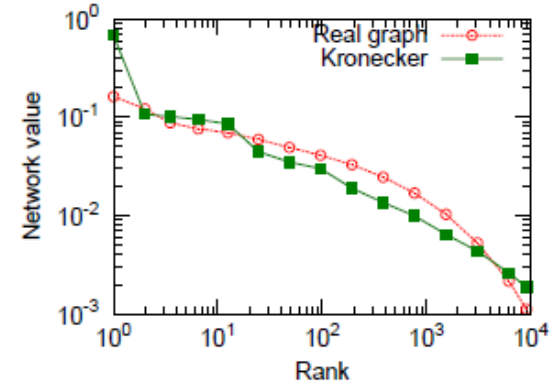
(c) Triangle participation



(d) Hop plot



(e) Scree plot

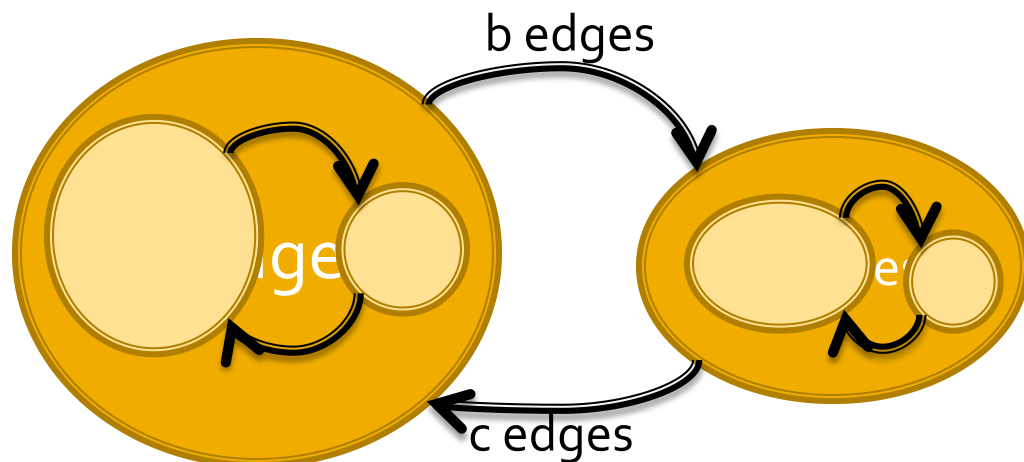


(f) "Network" value

Kronecker & Network Structure

- What do estimated parameters tell us about the network structure?

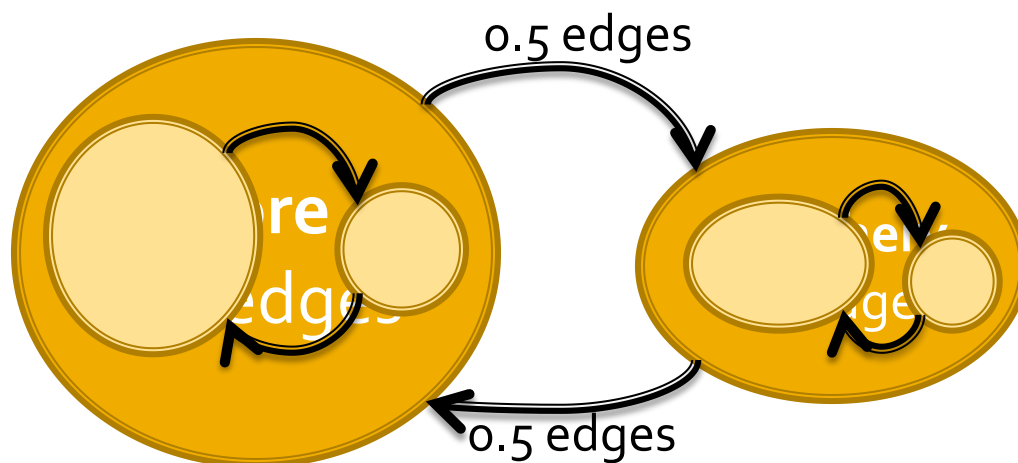
$$\Theta = \begin{bmatrix} a & b \\ c & d \end{bmatrix}$$



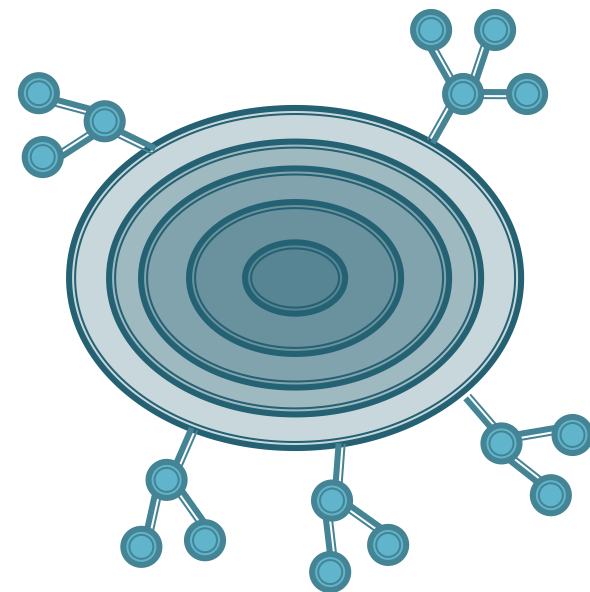
Kronecker & Network structure

- What do estimated parameters tell us about the network structure?

$$\Theta = \begin{bmatrix} 0.9 & 0.5 \\ 0.5 & 0.1 \end{bmatrix}$$

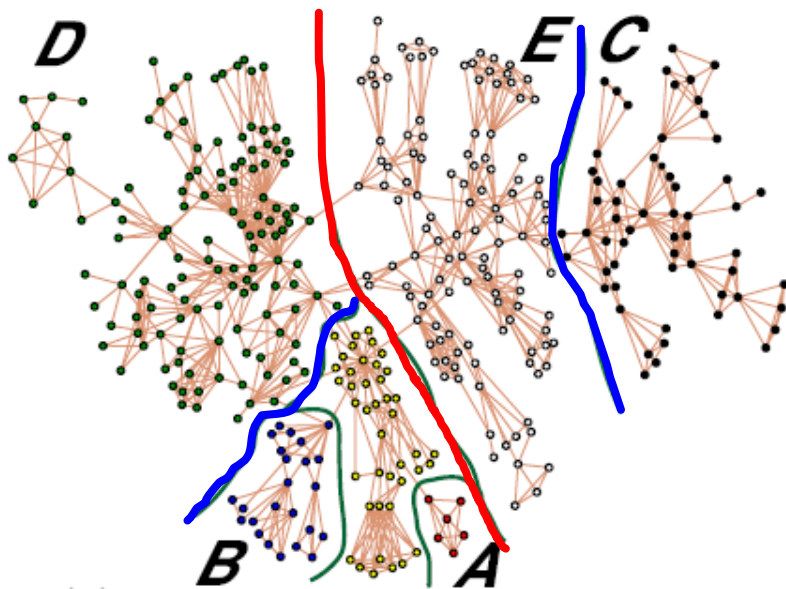


Nested Core-periphery

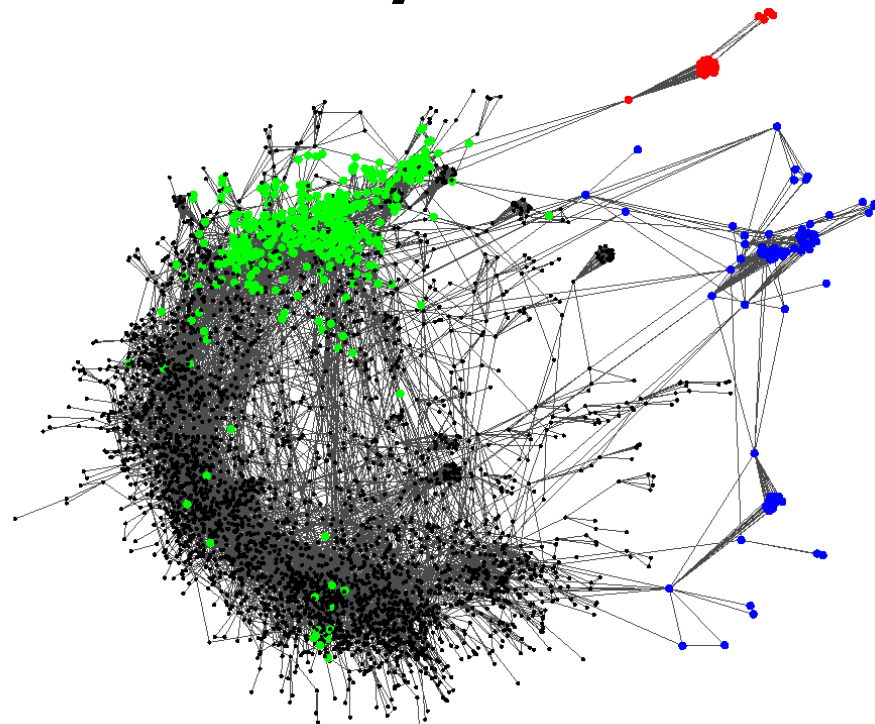


Small vs. Large Networks

- Small and large networks are very different:



$$\Theta = \begin{bmatrix} 0.99 & 0.17 \\ 0.17 & 0.82 \end{bmatrix}$$

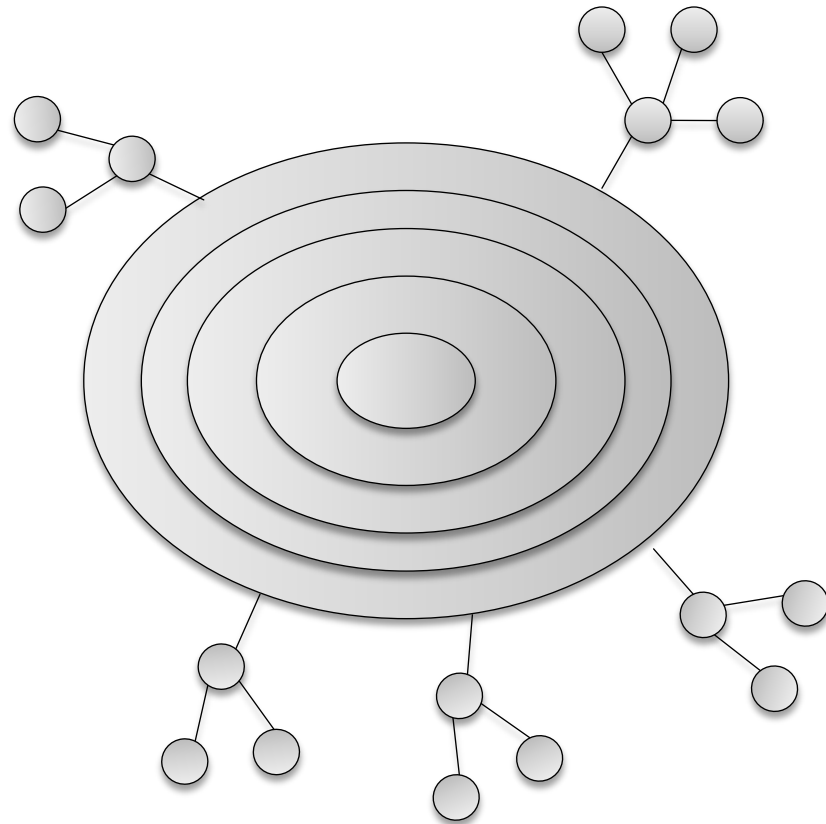


$$\Theta = \begin{bmatrix} 0.99 & 0.54 \\ 0.49 & 0.13 \end{bmatrix}$$

Implications (1)

Large scale network structure:

- Large networks are **different** from **small networks** and **manifolds**
- **Nested Core-periphery**
 - Recursive onion-like structure of the network where each layer decomposes into a core and periphery



Implications (2)

- Remember the SKG theorems:

- Connected, if $b+c > 1$:

- $0.55+0.15 > 1$. **No!**

- Giant component, if $(a+b) \cdot (b+c) > 1$:

- $(0.99+0.55) \cdot (0.55+0.15) > 1$. **Yes!**

- Real graphs are in the parameter region analogous to the giant component of an extremely sparse G_{np}

 $\Theta =$

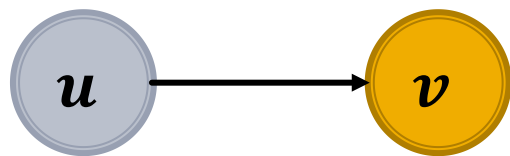
0.99	0.55
0.55	0.15



A Different Model: MAG Model

Nodes with Attributes

- Each node has a set of *categorycal attributes*
 - Example:
 - Gender: Male, Female
 - Home country: US, Canada, Russia, etc.
- How do node attributes influence link formation?



u is friends with v

		v 's gender	
u 's gender	u \ v	FEMALE	MALE
	FEMALE	0.3	0.6
	MALE	0.6	0.2

Link probability

Link-Affinity Matrix

- Let the values of the ***i -th attribute*** for node u and v be **$a_i(u)$** and **$a_i(v)$**
 - $a_i(u)$ and $a_i(v)$ can take values $\{0, \dots, d_i - 1\}$
- **Question:** How can we capture the influence of the attributes on link formation?
 - ***Attribute matrix Θ***

	$a_i(v) = 0$	$a_i(v) = 1$
$a_i(u) = 0$	$\Theta[0, 0]$	$\Theta[0, 1]$
$a_i(u) = 1$	$\Theta[1, 0]$	$\Theta[1, 1]$

$$P(u, v) = \Theta[a_i(u), a_i(v)]$$

Each entry of the attribute matrix captures the ***probability of a link*** between two nodes associated with the attributes of them

Approach: Great flexibility

- **Flexibility** in the network structure:

- **Homophily** : love of the *same*

- e.g., political parties, hobbies

0.9	0.1
0.1	0.8

- **Heterophily** : love of the *opposite*

- e.g., genders

0.2	0.9
0.9	0.1

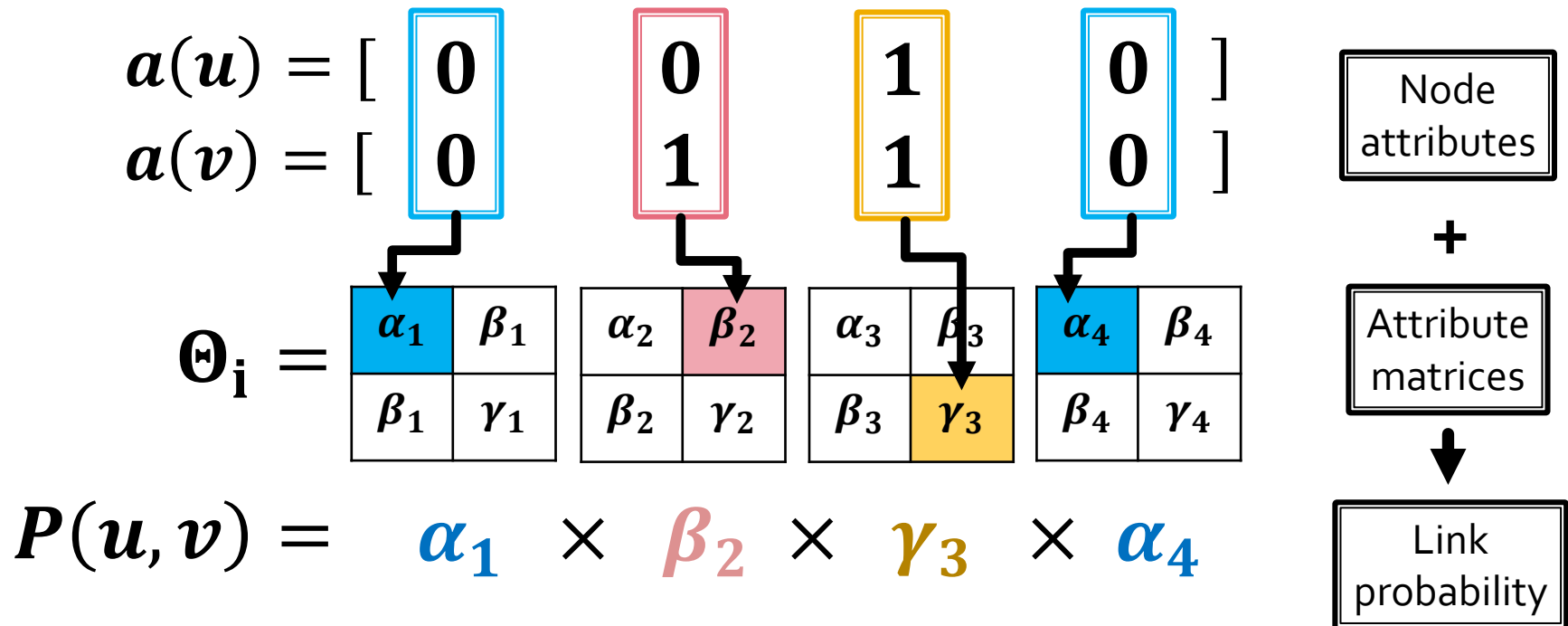
- **Core-periphery** : love of the *core*

- e.g. extrovert personalities

0.9	0.5
0.5	0.2

Combining attributes

- How do we combine the effects of multiple attributes?
 - Multiply the probabilities** from all attributes



Multiplicative Attribute Graph

- **Multiplicative Attribute Graph $M(n, l, \vec{a}, \vec{\Theta})$:**
 - A network contains **n nodes**
 - Each node has **l categorical attributes**
 - $a_i(u)$ represents the **i -th attribute of node u**
 - Each attribute $a_i(\cdot)$ is linked to a **$d_i \times d_i$ attribute link-affinity matrix Θ_i**
 - Edge probability between nodes u and v

$$P(u, v) = \prod_{i=1}^l \Theta_i[a_i(u), a_i(v)]$$

Connection to Kronecker Graphs

- Initiator matrix K_1 acts like an **affinity matrix**
- Probability of a link** between nodes u, v :

$$P(u, v) = \prod_{i=1}^k K_1(A_u(i), A_v(i))$$

$$K_1 = \begin{array}{cc|c} & 0 & 1 & \\ \hline & a & b & 0 \\ \hline & c & d & 1 \end{array}$$



	v_1	v_2	v_3	v_4
v_1	$a \cdot a$	$a \cdot b$	$b \cdot a$	$b \cdot b$
v_2	$a \cdot c$	$a \cdot d$	$b \cdot c$	$b \cdot d$
v_3	$c \cdot a$	$c \cdot b$	$d \cdot a$	$d \cdot b$
v_4	$c \cdot c$	$c \cdot d$	$d \cdot c$	$d \cdot d$

=

	v_1	v_2	v_3	v_4
v_1	a b a b			
v_2	c d c d			
v_3	a b a b			
v_4	c d c d			

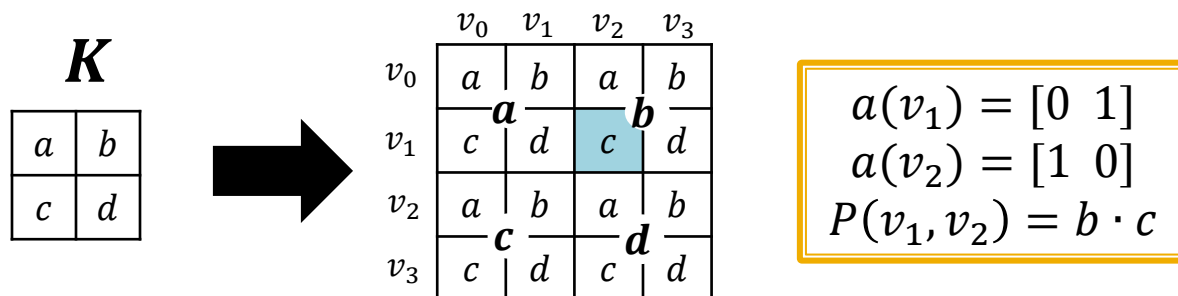
$$K_2 = K_1 \otimes K_1$$

$$v_2 = (0, 1)$$

$$v_3 = (1, 0) \quad P(v_2, v_3) = b \cdot c$$

Connection to Kronecker Graphs

- Each node in a Kronecker graph has a node id (e.g. $0, \dots, 2^l - 1$)
- A binary representation of node id is its attribute vector in a MAG model
- Then, the (stochastic) adjacency matrices of two models are equivalent
- **Example:**



Feature vector view: Question

■ 2 ingredients of Kronecker model:

- (1) Each of 2^k nodes has a unique binary vector of length k
 - Node id expressed binary is the vector
- (2) The initiator matrix K

	v_1	v_2	v_3	v_4
v_1	a	b	a	b
v_2	c	d	c	d
v_3	a	b	a	b
v_4	c	d	c	d

The table illustrates a 4x4 grid of features. The columns are labeled v_1, v_2, v_3, v_4 and the rows are labeled v_1, v_2, v_3, v_4 . The cells contain the letters 'a', 'b', 'c', and 'd'. The cell at row v_1 , column v_2 contains a bold 'a'. The cell at row v_1 , column v_3 contains a bold 'b'. The cell at row v_3 , column v_1 contains a bold 'c'. The cell at row v_3 , column v_3 contains a bold 'd'. The cell at row v_2 , column v_3 is shaded gray.

■ Question:

- What if ingredient (1) is dropped?
 - i.e., do we need high variability of feature vectors?

Comparison: Adjacency matrices

■ Adjacency matrices:

