

Generative Models for Complex Network Structure

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• **what is structure?**

- generative models for complex networks
 - general form
 - types models
 - opportunities and challenges
- weighted stochastic block models
 - a parable about thresholding
 - checking our models
 - learning from data (approximately)

what is structure?

- makes data different from noise
 - makes a network different from a random graph

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- helps us compress the data
 - describe the network succinctly
 - capture most relevant patterns

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- helps us compress the data
 - describe the network succinctly
 - capture most relevant patterns
- helps us generalize,
from data we've seen to data we haven't seen:
 - i. from one part of network to another
 - ii. from one network to others of same type
 - iii. from small scale to large scale (coarse-grained structure)
 - iv. from past to future (dynamics)

statistical inference

- imagine graph G is drawn from an ensemble or **generative model**: a probability distribution $P(G | \theta)$ with parameters θ
- θ can be continuous or discrete; represents structure of graph

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-

- if θ is partly known, constrain inference and determine the rest
- if G is partly known, infer θ and use $P(G | \theta)$ to generate the rest
- if model is good fit (application dependent), we can generate synthetic graphs structurally similar to G
- if part of G has low probability under model, flag as possible anomaly

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generative models for complex networks

general form

$$P(G \mid \theta) = \prod_{i < j} P(A_{ij} \mid \theta)$$

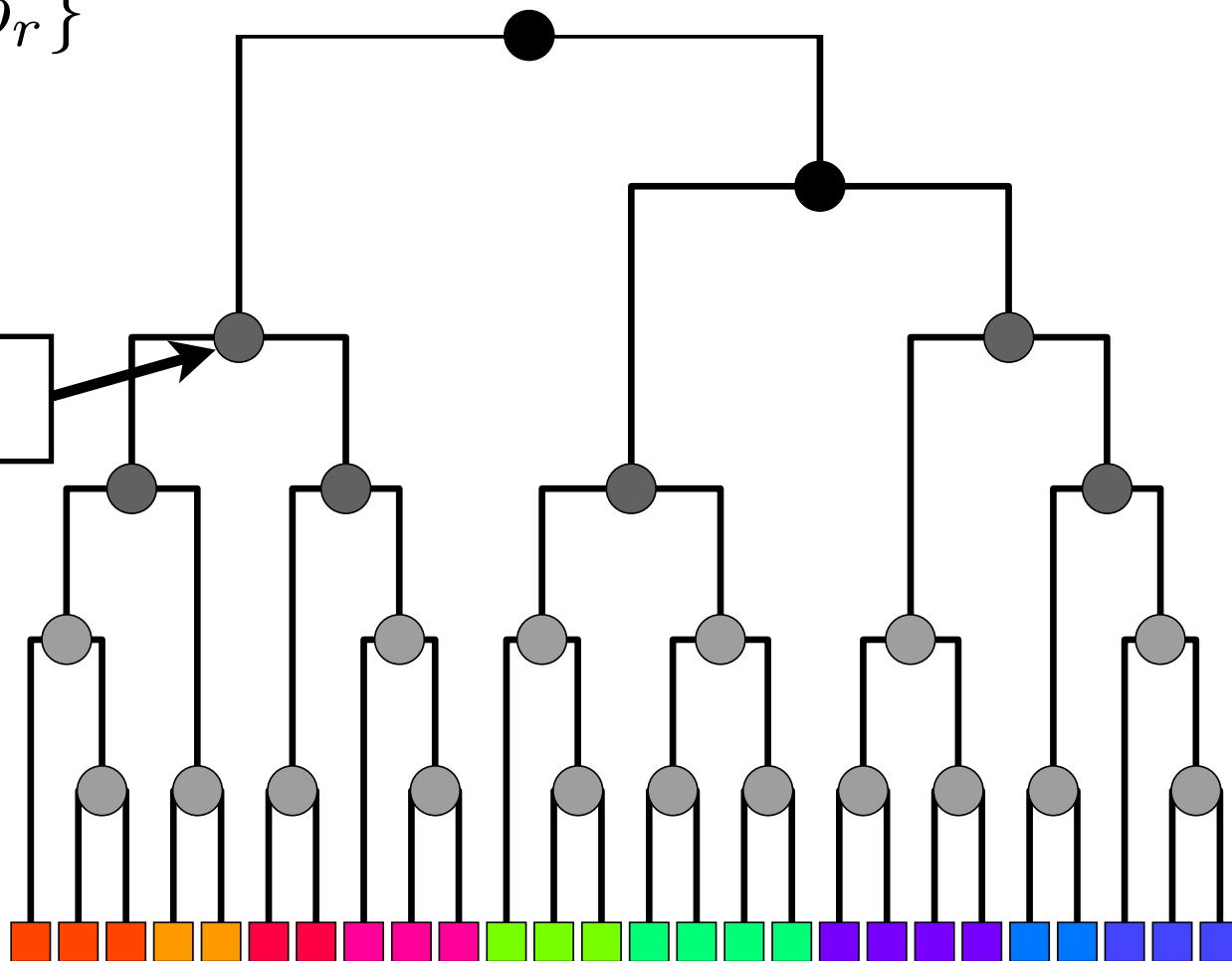
assumptions about “structure” go into $P(A_{ij} \mid \theta)$

consistency $\lim_{n \rightarrow \infty} \Pr(\hat{\theta} \neq \theta) = 0$

requires that edges be conditionally independent [Shalizi, Rinaldo 2011]

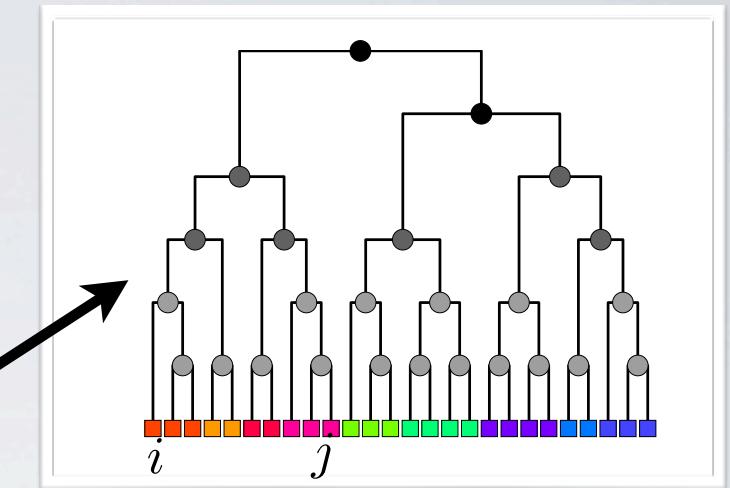
$\mathcal{D}, \{p_r\}$

probability p_r

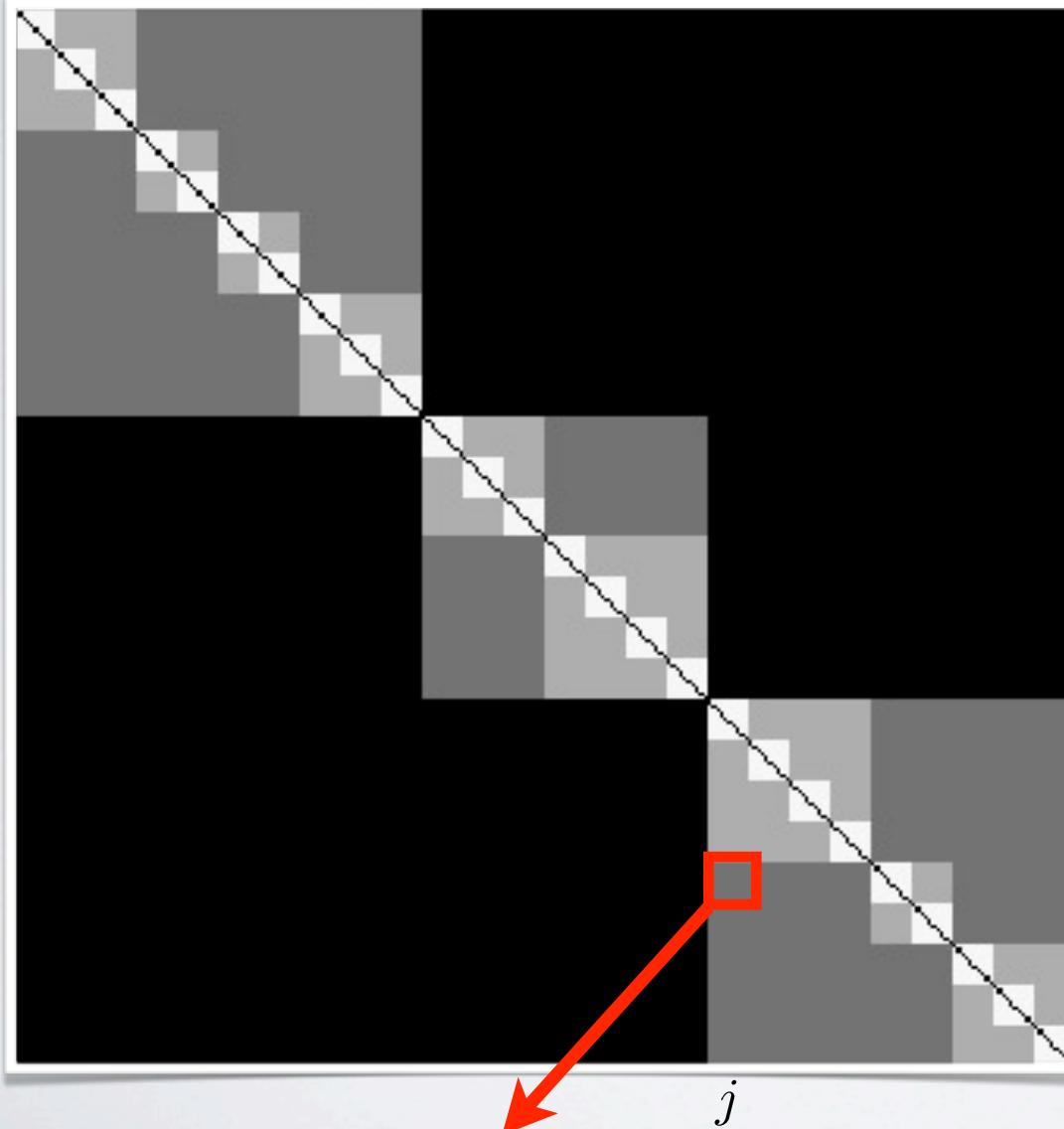


assortative modules

model



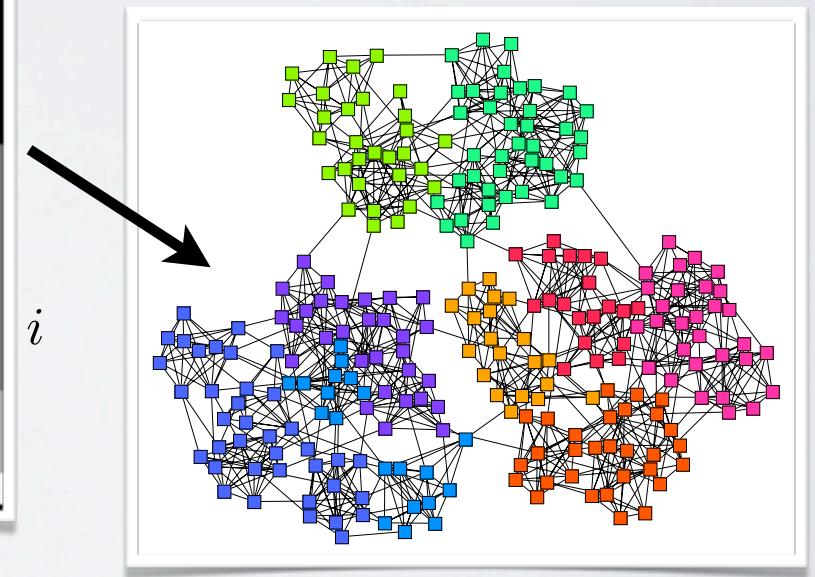
hierarchical random graph



$$\Pr(i, j \text{ connected}) = p_r$$

$$= p_{(\text{lowest common ancestor of } i, j)}$$

instance

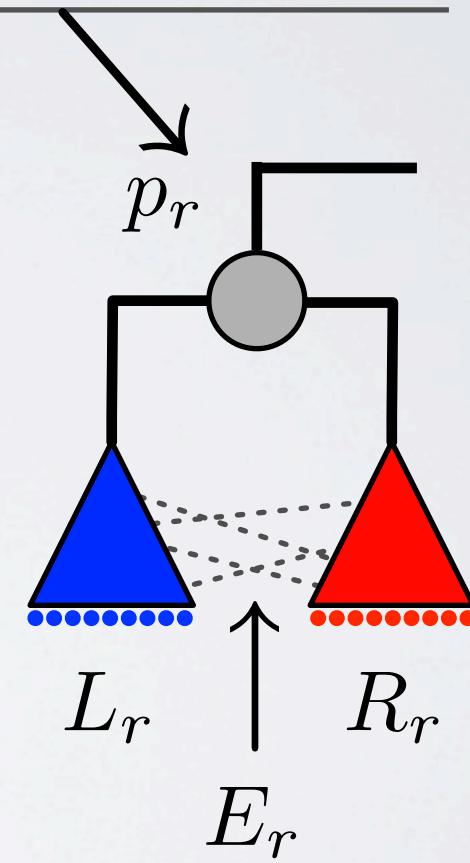


$$\mathcal{L}(\mathcal{D}, \{p_r\}) = \prod_r p_r^{E_r} (1 - p_r)^{L_r R_r - E_r}$$

L_r = number nodes in left subtree

R_r = number nodes in right subtree

E_r = number edges with r as lowest common ancestor



classes of generative models

- stochastic block models

k types of vertices, $P(A_{ij} | z_i, z_j)$ depends only on types of i, j
originally invented by sociologists [Holland, Laskey, Leinhardt 1983]

many, many flavors, including

mixed-membership SBM [Airoldi, Blei, Feinberg, Xing 2008]

hierarchical SBM [Clauset, Moore, Newman 2006,2008]

restricted hierarchical SBM [Leskovec, Chakrabarti, Kleinberg, Faloutsos 2005]

infinite relational model [Kemp, Tenenbaum, Griffiths, Yamada, Ueda 2006]

restricted SBM [Hofman, Wiggins 2008]

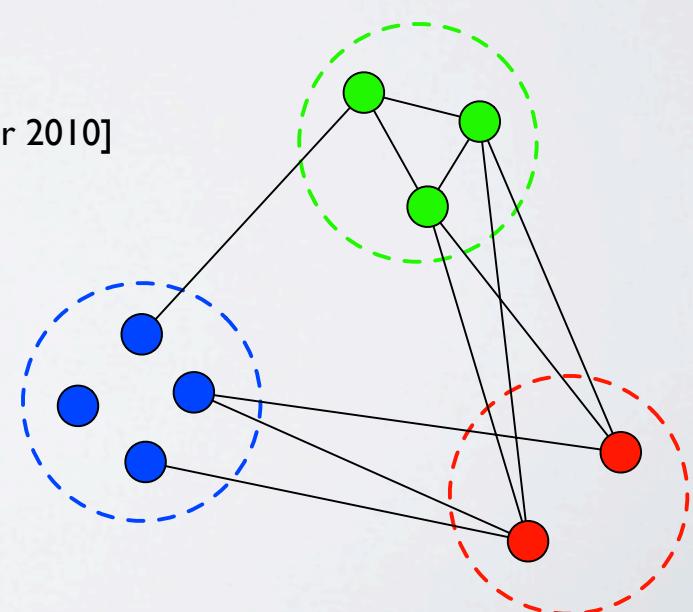
degree-corrected SBM [Karrer, Newman 2011]

SBM + topic models [Ball, Karrer, Newman 2011]

SBM + vertex covariates [Mariadassou, Robin, Vacher 2010]

SBM + edge weights [Aicher, Jacobs, Clauset 2013]

+ many others



classes of generative models

- latent space models

nodes live in a latent space, $P(A_{ij} | f(x_i, x_j))$ depends only on vertex-vertex proximity

many, many flavors, including

logistic function on vertex features [Hoff, Raftery, Handcock 2002]

social status / ranking [Ball, Newman 2013]

nonparametric metadata relations [Kim, Hughes, Sudderth 2012]

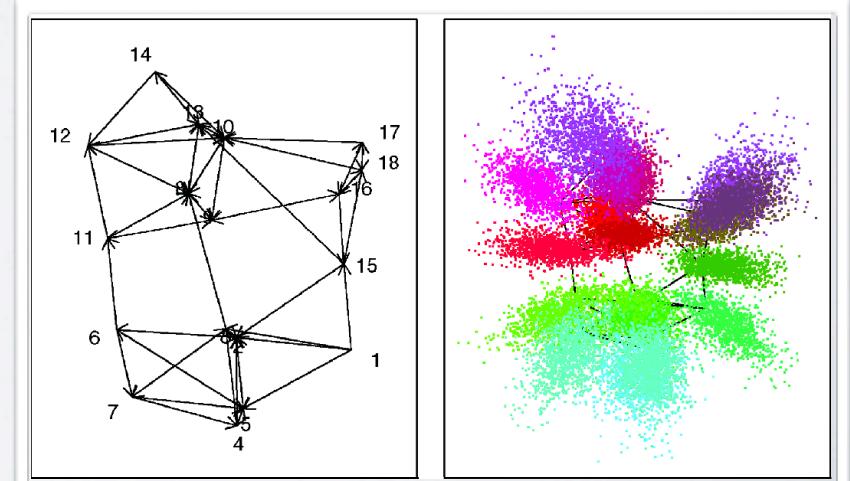
multiple attribute graphs [Kim, Leskovec 2010]

nonparametric latent feature model [Miller, Griffiths, Jordan 2009]

infinite multiple memberships [Morup, Schmidt, Hansen 2011]

ecological niche model [Williams, Anandanadesan, Purves 2010]

hyperbolic latent spaces [Boguna, Papadopoulos, Krioukov 2010]



opportunities and challenges

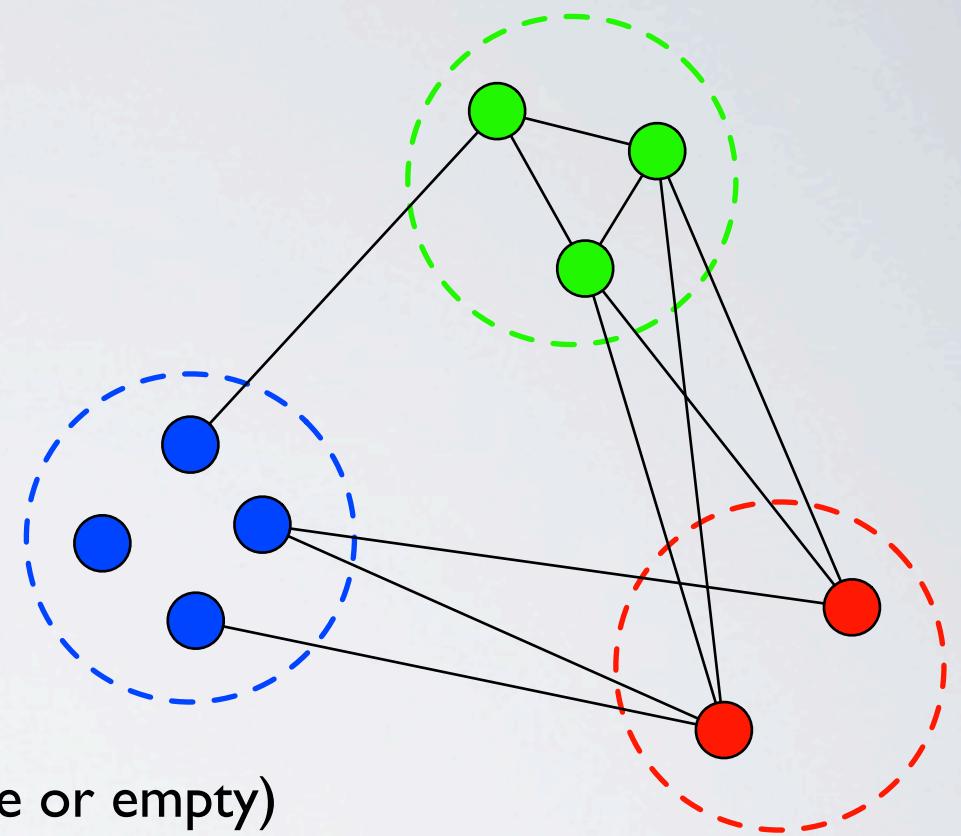
- richly annotated data
 - edge weights, node attributes, time, etc.
 - = new classes of generative models
- generalize from $n = 1$ to ensemble
 - useful for modeling checking, simulating other processes, etc.
- many familiar techniques
 - frequentist and Bayesian frameworks
 - makes probabilistic statements about observations, models
 - predicting missing links \approx leave- k -out cross validation
 - approximate inference techniques (EM, VB, BP, etc.)
 - sampling techniques (MCMC, Gibbs, etc.)
- learn from partial or noisy data
 - extrapolation, interpolation, hidden data, missing data

opportunities and challenges

- only two classes of models
 - stochastic block models
 - latent space models
- bootstrap / resampling for network data
 - critical missing piece
 - depends on what is independent in the data
- model comparison
 - naive AIC, BIC, marginalization, LRT can be wrong for networks
 - what is goal of modeling: realistic representation or accurate prediction?
- model assessment / checking?
 - how do we know a model has done well? what do we check?
- what is v -fold cross-validation for networks?
 - Omit n^2/v edges? Omit n/v nodes? What?

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- **weighted stochastic block models**
 - a parable about thresholding
 - learning from data (approximately)
 - checking our models

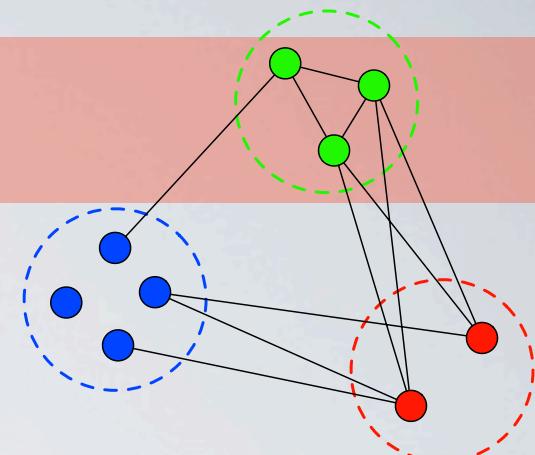
stochastic block models



functional groups, not just clumps

- social “communities” (large, small, dense or empty)
- social: leaders and followers
- word adjacencies: adjectives and nouns
- economics: suppliers and customers

classic stochastic block model



nodes have discrete attributes

each vertex i has type $t_i \in \{1, \dots, k\}$

$k \times k$ matrix p of connection probabilities

if $t_i = r$ and $t_j = s$, edge $(i \rightarrow j)$ exists with probability p_{rs}

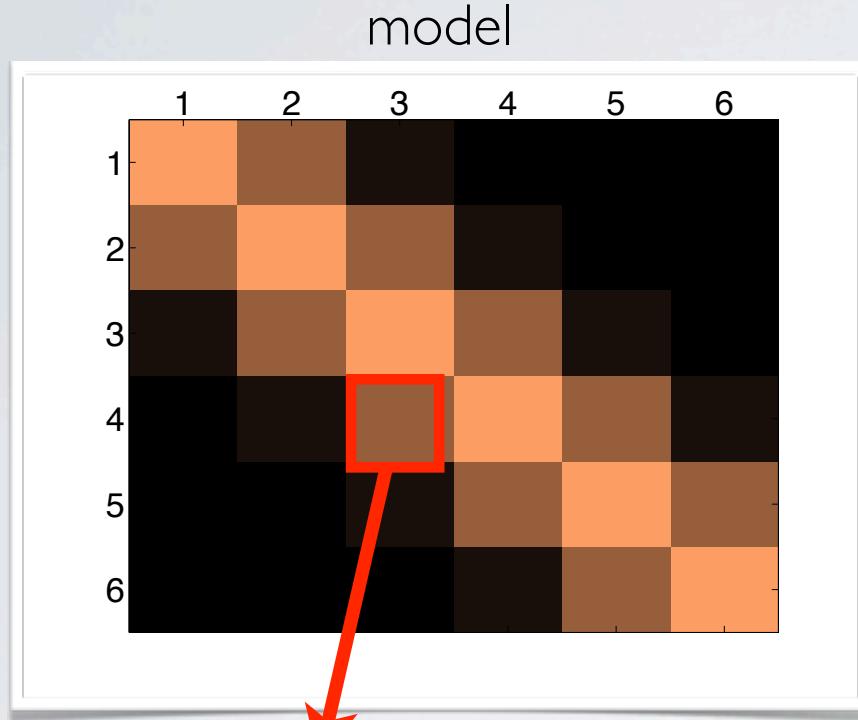
p not necessarily symmetric, and we do not assume $p_{rr} > p_{rs}$

given some G , we want to simultaneously

label nodes (infer type assignment $t : V \rightarrow \{1, \dots, k\}$)

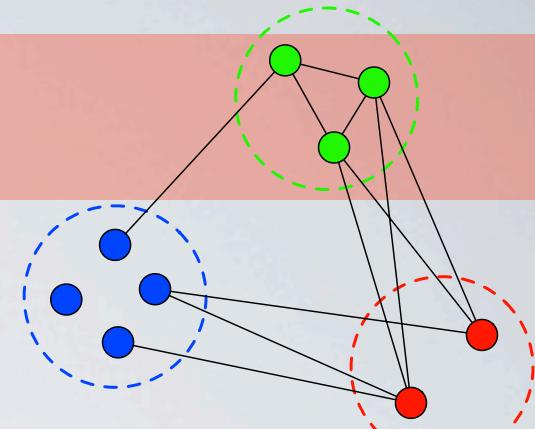
learn the latent matrix p

classic stochastic block model

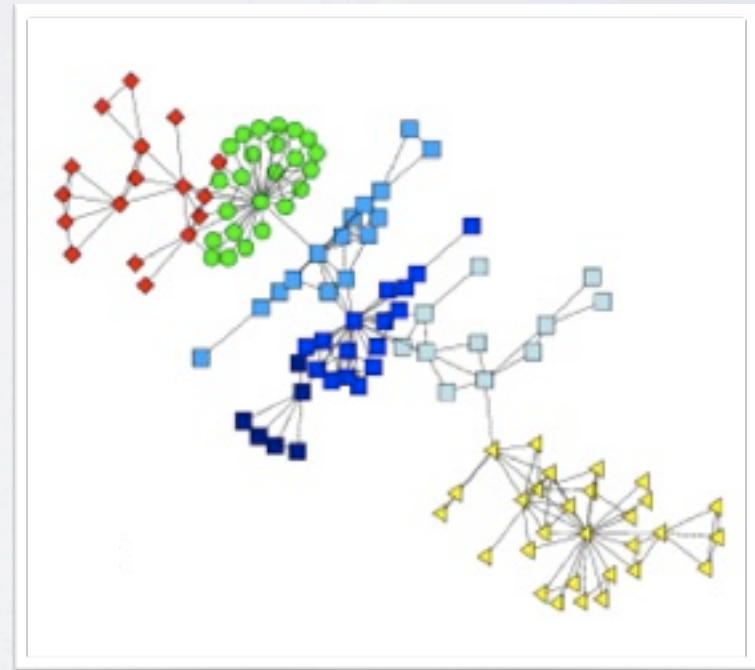


$$\Pr(\text{node in } i \text{ connected to node in } j) = p_{i,j}$$

assortative modules



instance

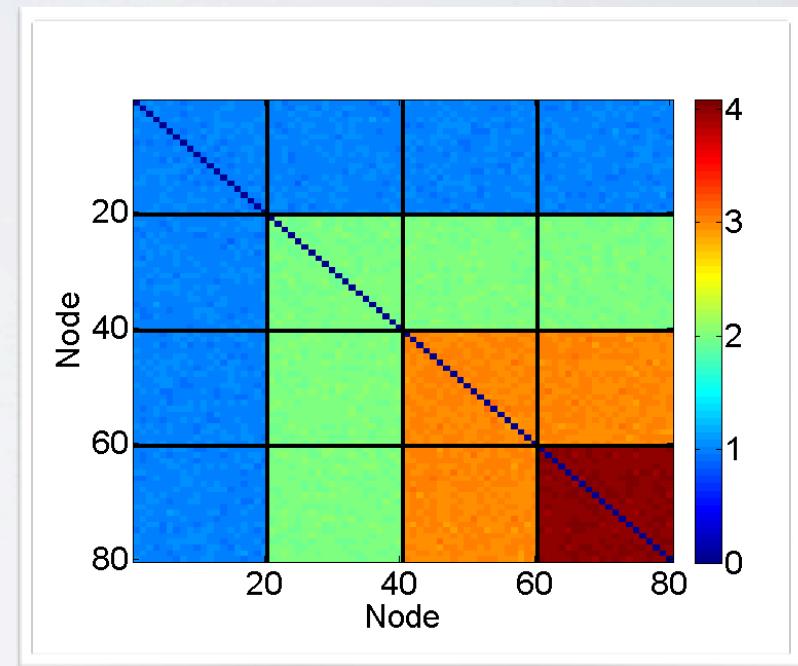
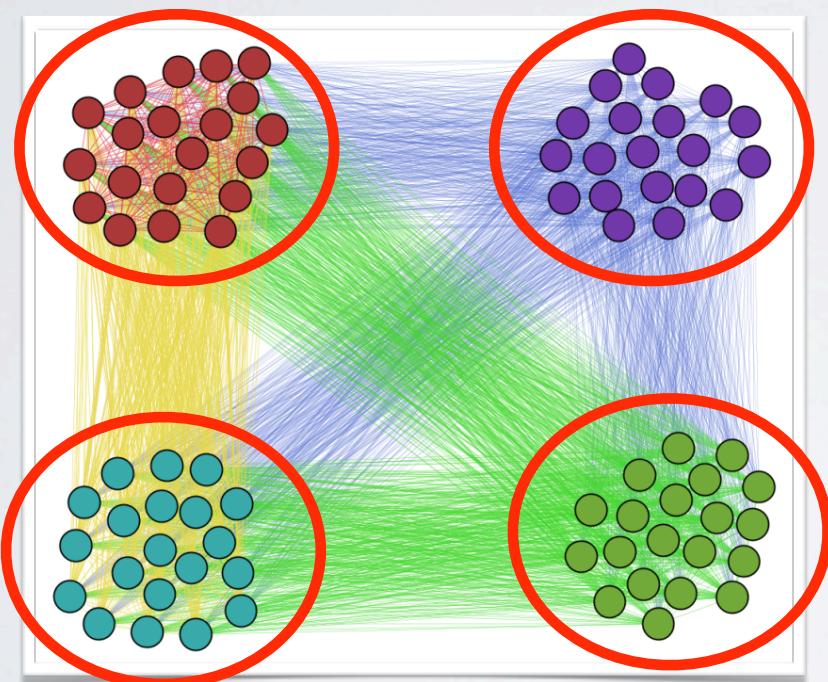


likelihood

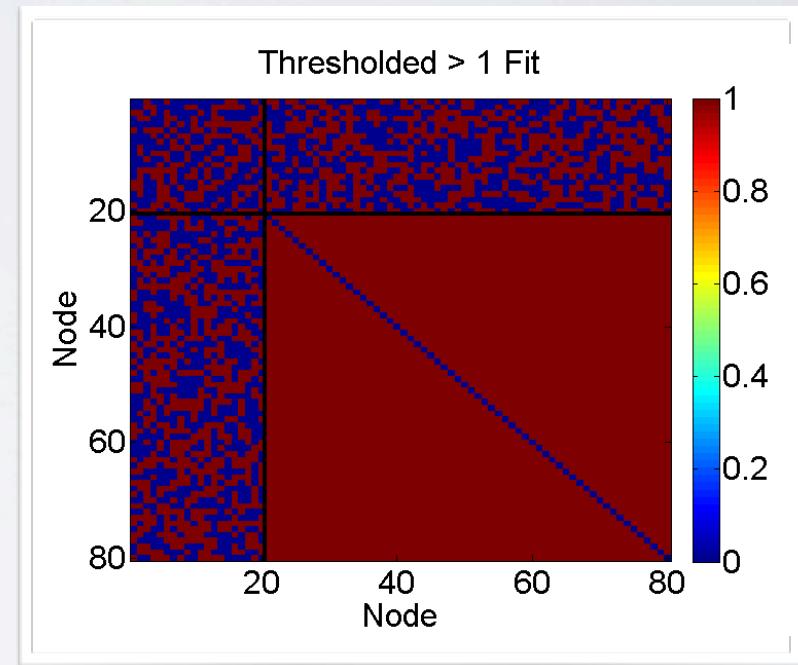
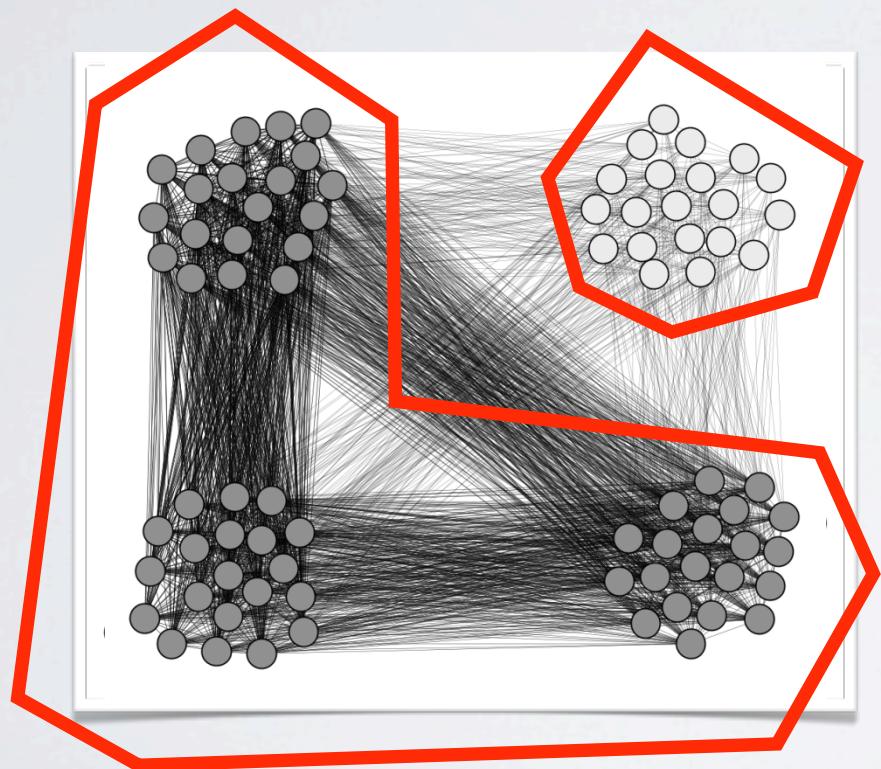
$$P(G | t, \theta) = \prod_{(i,j) \in E} p_{t_i, t_j} \prod_{(i,j) \notin E} (1 - p_{t_i, t_j})$$

thresholding edge weights

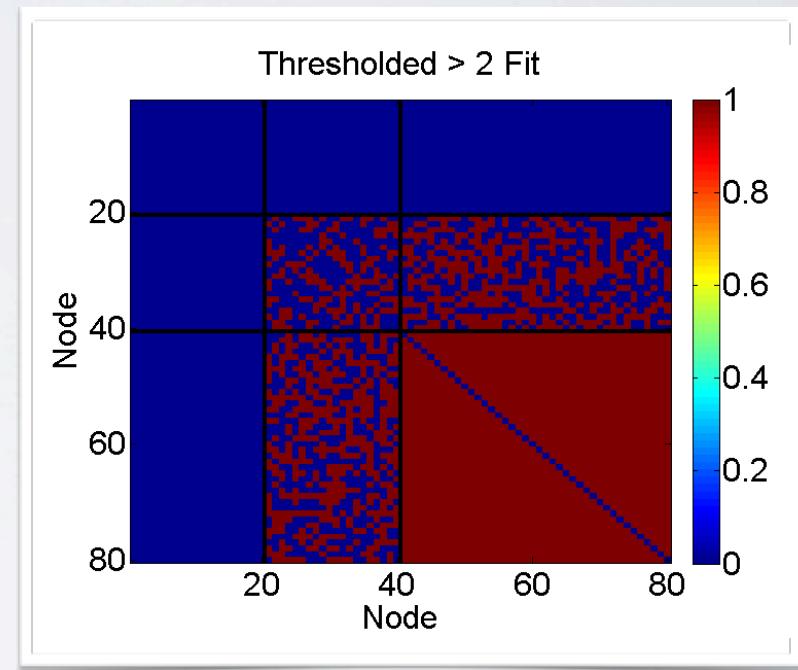
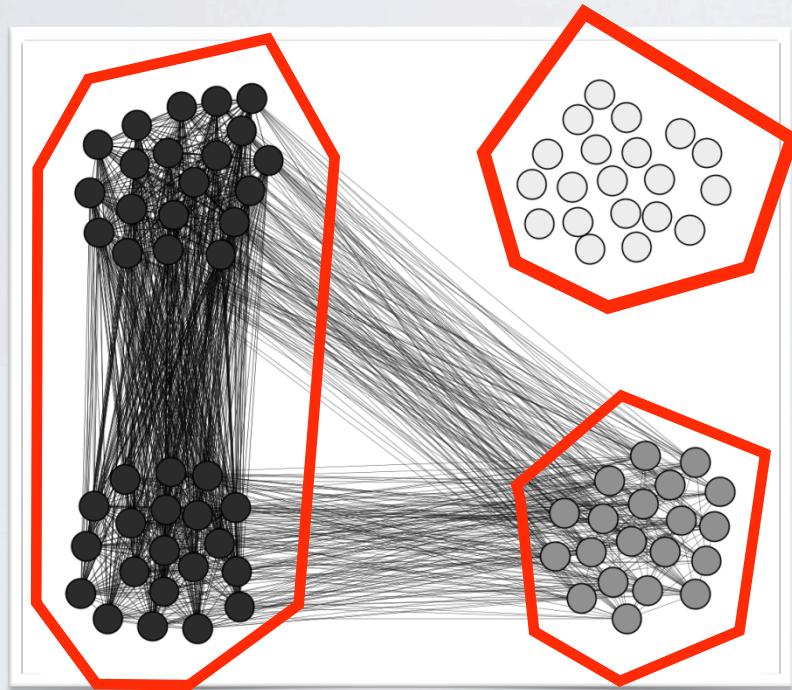
- 4 groups
- edge weights $\sim N(\mu_i, \sigma^2)$ with $\mu_1 < \mu_2 < \mu_3 < \mu_4$
- what threshold t should we choose? $t = 1, 2, 3, 4$



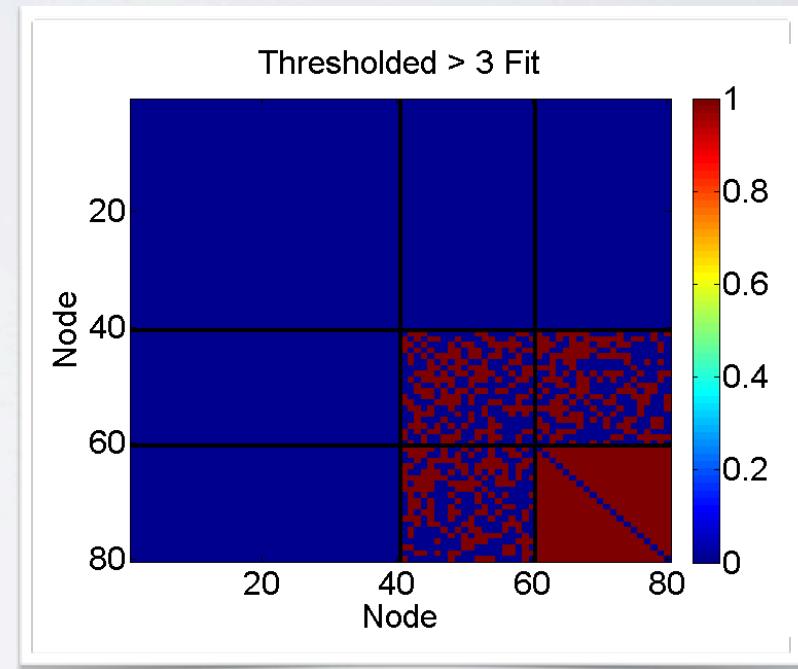
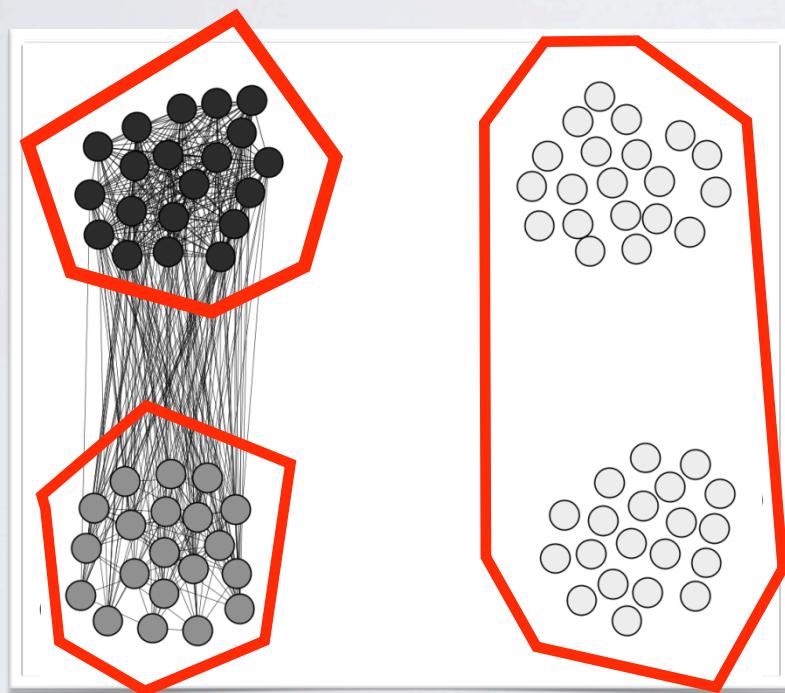
- 4 groups
- edge weights $\sim N(\mu_i, \sigma^2)$ with $\mu_1 < \mu_2 < \mu_3 < \mu_4$
- set threshold $t \leq 1$, fit SBM



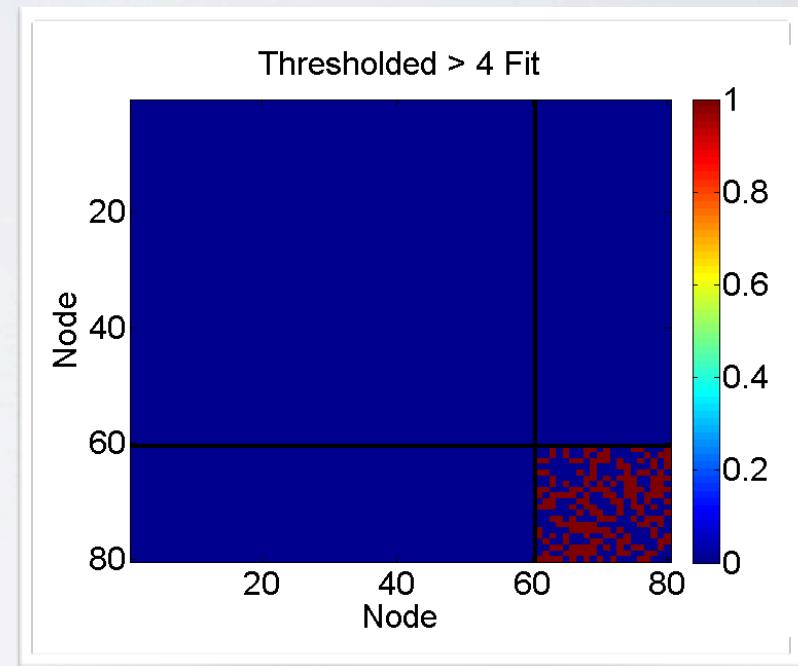
- 4 groups
- edge weights $\sim N(\mu_i, \sigma^2)$ with $\mu_1 < \mu_2 < \mu_3 < \mu_4$
- set threshold $t = 2$, fit SBM



- 4 groups
- edge weights $\sim N(\mu_i, \sigma^2)$ with $\mu_1 < \mu_2 < \mu_3 < \mu_4$
- set threshold $t = 3$, fit SBM



- 4 groups
- edge weights $\sim N(\mu_i, \sigma^2)$ with $\mu_1 < \mu_2 < \mu_3 < \mu_4$
- set threshold $t \geq 4$, fit SBM



weighted stochastic block model

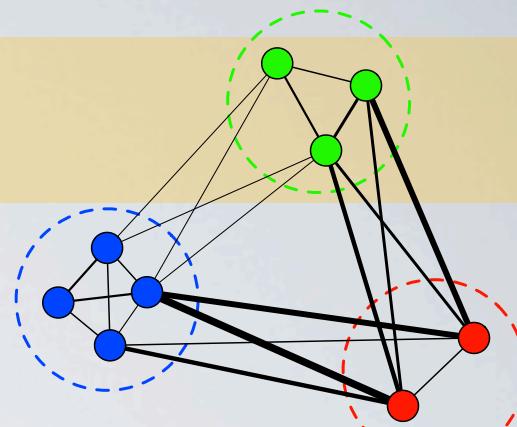
adding auxiliary information:

each edge has weight $w(i, j)$

$$\text{let } w(i, j) \sim f(x|\theta)$$

$$= h(x) \exp(T(x) \cdot \eta(\theta))$$

- covers all exponential-family type distributions:
 - bernoulli, binomial (classic SBM), multinomial
 - poisson, beta
 - exponential, power law, gamma
 - normal, log-normal, multivariate normal



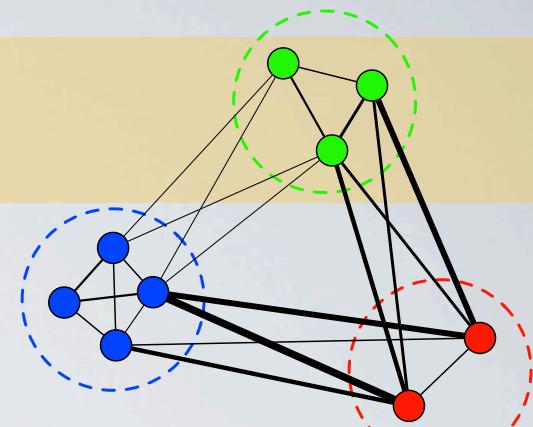
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► examples of weighted graphs:

frequency of social interactions (calls, txt, proximity, etc.)

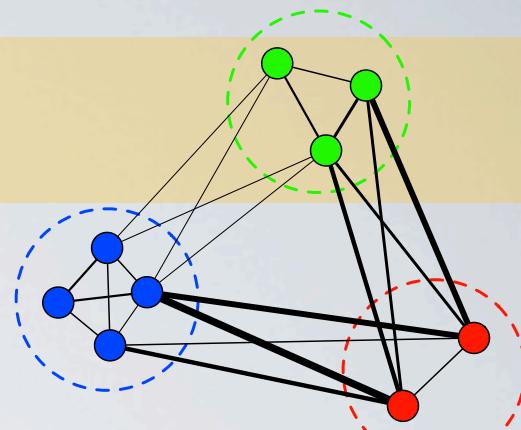
cell-tower traffic volume

other similarity measures

time-varying attributes

missing edges, active learning, etc.

weighted stochastic block model



block structure

$$\mathcal{R} : k \times k \rightarrow \{1, \dots, R\}$$

weight distribution

$$f$$

block assignment

$$z$$

weighted graph

$$G$$

likelihood function:

$$P(G | z, \theta, f) = \prod_{i < j} f(G_{i,j} | \theta_{\mathcal{R}(z_i, z_j)})$$

► given G and choice of f , learn z and θ

technical difficulties:

degeneracies in likelihood function

(variance can go to zero. oops)

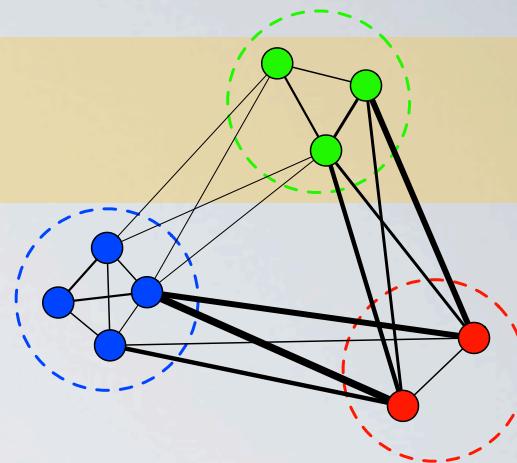
approximate learning

- edge generative model $P(G | z, \theta, f)$

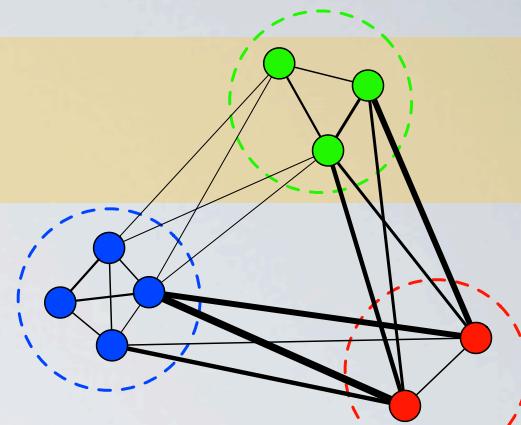
- estimate model via variational Bayes

conjugate priors solve degeneracy problem

algorithms for dense and sparse graphs



dense weighted SBM



approximate posterior distribution

$$\pi^*(z, \theta | G) \approx q(z, \theta) = \prod_i q_i(z_i) \prod_r q(\theta_r)$$

► estimate q by minimizing

$$D_{\text{KL}}(q || \pi^*) = \ln P(G | z, \theta, f) - \mathcal{G}(q)$$

where $\mathcal{G}(q) = \mathbb{E}_q(\mathcal{L}) + \mathbb{E}_q\left(\log \frac{\pi(z, \theta)}{q(z, \theta)}\right)$

for (conjugate) prior π for exponential family distribution f

► taking derivative yields update equations for z, θ

► iterating equations yields local optima

checking the model

synthetic network with known structure

- given synthetic graph with known structure
- run VB algorithm to convergence
- compare against choose threshold + SBM (and others)

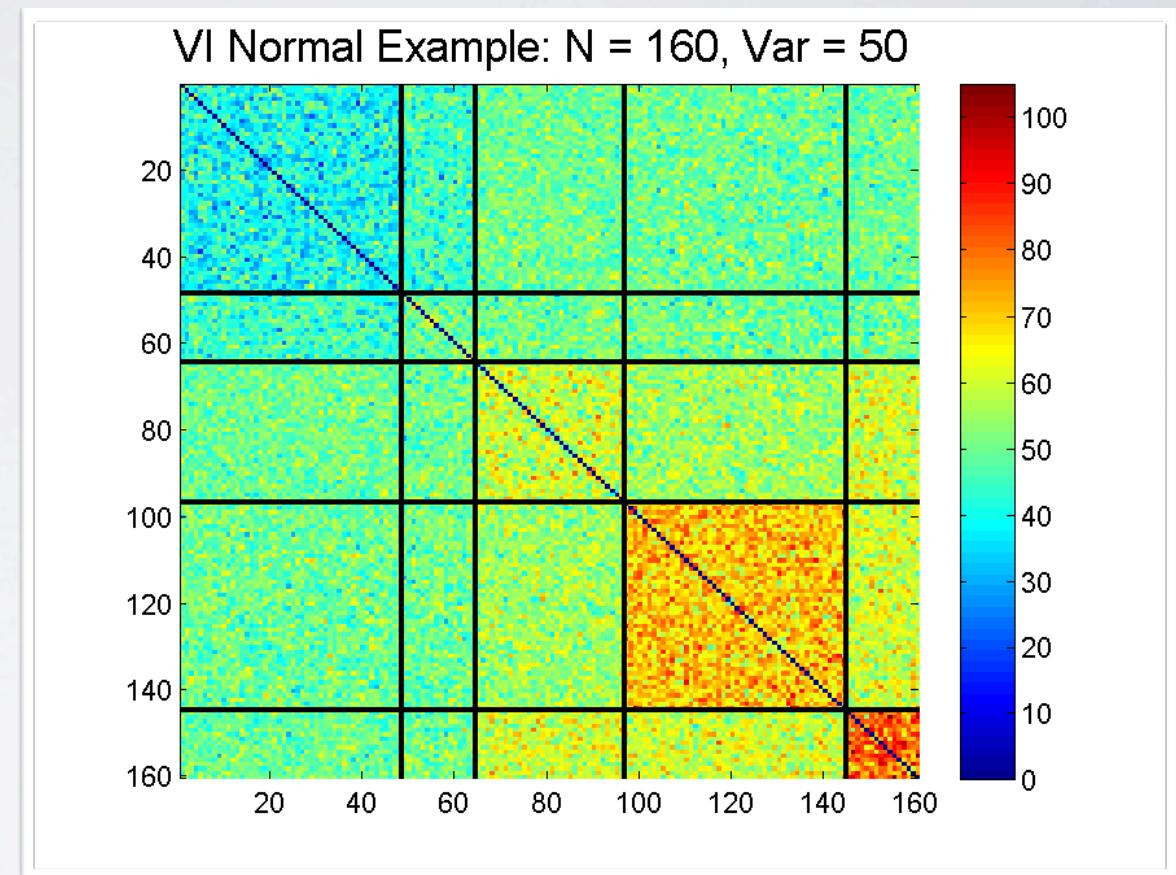
compute Variation of Information (partition distance)

$$\text{VI}(P_1, P_2) \in [0, \ln N]$$

checking the model

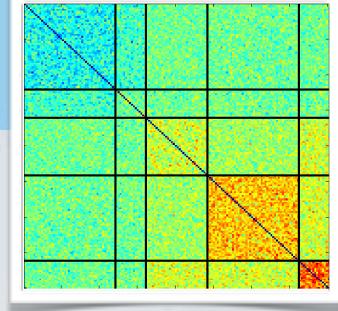
synthetic network with known structure

- variation of Newman's four-groups test
- $k^* = 5$ latent groups
 $n_r = [48, 16, 32, 48, 16]$
- Normal edge weights:
 $f = \mathcal{N}(\mu_r, \sigma_r^2)$



in this case $\text{VI}(P_1, P_2) \in [0, \ln k^* + 1.5] = [0, 3.1]$

learn better with more data

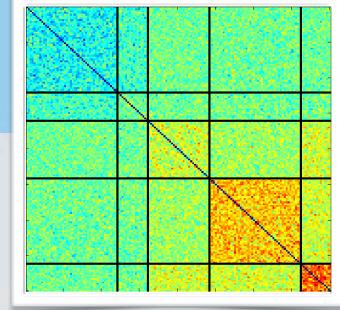


increase network size N

- fix $k = k^*$, $f = \mathcal{N}$
- bigger network, more data

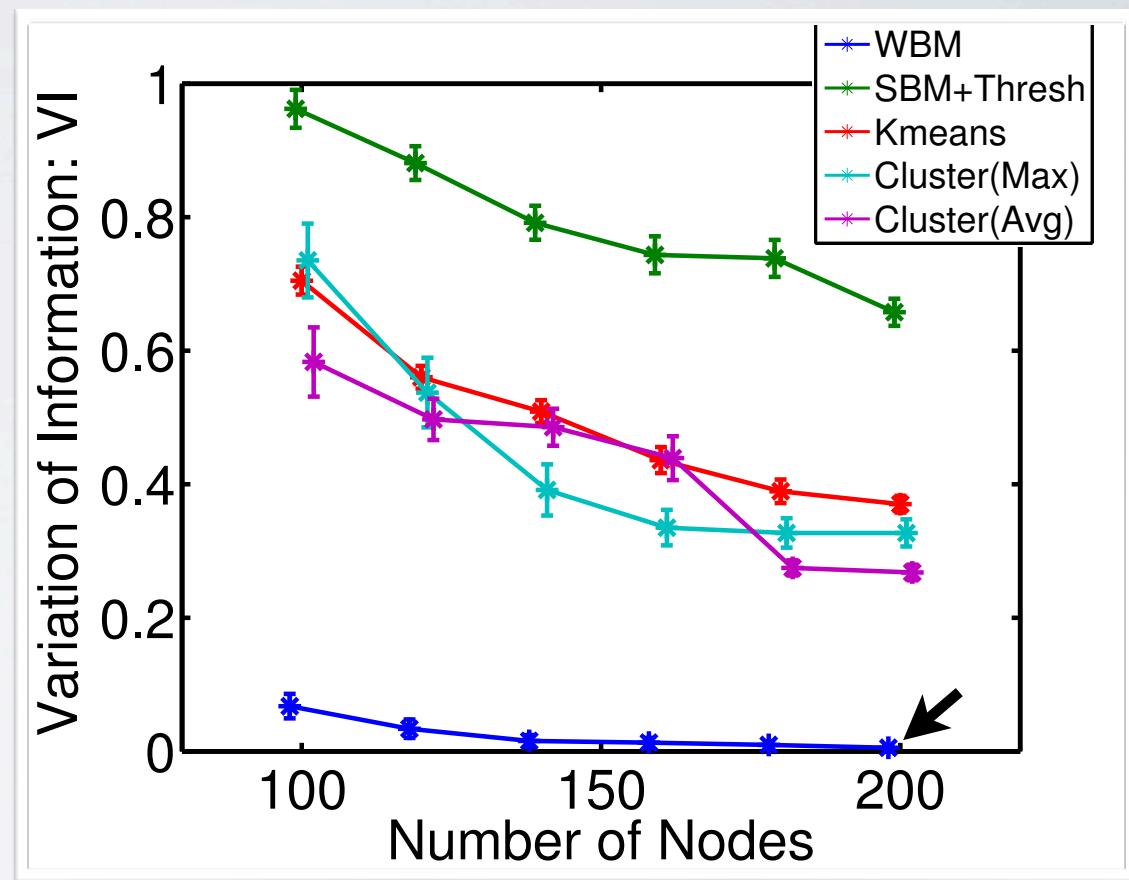
we keep the n_r/N constant

learn better with more data



increase network size N

- fix $k = k^*, f = \mathcal{N}$
- bigger network, more data
- WSBM converges on correct solution more quickly
- thresholding + SBM particularly bad

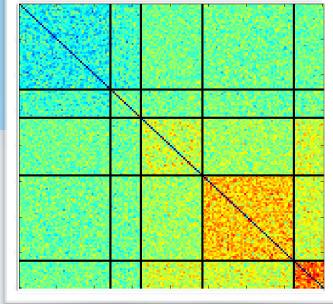


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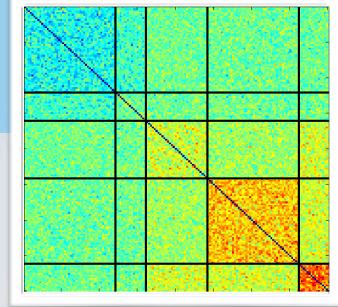
learning the number of groups

vary number of groups found k

- fix $f = \mathcal{N}$
- too few / many blocks?

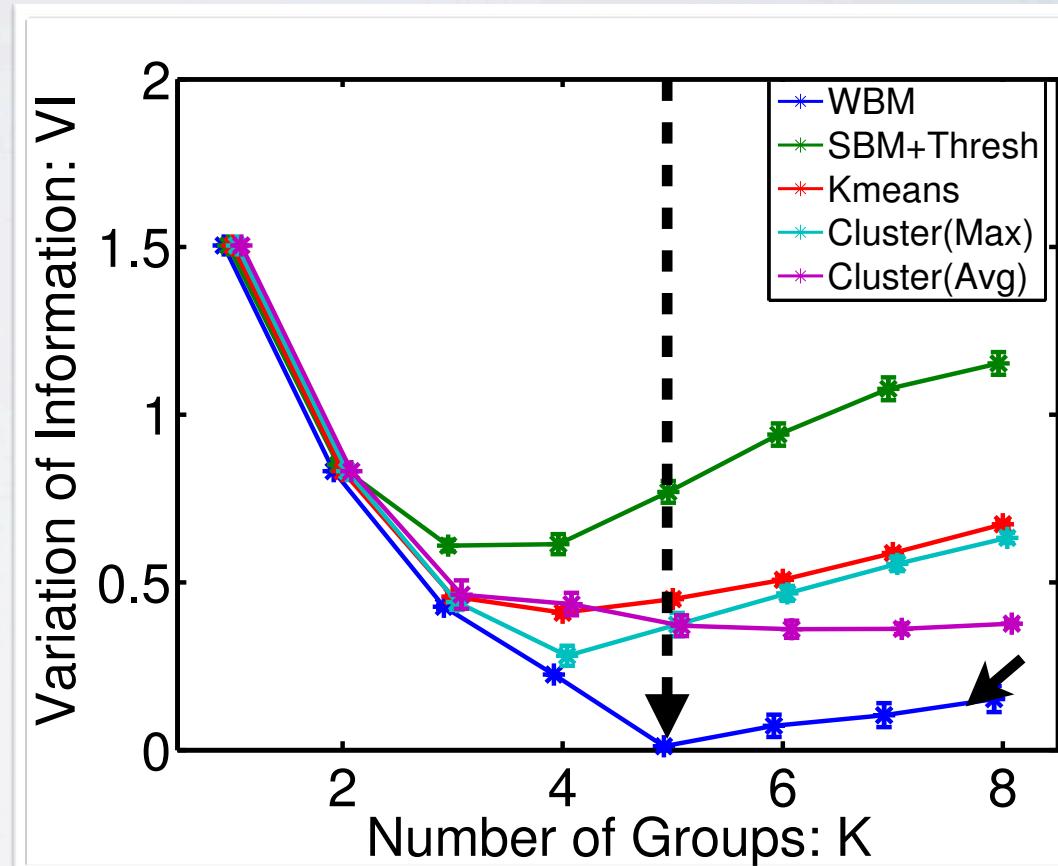


learning the number of groups



vary number of groups found k

- fix $f = \mathcal{N}$
- too few / many blocks?
- WSBM converges on correct solution
- WSBM fails gracefully when $k > k^*$
- others do poorly

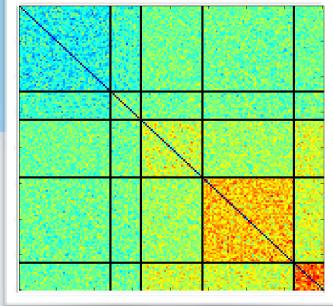


In fact, Bayesian marginalization will correctly choose $k=k^*$ in this case.

learning despite noise

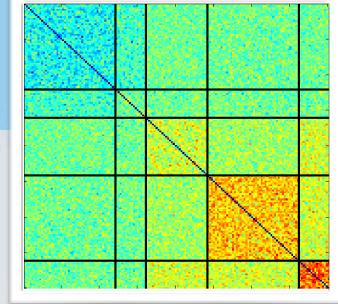
increase variance in edge weights σ_r^2

- fix $k = k^*, f = \mathcal{N}$
- bigger variance, less signal

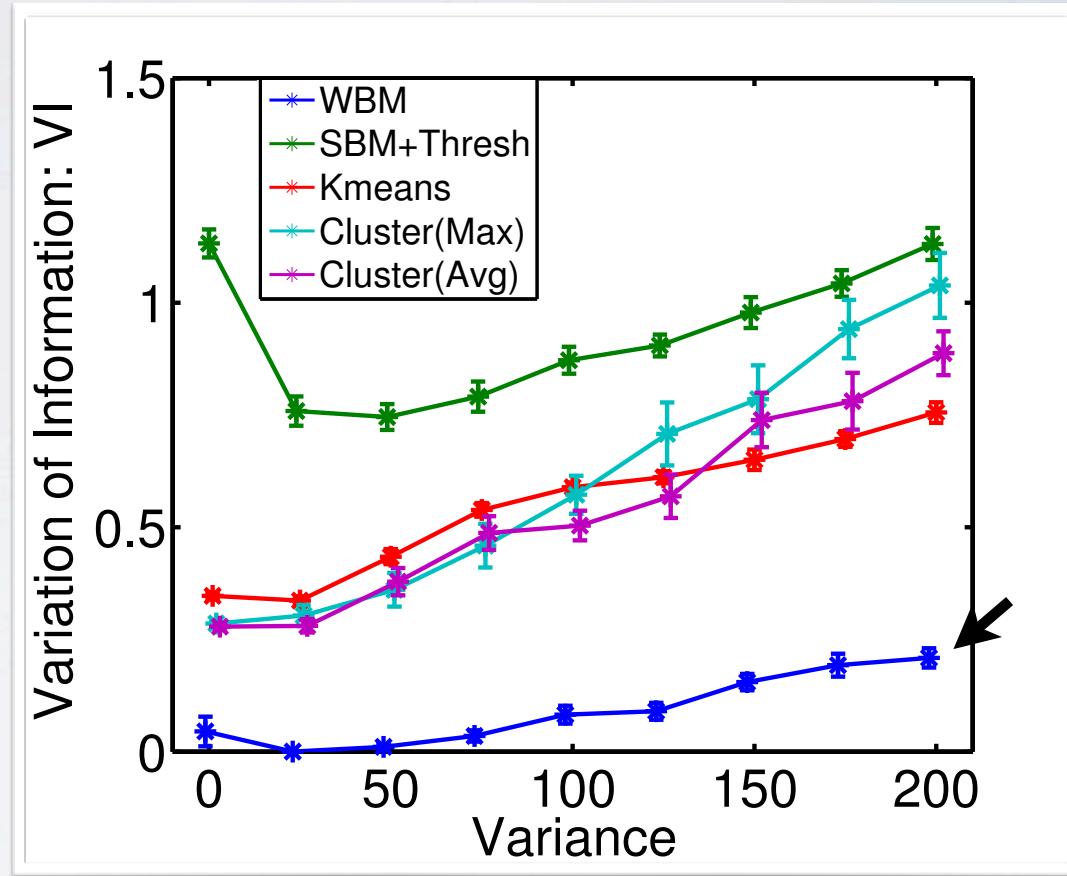


learning despite noise

increase variance in edge weights σ_r^2



- fix $k = k^*$, $f = \mathcal{N}$
- bigger variance, less signal
- WSBM fails more gracefully than alternatives, even for very high variance
- thresholding + SBM particularly bad



comments

- **single-scale structural inference**
mixtures of assortative, disassortative groups
- **inference is cheap (VB)**
approximate inference works well
- **thresholding edge weights is bad, bad, bad**
one threshold (SBM) vs. many (WSBM)
- **generalizations also for sparse graphs, degree-corrections, etc.**

generative models

- auxiliary information
 - node & edge attributes, temporal dynamics (beyond static binary graphs)
- scalability
 - fast algorithms for fitting models to big data (methods from physics, machine learning)
- model selection
 - which model is better? is this model bad? how many communities?
- model checking
 - have we learned correctly? check via generating synthetic networks
- partial or noisy data
 - extrapolation, interpolation, hidden data, missing data
- anomaly detection
 - low probability events under generative model

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acknowledgments



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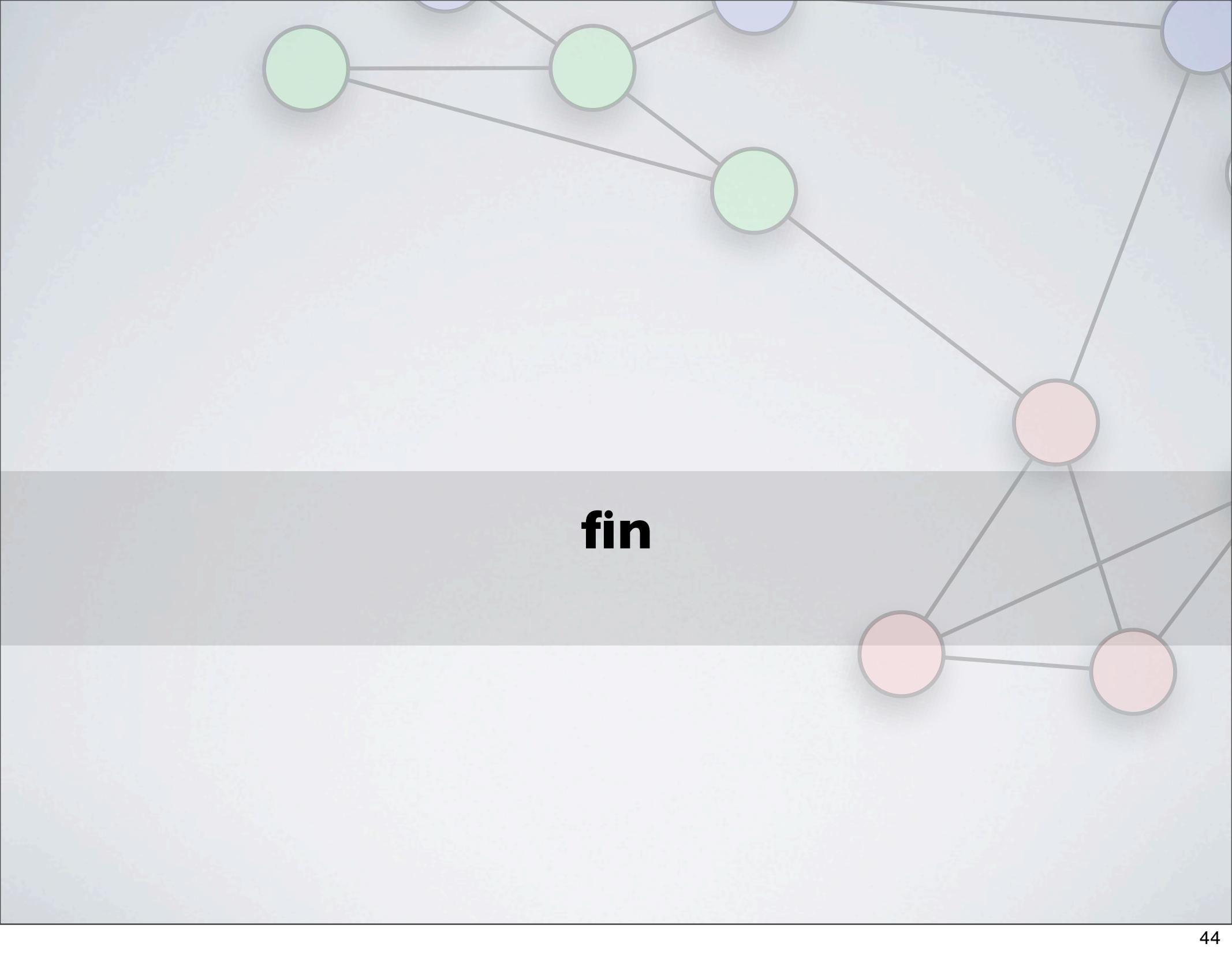
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some references

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- Park, Moore and Bader, "Dynamic Networks from Hierarchical Bayesian Graph Clustering." *PLoS ONE* 5(1): e8118 (2010).
- Clauset, Newman and Moore, "Hierarchical structure and the prediction of missing links in networks." *Nature* 453, 98-101 (2008)
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A network graph illustrating connections between nodes. The graph consists of several nodes represented by circles. Some nodes are filled with a light green color, while others are filled with a light pink color. All nodes have a thin gray outline. The connections are shown as thin gray lines. A cluster of three green nodes is located in the upper left. A single pink node is positioned centrally below them. To the right, there is a group of three pink nodes. A pink node is also located near the bottom center. Several lines connect the green nodes to each other and to the pink nodes. There are also internal connections within the pink node cluster.

fin