# Estimation methods in network models

#### Purnamrita Sarkar

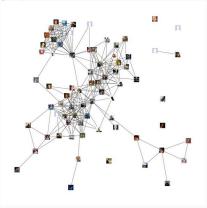
Department of Statistics and Data Sciences University of Texas, Austin

## **Clustering or Community detection**

- 1. A fundamental problem in exploratory analysis
- 2. Communities groups of nodes which behave similarly

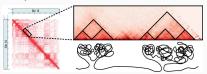
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  - 2.1 Networks: nodes are entities, links represent interactions between nodes. Communities could be
    - 2.1.1 groups of users in Facebook



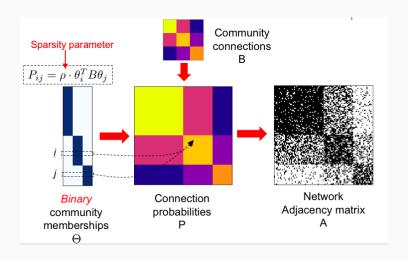
## **Clustering or Community detection**

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- 2. Communities groups of datapoints which behave similarly
  - 2.1 Networks: nodes are entities, links represent interactions between nodes. Communities could be
    - positions of a chromatin which are associated via 3D looping (Weinreb and Raphael 2005, Wang, S., Ursu, Kundaje and Bickel, 2018)



- Nodes denote a position on the chromatin
- edges measure how close they are in a 3D arrangement
- Goal is to find loops or hairballs, formally known as Topologically associated domains (TADs)
- These are preserved across cell types and also different species

# The stochastic block model (Holland, Laskey and Leinhardt 1983)



- 1. We will start by writing down the log likelihood for a fixed  $\Theta$ .
- 2. First note that the conditional expectation matrix  $E[A|\Theta] = \Theta B \Theta^T$  is blockwise constant.

$$\log P(A; \Theta, B)$$

$$= \sum_{i,j} A_{ij} \log P_{ij} + (1 - A_{ij}) \log(1 - P_{ij})$$

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$$\begin{split} &\log P(A;\Theta,B) \\ &= \sum_{i,j} A_{ij} \log P_{ij} + (1-A_{ij}) \log(1-P_{ij}) \\ &= \sum_{i,j} \sum_{k,\ell} \underbrace{\Theta_{ik}\Theta_{j\ell}}_{\mathbf{1} \text{ if } i \in C_k, j \in C_\ell} (A_{ij} \log B_{k\ell} + (1-A_{ij}) \log(1-B_{k\ell})) \end{split}$$

- 1. Let  $B_{ii} = p$  and  $B_{ij} = q$  for  $i \neq j$
- 2. Let cluster sizes be equal and n/k
- 3. Let  $X = \Theta \Theta^T$

$$\log P(A; \Theta, B) = \sum_{i,j} A_{ij} \log P_{ij} + (1 - A_{ij}) \log(1 - P_{ij})$$

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$$= \sum_{i,j} \left( X_{ij} A_{ij} \log \frac{p}{1 - p} + (1 - X_{ij}) A_{ij} \log \frac{q}{1 - q} \right) + \log(1 - p) n^2 / k + \log(1 - p) n^2$$

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# **Optimization perspective**

1. Optimization goal -

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- 2. Question is, what is  $\mathcal{F}$ ?
- 3. For k equal communities,  $\Theta^T \Theta = \frac{n}{k} I$ , where I is the identity matrix.

# **Spectral Clustering**

1. Consider

$$\arg\max_{\boldsymbol{\Theta}^T\boldsymbol{\Theta}=\frac{n}{k}}\langle\boldsymbol{\Theta}\boldsymbol{\Theta}^T,\boldsymbol{A}\rangle$$

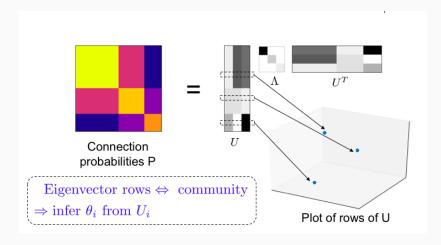
## **Spectral Clustering**

1. Consider

$$\arg\max_{\boldsymbol{\Theta}^T\boldsymbol{\Theta}=\frac{n}{k}}\langle\boldsymbol{\Theta}\boldsymbol{\Theta}^T,\boldsymbol{A}\rangle$$

- 2. As it turns out, this returns the top k eigenvectors of A (suitably scaled).
  - 2.1 Widely used in ML (Ng et al 2002, Shi and Malik 2001)
  - 2.2 Often you compute top k eigenvectors of the normalized adjacency matrix. (for consistency results, see Rohe et al 2010)

# **Spectral Clustering - why this works**



- 1. Lets generate a network from a blockmodel
- 2. First figure out parameters

```
K=2
m=50
n=K*m
Z=numpy.zeros([n,2])
Z[0:m,0]=1;
Z[m:n,1]=1;
B=np.array([[.3, .1], [.1, .3]])
```

1. Now build a symmetric random uniform matrix

P=Z.dot(B).dot(numpy.transpose(Z))

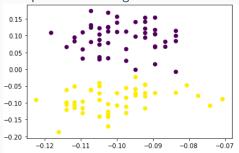
```
R=numpy.random.uniform(size=[n,n])
Rl=triu(R)+numpy.transpose(triu(R))
A=1*(R1<P)
A=A-np.diag(np.diag(A))
```

```
Dl=np.diag(sum(A,axis=0)**(-.5))
Kl=Dl.dot(A).dot(Dl)
u,s,vt=svd(Kl)
```

1. Do Spectral Clustering

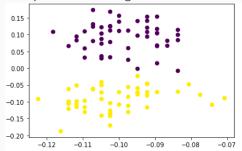
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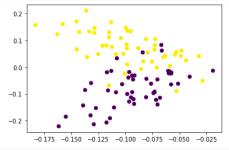
2. Accuracy of kmeans 90%

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1. Why would this work?

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- 5. Next Convex relaxations

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- 3. Natural feasible set is

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$$\sum_{a} \Theta_{ia} = 1, \forall i \in [n]\}$$

4. Instead of the above nonconvex objective, we will consider:

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4. Instead of the above nonconvex objective, we will consider:

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This is a convex objective function as long as we are careful about  $\mathcal{F}'$ 

1. We can think of the ideal X as a clustering matrix

$$X_{ij} = \begin{cases} 1 & \text{if } i, j \text{ belong to same class} \\ 0 & \text{otherwise} \end{cases}$$

2. We can use a slightly different feasible set, namely

$$\mathcal{F}' = \{X_{ij} \in [0, 1], \forall i, j \in [n] \$$
 $X_{ii} = 1$ 

$$\sum_{j} X_{ij} = 1, \forall i \in [n] \$$
 $X \succeq 0\}$ 

## **Semidefinite relaxations -pros**

- Variety of feasible sets for blockmodels and degree corrected blockmodels
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- 2. Robust to outliers (Cai et al 2014, Yan and S. 2016)
- 3. Has superior performance for sparse networks (Guedon et al 2014)

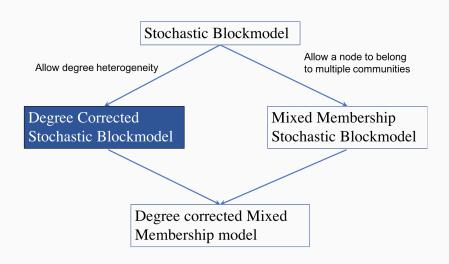
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#### Semidefinite relaxations

- 1. Very slow, scales to a few thousands of nodes
- 2. Requires to store  $n \times n$  clustering matrix—may become prohibitive for large networks
  - 2.1 Recently there have been a lot of effort on optimizing quantities like  $\langle A, YY^T \rangle$
  - 2.2 Known as Burer Monteiro methods, these have been shown to enjoy nice theoretical properties, e.g. the local optima in fact are the global optima, and saddle points can be escaped [Mei et al, 2017, Boumal et al 2018].

#### Generalizations of a blockmodel



## A real clustering dataset

- 1. Lets talk about the political blogs dataset.
- 2. Here, every node is a political blog, a link signifies which blog points to which other blog.
- 3. The labels (blue and red) signify political orientation of the blogs

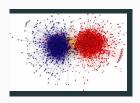


Figure 1: Lada Adamic and Natalie Glance." The political blogosphere and the 2004 US election: divided they blog." Proceedings of the 3rd international workshop on Link discovery. ACM, 2005.

# Political blogs degree distribution

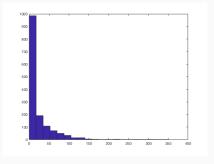


Figure 2: Histogram of degrees of nodes (after removing directions on edges)

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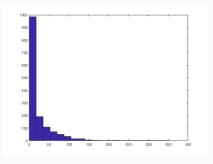


Figure 2: Histogram of degrees of nodes (after removing directions on edges)

1. Spectral Clustering using the top 2 eigenvectors of A fails here – clustering accuracy 60%

# Degree-corrected SBM (Karrer, Newman 2010)

#### Degree homogeneity in SBM models

- 1. Expected degrees are equal among different nodes
- 2. Real networks, there are usually "hub" nodes with very large degrees

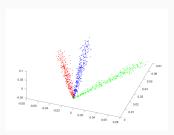
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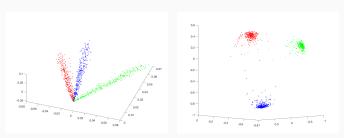
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#### Easy to fix-

- 1. Add degree parameter to each node that encodes the popularity
- 2.  $P(A_{ij} = 1) = \rho_n \gamma_i \gamma_j \theta_i^T B \theta_j$ , where  $\gamma_i$  is the degree parameter of node i.
- 3. Put constraints on the sum of them to make things identifiable.





Un-normalized and

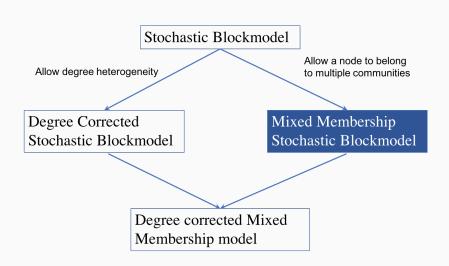
Row normalized top K eigenvectors

- 1. Normalize top K eigenvectors and do clustering (Chaudhuri et al 2012, Qin et al 2013)
- k-median based clustering algorithm on a low rank approximation of A followed by a refinement procedure. (Gao et al 2016).
- 3. SDP-based methods with regularization (Chen et al 2017)

## Political blogs again

- 1. If we take the top eigenvectors of the adjacency matrix and row normalize them prior to kmeans, the accuracy is around 80%
- If we just do normalized Spectral clustering on the largest connected component, then the error is nearly 50% (whether we row normalize or not)
- 3. If we do regularized spectral clustering without row normalization, error is about 30%.
- 4. If we do regularized spectral clustering with row normalization, error is about 5%.

#### Generalizations of SBM



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  - 4.1 Bayesian variant of NMF (Psorakis 2011)
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- For degree corrected mixed membership models, one needs to adapt the Spectral algorithms further (see Zhang and Levina 2014, Jin et al 2017, Mao et al 2019).

## The Mixed Membership Blockmodel (Airoldi et al, 2008)

- 1. Number of communities K
- 2.  $K \times K$  matrix of connection probabilities **B**
- 3.  $\theta_i \sim \text{Dirichlet}(\alpha_1, \dots, \alpha_K)$
- 4.  $A_{ij} \sim E[A_{ij}|\theta] = \rho_n \theta_i^T B \theta_j = \text{Call this } P$
- 5. Special case : Stochastic blockmodel when  $\alpha_a \to 0$  5.1 All  $\theta_i \in \{0,1\}^K$  have exactly one 1.
- 6. Large  $\alpha_a \equiv$  more overlap and small  $\alpha_a \equiv$  less overlap
- 7. Goal: Given **A**, infer  $\{\theta_i\}$  and **B**

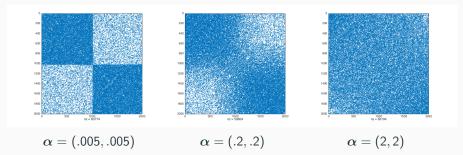
#### Dirichlet distribution

- 1. Parameters:  $\alpha_1, \ldots, \alpha_K > 0$
- 2. Density  $f(x_1, \ldots, x_K) = \frac{\prod_i x_i^{\alpha_i 1}}{B(\alpha)}$
- 3. Where  $x_1, \ldots x_k \ge 0$  belong to the K-1 simplex, i.e.

$$\sum_{i} x_i = 1, x_i \ge 0$$

https://upload.wikimedia.org/wikipedia/commons/thumb/ 5/54/LogDirichletDensity-alpha\_0.3\_to\_alpha\_2.0.gif/ 500px-LogDirichletDensity-alpha\_0.3\_to\_alpha\_2.0.gif

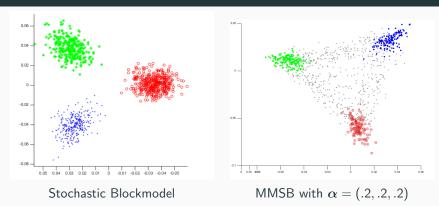
## The Mixed Membership Stochastic Blockmodel (MMSB)



lpha is the Dirichlet parameter for  $heta_i$ 

Given **A**, infer  $\{\theta_i\}$  and **B** 

## **Eigenvectors for 3 blocks**



- 1. Highlighted are  $S = \{i : \max_a \theta_{ia} > .9\}$ .
- 2. These are "pure" nodes

- Notable methods include Variational inference (Airoldi et al 2008, Gopalan et al 2013)
- 2. Tensor based methods (Anandkumar 2014, Hopkins et al 2018)
- 3. If B is positive semidefinite, then one can pose this as a symmetric non-negative matrix factorization problem (SNMF).
  - 3.1 Bayesian variant of NMF (Psorakis 2011)
  - $\it 3.2$  Use geometric intuition to solve the SNMF problem (Mao, S. and Chakrabarti 2017).

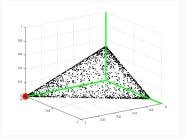
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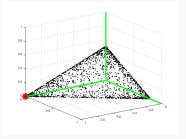
- 1. Eigenvectors fall on a simplex
  - We are essentially looking for a way to learn with K simplexes in K dimensional space
  - All points are convex combinations of the corners
  - Once you find the corners, all the parameters can be learned using a simple regression step
- 2. Let us try some simple ideas to find corners.
- 3. What if I find the node with maximum length?
  - 3.1 Indeed, it gives you "a nearly pure node" (with high probability).

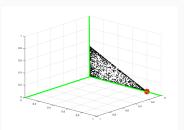
- 1. Scalable methods (Gillis et al 2014) in computational geometry to find corners of a noisy simplex with K corners in K dimensions.
  - 1.1 Find a node with largest  $\ell_2$  norm
  - 1.2 Remove its projection from the other rows.
  - 1.3 Repeat for K times.

# Building the geometric intuition with eigenvectors of P

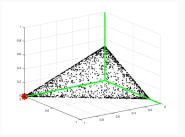


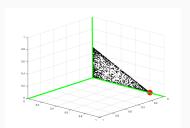
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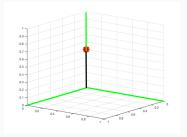




# Building the geometric intuition with eigenvectors of P







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- 3. As it turns out, in the mixed membership model, we have:

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1. Simple: use

$$\hat{V} = \hat{\Theta} \hat{V}_{\hat{S}} \Rightarrow \hat{\Theta} = \hat{V} \hat{V}_{\hat{S}}^{-1} \tag{1}$$

2. Recall

$$\Theta B \Theta^T = V E V^T \Rightarrow B = (\Theta^T \Theta)^{-1} \Theta^T V E V^T \Theta (\Theta^T \Theta)^{-1}$$

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$$\hat{B} = \hat{V}_{\hat{S}} \hat{E} \hat{V}_{\hat{S}}^T$$

# Estimation in Random Dot Product Graphs (RDPG models)

1. Edge probabilities [Young and Scheinerman 2007]

$$P(A_{ij} = 1 \mid Y) = \langle \underbrace{X_i}_{\text{Latent positions}}, X_j \rangle$$

- 2. The generalized RDPG model [Rubin-Delanchy et al, 2017] encompasses the Stochastic Blockmodel and its variants
- Popular methods include network embedding approaches using the adjacency matrix and its variants (Sussman et al 2012, Fishkind et al 2013, Tang et al 2013, Le et al 2017, Athreya et al 2016).

# Estimation in Random Dot Product Graphs (RDPG models)

- 1. If we can just use Spectral Clustering, why go into all the trouble to do all we did for MMSB?
- 2. We could have just used  $\hat{X} = \hat{V}\hat{E}^{1/2}$
- 3. But note that,  $\hat{X}$  by itself does not mean anything. It can be used to cluster nodes for clustering models like blockmodels.
- 4. But when there is mixed memberships,  $\hat{X}$  is essentially a transformed version of  $\Theta$ .
- 5. But in order to get to  $\Theta$ , we need to do more work.

## Triangle formation and block models

- 1. Real social networks have many triangles, even if they are sparse.
- To be particular, the global clustering coefficient, defined as number of triangles divided by number of closed or open triplets is often used to measure "clustered" networks are.
- 3. But for blockmodels or its variants, as the network gets sparser, the network becomes more treelike.

## Triangle formation and block models

- 1. Real social networks have many triangles, even if they are sparse.
- To be particular, the global clustering coefficient, defined as number of triangles divided by number of closed or open triplets is often used to measure "clustered" networks are.

#### Latent distance models

- 1. Latent distance models model homophily or transitivity by introducing a latent space where nodes lie.
- 2. Two nodes close in the latent space are more likely to be connected.
- 3. Why do you think this leads to transitivity and reciprocity (what is that)?

#### Latent distance models

- 1. Reciprocity:
  - 1.1 If  $i \rightarrow j$ , then the event  $j \rightarrow i$  is more likely.
  - 1.2 Why?

#### Latent distance models

- 1. Reciprocity:
  - 1.1 If  $i \rightarrow j$ , then the event  $j \rightarrow i$  is more likely.
  - 1.2 Why?
- 2. Transitivity:
  - 2.1 If  $i \to j$  and  $j \to k$ , then  $i \to k$  is more likely.
  - 2.2 Why?

## Latent Distance Models [Hoff et al 2002]

1. Log likelihood:

$$\begin{split} \log P(A \mid \eta) &= \sum_{i \neq j} \left\{ \eta_{ij} \cdot A_{ij} - \log \left( 1 + e^{\eta_{ij}} \right) \right\}, \\ \text{where } \eta_{ij} &= \log \ \textit{odds} (A_{ij}, z_i, z_j) = \alpha - \|z_i - z_j\|_2 \end{split}$$

- Two stage approaches, which initialize with Spectral methods [S. and Moore, 2005] no guarantee for global optima
- Recently there has been some work on consistency of convex relaxation based inference, and non-convex inference methods [Ma et al 2020].

## Estimation for Latent Space Models: Bayesian Approach

1. Log likelihood:

$$\begin{split} \log P(A \mid \underbrace{z_i, i \in [n]}_{\text{Latent positions}}) &= \sum_{i \neq j} \left\{ \eta_{ij} \cdot A_{ij} - \log \left( 1 + e^{\eta_{ij}} \right) \right\}, \\ \text{where } \eta_{ij} &= \log \ \textit{odds}(A_{ij}, z_i, z_j) = \alpha - \|z_i - z_j\|_2 \end{split}$$

- 2. Alternatively, place priors on  $\alpha$ , Z.
- 3. Use Metropolis-Hastings to update Z and  $\alpha$  serially.
- The Bayesian approach can be computationally prohibitive for larger networks.