

Graphs and Community Detection



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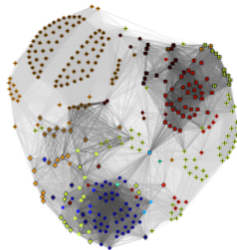
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MATH 373



- Networks
- Exploratory analysis of networks via community detection
 - Spectral clustering
 - Modularity optimization methods
 - Stochastic block modeling methods
 - Extraction-based methods
- Inference on networks via random graph models
- Software and useful resources

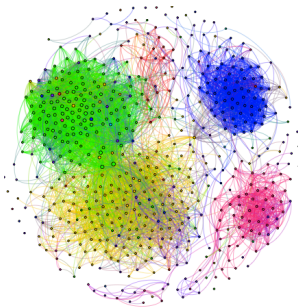


- Means to visualize and model interactions of a complex system
- Treat an “actor” of the system as a **vertex** and place **edges** between actors that interact



- Interactions of virulence factor genes of *E. coli* from UTI patients

Reference: Parker et al. "Network analysis reveals sex- and antibiotic resistance-associated antivirulence targets in clinical uropathogens"
American Chemical Society: Infectious Diseases (2015)



- Friendships among my friends on Facebook

Reference: Wilson et al. "A testing based extraction algorithm for identifying significant communities in networks" *Annals of Applied Statistics* (2014)



Primary goal: Analyze and interpret *relational* data

- Exploratory analysis and visualization
- Simulation and inference of graphical models
- Development of scalable algorithms
- Comparison of statistical algorithms and methodology



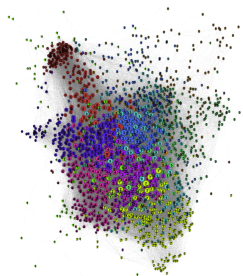
Graph $G = ([n], E)$:

- **Vertices** ($[n] = \{1, \dots, n\}$): actors of the system
- **Edges** ($E \subseteq [n] \times [n]$): placed between actors w/ relationship
- **Adjacency Matrix** $A \in \mathcal{R}^{n \times n}$:

$A(u, v)$ = edge weight between nodes $u, v \in [n]$

- **Degree sequence**: $\mathbf{d} = (d(1), \dots, d(n))$ where

$$\begin{aligned} d(i) &= \# \text{ of edges incident on node } i \\ &= \sum_{v \in [n]} A(u, v) \end{aligned}$$



Informally: Identify **communities** $C_1, \dots, C_k \subseteq [n]$ such that

- Edge density within sets C_i is large
- Edge density between sets C_i is small

Aim: Capture relevant structure of a complex system



Min-cut

- Identify cut of vertices that "cuts" the fewest edges

Modularity

- Partition that deviates most from organization in random graph

Spectral

- Focus on spectral properties of graph Laplacian

Stochastic Block Model

- Approximate Maximum Likelihood Estimation

Extraction

- Identify dense communities one at a time



$G = (V, E)$ undirected graph on $V = \{1, \dots, n\}$ and adjacency matrix S

- Define $D = \text{diag}(d(1), \dots, d(n)) \in \mathcal{R}^{n \times n}$ where
- Graph Laplacian L :

$$L = D - S$$

- Normalized graph laplacian L_{norm} :

$$L_{norm} = D^{-1}L = I - D^{-1}S$$



- λ is an eigenvalue of L_{norm} with eigenvector v iff λ and v solve the eigenproblem $Lv = \lambda Dv$
- 0 is an eigenvalue of L and L_{norm} with eigenvector $\mathbf{1}$
- L and L_{norm} are nonnegative definite and have n real-valued eigenvalues $0 = \lambda_1 \leq \dots \leq \lambda_n$
- L is symmetric



Theorem 1.

Let G be an undirected graph with non-negative weights and let L_{norm} be its normalized graph laplacian.

Let k = the multiplicity of the eigenvalue 0 of L_{norm} . Then,

- (1) k is the number of connected components C_1, \dots, C_k in G*
- (2) The eigenspace of 0 is spanned by the indicator vectors $\mathbf{1}_{C_i}$*

Remark:

- If G clustered into k disjoint connected components, then we can perfectly identify the k clusters using the k smallest eigenvectors



Algorithm

Input: Adjacency matrix $A \in \mathcal{R}_+^{n \times n}$, number of communities k

1 Calculate normalized graph laplacian L_{norm}

2 Compute

X = the $n \times k$ matrix of the k smallest eigenvectors of L_{norm}

3 Cluster the rows of X using k-means

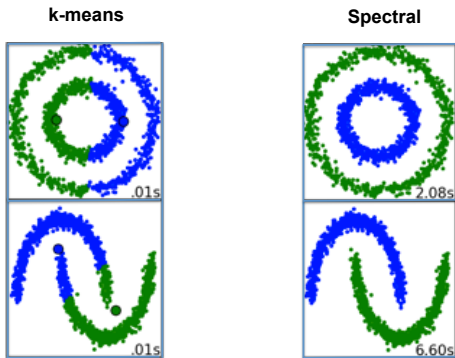
Output: Clusters C_1, \dots, C_k



- Requires a prespecified number of clusters k
- Works perfectly in an ideal scenario
- Requires the use of another clustering method (k-means)
- The solution to a relaxed version of the [normalized-cut problem](#)

Reference: Ulrike Von Luxburg "A tutorial on spectral clustering"
(2006)

A Quick Comparison



- Spectral can find special structure due to connectivity/similarity
- k-means is faster and better suited for compact clusters



- **Model-based** approach to community detection
- $G = (V = [n], E)$ with binary adjacency matrix A
- Assumes that G has k blocks generated as follows:

- 1 Community labels $\mathbf{c} = (c_1, \dots, c_n)$ generated at random:

$$c_1, \dots, c_n \stackrel{iid}{\sim} \text{multinomial}(1, \pi = \{\pi_1, \dots, \pi_k\})$$

- 2 Conditional on \mathbf{c} , $A(u, v)$ are independent Bernoulli rvs with

$$\mathbb{E}[A(u, v) | \mathbf{c}] = P_{c_u, c_v}$$

Reference: Holland, et al. "Stochastic block models: first steps" (1983)



- Observe $G = G_o$, calculate likelihood $\mathcal{L}(\Theta|G_o, k)$ with $\Theta = \{\mathbf{P}, c\}$
- Finding c becomes an **estimation** problem:

$$\hat{\Theta} = \arg \max_{\Theta} \mathcal{L}(\Theta|G_o, k)$$

- Requires approximate algorithms like MCMC or variational EM
- **Issue:** Approximate algorithms can be slow!



- Requires pre-specified k
- “Best” performance on identifying disjoint communities
- Block labels are consistent (as $n \rightarrow \infty$) if for all $i \neq \ell$:

$$nP_{i,i} - nP_{i,\ell} \geq \sqrt{k(nP_{i,i} + (k-1)nP_{i,\ell})}$$

- Algorithms like MCMC and variational EM can be slow!



- **Aim:** find the partition of G whose communities contain the highest density of edges relative to the expected density of edges

Remarks:

- Requires a notion of what a *random* network looks like
- The choice of a null network model affects resulting communities

Reference: Mark E Newman "Modularity and community structure in networks" (2004)



- Graph $G = (V = [n], E)$, adjacency matrix $A = [A_{u,v}]$
- **Modularity (Q)**: Measures the “significance” of partition \mathbf{c} :

$$Q(\mathbf{c}) = \frac{1}{2|E|} \sum_{u,v} \left[\left(A(u,v) - \frac{d(u)d(v)}{2|E|} \right) \mathbb{I}\{c_u = c_v\} \right]$$

- Measures the average departure of **observed** edge density from **expected** edge density



Aim: Find the labels $c^* \in \{1, \dots, k\}^n$ that maximizes modularity:

$$c^* = \arg \max_c \{Q\}$$

- NP hard optimization problem
- *Many* approximate algorithms developed

Reference: Santo Fortunato, "Community detection in graphs" (2009).
[100+ page review paper]



Basic Idea:

- Identify communities $C_i \subseteq V$ one at a time via iterative search
- Remove/avoid C_1, \dots, C_i when searching for C_{i+1}

Virtues:

- Possible to accommodate overlap
- Automatic selection of number of communities
- Parallelizable! Can easily scale to large networks.



Methods:

- **OSLOM**: Lancichinetti, et al. "Finding statistically significant communities in networks" (2011) – resampling based method
- **Extraction**: Zhao, et al. "Community extraction for social networks" (2011) – score-based residualizing
- **ESSC**: Wilson, et al. "A testing based extraction algorithm for identifying significant communities in networks" (2014) – hypothesis testing based extraction



Aim: Model the occurrence of an observed graph $G = ([n], E)$ from a family of graphs \mathcal{G} .

Use: the distribution on \mathcal{G} gives a means to make inference on complicated network systems.

Reference: Anna Goldenberg et al. "A Survey of Statistical Network Models" (2009)



- "Small-world brain networks" (2006)
- "Exponential random graphs for social networks" (2012)
- "The structure of adolescent romantic and sexual networks" (2004)
- "Childhood peer network characteristics: genetic influences and links with early mental health trajectories" (2015)
- "Network biology: understanding the cell's functional organization" (2004)



- **Erdős-Rényi model:** Independent edges with probability

$$\mathbb{P}(\{u, v\} \in E) = p$$

- **Configuration model:** Independent edges with probability

$$\mathbb{P}(\{u, v\} \in E) = \frac{d(u)d(v)}{\sum_w d(w)}$$

- **Beta model:** Let $\beta \in (0, \infty)^n$. Independent edges with probability

$$\text{logit}(\mathbb{P}(\{u, v\} \in E)) = \beta_u + \beta_v$$



- **Latent space model**: Independent edges with

$$\text{logit}(\mathbb{P}(\{u, v\} \in E)) = X\beta - \delta_{u,v}$$

- X = design matrix of covariates
- $\delta_{u,v}$ = latent distance between nodes $u, v \in [n]$
- **Stochastic block model**: Conditional on community labels \mathbf{c} , edges are independent with probability

$$\mathbb{P}(\{u, v\} \in E) = P_{c_u, c_v}$$



- **Exponential Random Graph model (ERGM)**: let A denote the observed binary adjacency matrix. Then,

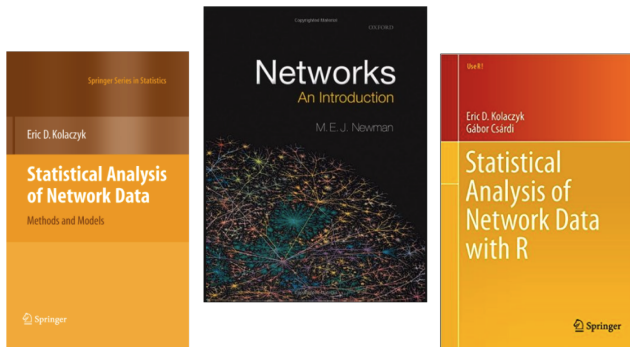
$$P(\mathcal{A} = A) \propto \exp\{\theta^T h(A)\}$$

- $h(A) : \{0, 1\}^{\binom{n}{2}} \rightarrow \mathcal{R}^p$ = network statistics (both endogenous and exogenous).
- **Generalized Exponential Random Graph model (GERGM)**: extension of ERGM to general weighted networks.

Reference: Wilson et al. "Stochastic weighted graphs: flexible model specification and simulation" (2015)



- Appropriately incorporating edge dependencies
- Models that do incorporate dependencies are intractable and rely on approximation / MCMC methods (e.g, ERGM and GERGM)
- Scalability
- Dynamic random graph models



Useful R packages: *igraph* (also available in python); *statnet*; *gergm*