

# Chapter 11. Network Community Detection

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PubH 7475/8475

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# Outline

- ▶ Introduction
- ▶ Spectral clustering
- ▶ Hierarchical clustering
- ▶ Modularity-based methods
- ▶ Model-based methods
- ▶ Key refs:
  1. Newman MEJ
  2. Zhao Y, Levina E, Zhu J (2012, Ann Statist 40:2266-2292).
  3. Fortunato S (2010, Physics Reports 486:75-174).
- ▶ R package `igraph`: drawing networks, calculating some network statistics, some community detection algorithms, ...

# Introduction

- ▶ Given a binary (undirected) network/graph:  $G = (V, E)$ ,  
 $V = \{1, 2, \dots, n\}$ , set of nodes;  $E$ , set of edges.  
Adjacency matrix  $A = (A_{ij})$ :  $A_{ij} = 1$  if there is an edge/link  
b/w nodes  $i$  and  $j$ ;  $A_{ij} = 0$  o/w. ( $A_{ii} = 0$ )
- ▶ Goal: assign the nodes into  $K$  “homogeneous” groups.  
often means dense connections within groups, but sparse b/w  
groups.
- ▶ Why? Figs 1-4 in Fortunato (2010).

# Spectral clustering

- ▶ Laplacian  $L = D - A$ , or ...

$$D = \text{Diag}(D_{11}, \dots, D_{nn}), D_{ii} = \sum_j A_{ij}.$$

- ▶ Intuition:

If a network separates perfectly into  $K$  communities, then  $L$  (or  $A$ ) is block diagonal (after some re-ordering of the rows/columns).

If not perfectly but nearly, then the eigenvectors of  $L$  are (nearly) linear combinations of the indicator vectors.

- ▶ Apply K-means (or ..) to a few ( $K$ ) eigenvectors corresponding to the smallest eigenvalues of  $L$ .

Note: the smallest eigen value is 0, corresponding to eigenvector 1.

- ▶ Two clusters  $\implies$  spectral bisection: use the eigenvector of the second smallest eigen value; partition by its positive/negative elements.

Generally, repeatedly apply the above to each cluster... vs apply SC once?

- ▶ Widely used; some theory (e.g consistency):

## Modified spectral clustering

- ▶ SC may not work well for sparse networks.
- ▶ Regularized SC (Qin and Rohe): replace  $D$  with  $D_\tau = D + \tau I$  for a small  $\tau > 0$ .
- ▶ SC with perturbations (Amini, Chen, Bickel, Levina, 2013, Ann Statist 41: 2097-2122):  
regularize  $A$  by adding a small positive number on a random subset of off-diagonals of  $A$ .

# Hierarchical clustering

- ▶ Need to define some similarity or distance b/w nodes.
- ▶ Euclidean distance:  $A_i. = (A_{i1}, A_{i2}, \dots, A_{in})'$ ,

$$x_{ij} = \|A_i. - A_j.\|_2$$

- ▶ Or, Pearson's corr,

$$x_{ij} = \text{corr}(A_i., A_j.)$$

- ▶ Then apply a hierarchical clustering.  
can be used to re-arrange the rows/columns of  $A$  to get a nearly block-diagonal  $A$ .
- ▶ Fig 3 in Neuman.
- ▶ Fig 2 in Meunier et al (2010).

## Algorithms based on edge removal

- ▶ Divisive: edges are progressively removed.
- ▶ Which edges? "bottleneck" ones.
- ▶ *edge betweenness* is defined to be the number of shortest paths between all pairs of all nodes that run through the two nodes.
- ▶ Algorithm (Girvan and Neuman 2002, PNAS):
  - 1) calculate *edge betweenness* for each remaining edge in a network;
  - 2) remove the edge with the highest *edge betweenness*;
  - 3) repeat the above until ...
- ▶ A possible stopping criterion: *modularity*, to be discussed.
- ▶ Fig 4 in Neuman.
- ▶ Remarks: slow; some modifications, e.g. a Monte Carlo version in calculating *edge betweenness* using only a random subset of all pairs; or use a different criterion.
- ▶ R package igraph: `cluster_edge_betweenness()`

# Modularity-based methods

- ▶ Notation:

degree of node  $i$ :  $d_i = D_{ii} = \sum_{j=1}^n A_{ij}$ ,

(twice) total number of edges:  $m = \sum_{i=1}^n d_i$ ,

Community assignment:  $C = (C_1, C_2, \dots, C_n)$ ; **unknown**,

$C_i \in \{1, 2, \dots, K\}$ : community containing node  $i$ .

- ▶ Modularity: given  $C$ ,

$$Q = Q(C) = \frac{1}{2m} \sum_{i,j} \left( A_{ij} - \frac{d_i d_j}{m} \right) I(C_i = C_j).$$

- ▶ Intuition: obs'ed - exp'ed

- ▶ Goal:  $\hat{C} = \arg \max_C Q(C)$

Assumption: good to maximize  $Q$ , reasonable but ...



- ▶ Key: a **combinatorial** optimization problem!  
seeking exact solution will be too slow  $\implies$  many *approximate* algorithms, such as greedy searches (e.g. genetic algorithms, simulated annealing), relaxed algorithms, ...  
Newman (2003): repeat: combining two nodes  $i$  and  $j$  with  $A_{ij} = 1$  and the largest increase (or smallest decrease) in  $Q$ ;  
until all nodes in one community.  
 $\implies$  hierarchical; choose one with the largest  $Q$ .
- ▶ Very nonparametric?!
- ▶ Problems: resolution limit; too many local solutions.  
cannot detect relatively small communities; why? an implicit null model for the *whole network* (Fortunato 2010, p.40).
- ▶ R package igraph:  
greedy search, approx./fast: `cluster_fast()`;  
combinatorial search, exact/slow: `cluster_optimal()`;  
heuristic, hierarchical communities for large networks (e.g. millions of nodes); see Blondel et al (2008) in the manual: `cluster_louvain()`.

## Model-based methods

- ▶ Stochastic block model SBM (Holland et al 1983):
  - 1) a  $K \times K$  probability matrix  $P$ ;
  - 2)  $A_{ij} \sim \text{Bin}(1, P_{C_i, C_j})$  independently.
- ▶ Simple; can model dense/weak within-/between-community edges.

But, treat all nodes/edges in a community equally; cannot model *hub* nodes!

Scale-free network: node degree distribution  $Pr(k)$  is heavy-tailed; a power law.
- ▶ SBM with  $K = 1$ : Erdos-Renyi Random Graph.
- ▶ Degree-corrected SBM (DCSBM) (Karrer and Newman 2011):
  - 1)  $P$ ; each node  $i$  has a degree parameter  $\theta_i$  (with some constraints for identifiability);
  - 2)  $A_{ij} \sim \text{Bin}(1, \theta_i \theta_j P_{C_i, C_j})$  independently

- ▶ More notations:

$n_k(C) = \sum_{i=1}^n I(C_i = k)$ , number of nodes in community  $k$ ;

$O_{kl} = \sum_{i,j=1}^n A_{ij} I(C_i = k, C_j = l)$ , number of edges b/w communities  $k \neq l$ ;

$O_{kk} = \sum_{i,j=1}^n A_{ij} I(C_i = k, C_j = k)$ , (twice) number of edges within community  $k$ ;

$O_k = \sum_{l=1}^K O_{kl}$ , sum of node degrees in community  $k$ ;

$m = \sum_{i=1}^n d_i$ , (twice) the number of edges in the network.

- ▶ Objective function: A profile likelihood (profiling out nuisance parameters  $P$  and  $\theta$ 's based on a Poisson approximation to a binomial).

Given a likelihood  $L(C, P)$ ,

a profile likelihood  $L^*(C) = \max_P L(C, P) = L(C, \hat{P}(C))$ .

- ▶ SBM:

$$Q_{SB}(C) = \sum_{k,l=1}^K (O_{kl} \log \frac{O_{kl}}{n_k n_l}).$$

- ▶ DCSBM:

$$Q_{DC}(C) = \sum_{k,l=1}^K (O_{kl} \log \frac{O_{kl}}{O_k O_l}).$$

- ▶ Neuman-Girvan modularity:

$$Q_{NG}(C) = \frac{1}{2m} \sum_k (O_{kk} - \frac{O_k^2}{m}).$$

- ▶ Remarks: Still a combinatorial optimization problem; better theoretical properties.
- ▶ Numerical examples in Zhao et al (2012).

## Other topics

- ▶ Summary statistics for networks; e.g. clustering coefficient,...
- ▶ Weighted networks; with or without negative weights (e.g. Pearson's correlations).
- ▶ Overlapping communities.
- ▶ Time-varying (dynamic) networks.
- ▶ With covariates. How to model covariates?
- ▶ Fast (approximate) algorithms; theory.