

Introduction to statistical network analysis

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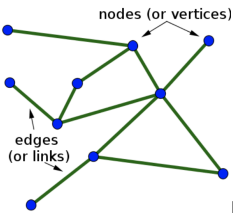
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Basic concepts

Basic concepts

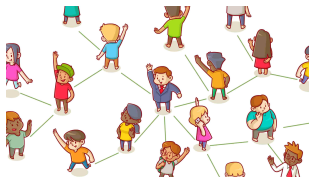


- **Nodes (Vertices):** participants in the network
Network size: number of nodes
- **Edges (Links):** connections or relationships between nodes
- **Examples [nodes (edges)]:**
 - people (friendship)
 - webpages (hyperlinks)
 - paper (citation)
 - genes (regulatory actions)
 - brain regions (oxygen level correlation)
- **Adjacency matrix:** $A_{i,j}$ is the edge between nodes (i, j)

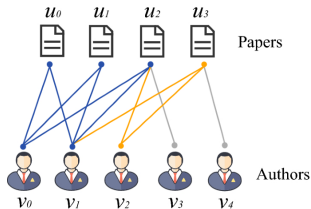
Basic concepts

Classification of network data: (1) partite

- **Unipartite network:** any node may connect to anyone else



- **Bipartite network:** nodes partitioned into two groups, only between-group edges are possible



Basic concepts

Classification of network data: (2) edge direction

- **Undirected (Symmetric) network:** $A_{i,j} = A_{j,i}$
- **Directed (Asymmetric) network:** $A_{i,j}$ and $A_{j,i}$ might not equal

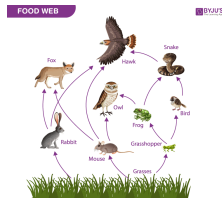
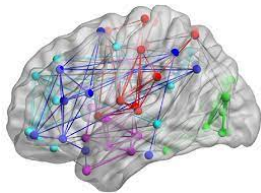


Figure: Left: undirected network; right: directed network

Classification of network data: (2) edge weight

- **Binary:** $A_{i,j} \in \{0, 1\}$
Example: friend (1) or non-friend (0)
- **Signed:** $A_{i,j} \in \{0, 1, -1\}$
Example: friend (1) or foe (-1) or no interaction (0)
- **Weighted:**
Example: trade surplus/deficit between countries

Statistical research aims:

- modeling network formation
- finding roles of individual nodes
Examples: community detection, node embedding
- stochastic behavior of network features
Examples: network moments
- link prediction

Network data vs conventional data:

- Networks: **no individual observation**, only relational data
- Deriving “**network analogues** of classical techniques”

Examples:

- one-/two-sample test
- cross-validation
- method-of-moments
- goodness-of-fit test
- re/subsampling

Popular network models

Now we survey Network models: Erdos-Renyi model

- Simple \rightarrow complex
- For simplicity, focus on **undirected**, **binary** networks
- Model **probability matrix**:

$$P_{i,j} := \mathbb{P}(A_{i,j} = 1)$$

Network models: Erdos-Renyi

Erdos-Renyi (ER) model (Newman 2018)

- Model formula:

$$P_{i,j} := \mathbb{P}(A_{i,j} = 1) \equiv p$$

- Definition:** network density is the (order of) average edge probability:

$$\rho_n \asymp \bar{P} := \binom{n}{2}^{-1} \sum_{1 \leq i < j \leq n} P_{i,j}$$

In ER model, $\rho_n \asymp p$

- Density is usually a crucial measure for assessing the **problem difficulty** of estimation/testing

Erdos-Renyi model

$$P_{i,j} := \mathbb{P}(A_{i,j} = 1) \equiv p \asymp \rho_n$$

Model estimation:

- **Moment estimator:**

$$\hat{p} := \binom{n}{2}^{-1} \sum_{1 \leq i < j \leq n} A_{i,j}$$

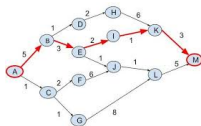
- **Consistency** (Chen et al. 2021): When $\binom{n}{2}p \rightarrow \infty$, $\hat{p}/p \xrightarrow{P} 1$
Roughly speaking

$$\frac{|\hat{p} - p|}{p} \underset{\text{approx.}}{\asymp} \frac{1}{\sqrt{n^2 p}}$$

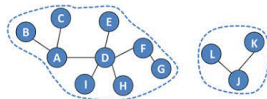
Erdos-Renyi model

ER model:

- Too simple to fit real-world data well
- But a good **null model** (“no structure/pattern”) for testing (Gao & Lafferty 2017a)
- Neat for studying some network properties
 - **Example:** average **shortest path** (Katzav et al. 2018)



- **Example:** is the network connected? (Erdos & Rényi 1960)

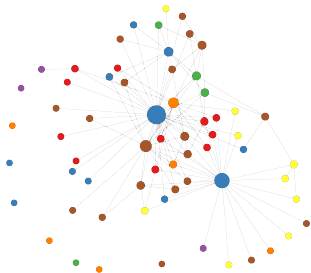


Answers to these questions are the same for: (i) ER model; and (ii) general model with all $P_{i,j} \asymp \rho_n$

- **Definition:** **degree** is the total number of edges of a node

$$d_i := \sum_{j:j \neq i} A_{i,j}$$

- Average degree: $\bar{d} := \frac{1}{n} \sum_{i=1}^n d_i \asymp \rho_n n$
- Degree describes the node's **popularity** (hub, leaf)



Degree-driven network model:

- Network structure entirely determined by degrees
- Network \rightarrow degrees: easy to compute
- Degree sequence $\xrightarrow{?}$ network model

Configuration model (Chung & Lu 2002):

- Not a probabilistic model for “network population”
- Given a degree sequence d_1, \dots, d_n , generate network
 - 1 Set $S := \{1, \dots, 1, 2, \dots, 2, \dots, n, \dots, n\}$, each i repeats d_i times
 - 2 Randomly select two entries from S , make an edge, delete them from S
 - 3 Repeat step 2 until S is exhausted

β -model (Chatterjee et al. 2011):

- **Formula:**

$$P_{i,j} = \frac{e^{\beta_i + \beta_j}}{1 + e^{\beta_i + \beta_j}}$$

Parameters $\beta_i \in \mathbb{R}$

- **Estimation: MLE**, negative log-likelihood:

$$L(\beta) := \sum_{1 \leq i < j \leq n} \log(1 + e^{\beta_i + \beta_j}) - \sum_{i=1}^n \beta_i d_i$$

Then

$$\hat{\beta} := \arg \min_{\beta} L(\beta)$$

- **Consistency:** if true β_i 's are $O(1)$, then $\|\hat{\beta} - \beta\|_{\infty} \xrightarrow{P} 0$

β -model:

$$P_{i,j} = \frac{e^{\beta_i + \beta_j}}{1 + e^{\beta_i + \beta_j}}$$

$$L(\beta) := \sum_{1 \leq i < j \leq n} \log(1 + e^{\beta_i + \beta_j}) - \sum_{i=1}^n \beta_i d_i$$

Pros and cons:

- (+) Scalable to very large and sparse networks (Shao et al. 2023)
- (+) Privacy protection (only publicize degrees) (Karwa & Slavković 2016)
- (–) Limited expressivity

Stochastic block model (SBM)

- Nodes are partitioned into K communities (blocks, groups)
- Formula** (Holland et al. 1983)

$$P_{i,j} := \rho_n P_{c_i, c_j}$$

c_i : which community node i belongs to

ρ_n : network sparsity rescaler

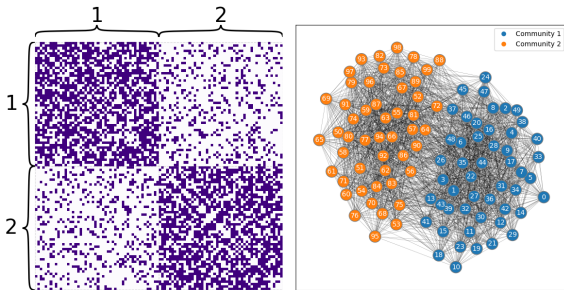


Figure: Left: heatmap of adjacency A ; right: network plot (nodes have been sorted)

Stochastic block model (SBM)

SBM:

$$P_{i,j} := \rho_n B_{c_i, c_j}$$

where B : community-level edge probability matrix

- **Estimation:** MLE:

$$L(c, B) := \sum_{1 \leq i < j \leq n} \left\{ A_{i,j} \log B_{c_i, c_j} + (1 - A_{i,j}) \log(1 - B_{c_i, c_j}) \right\}$$

$$(\hat{c}, \hat{B}) := \arg \max_{c_i \in [1:K], B \in [0,1]^{K \times K}} L(c, B)$$

- Estimating c_{true} is community detection (Abbe 2018)
- If $\rho_n \gg n^{-1} \log n$, then MLE is consistent: as $n \rightarrow \infty$,
 - $\hat{c} = c_{\text{true}}$ a.s. (Bickel & Chen 2009)
 - $\hat{B} \xrightarrow{P} B_{\text{true}}$

Question: How to understand this requirement on ρ_n ?

Stochastic block model (SBM)

$$L(c, B) := \sum_{1 \leq i < j \leq n} \left\{ A_{i,j} \log B_{c_i, c_j} + (1 - A_{i,j}) \log(1 - B_{c_i, c_j}) \right\}$$

- Exact MLE **infeasible** (combinatorial optimization for c)
- In practice: Tabu search (**Zhao et al. 2012**)
 - 1 Initialize c and B
 - 2 Iterate until convergence:
 - For $i \in [1 : n]$: update c_i
 - Update B

Stochastic block model (SBM)

SBM:

$$P_{i,j} := \rho_n P_{c_i, c_j}$$

MLE is too slow, can we do better?

- **Alternative formula:**

$$P = \rho_n Z B Z^T$$

where **membership matrix** Z may look like:

$$\begin{pmatrix} 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 \\ \vdots & \vdots & \vdots & \vdots \\ 0 & 1 & 0 & 0 \end{pmatrix}$$

corresponding to $c = (1, 1, 4, \dots, 2)$, i.e., each row of Z is a one-hot vector indicating membership

- Verify: $P_{i,j} = \rho_n Z_{i,\cdot} B Z_{j,\cdot}^T = \rho_n B_{c_i, c_j}$

Stochastic block model (SBM)

SBM (matrix form):

$$P_{n \times n} = \rho_n Z_{n \times K} B_{K \times K} (Z^T)_{K \times n}$$

Therefore:

- P is rank- K
- **SVD** of P :

$$P = USU^T$$

where $S : K \times K$ diagonal; $U : n \times K$ orthonormal: $U^T U = I$

- U only has K **different rows**, one for each community (Qin & Rohe 2013)
- cluster rows of $U \Rightarrow$ true community labels

Stochastic block model (SBM)

SBM (matrix form):

$$P = \rho_n Z B Z^T$$

In practice, we only observe A ...

- Suppose we know K
- **Leading- K SVD** of A :

$$A \approx \hat{U} \hat{S} \hat{U}^T$$

(network sparsity absorbed into \hat{S})

- Cluster the rows of $\hat{U} \Rightarrow$ community detection
- The above method is called **spectral clustering** (Lei & Rinaldo 2015)
- **Consistency:** $\rho_n \gg n^{-1} \log n \Rightarrow$ misclassification rate $\xrightarrow{p} 0$

Stochastic block model (SBM)

SBM:

$$P_{i,j} := \rho_n P_{c_i, c_j}$$

- We only explained “block model”
- What about “stochastic”?
- **Two-stage** generation (Bickel & Chen 2009):
 - 1 “Stochastic”: each node chooses its community:

$$c_1, \dots, c_n \stackrel{\text{i.i.d.}}{\sim} \text{Multinomial}(q_1, \dots, q_K)$$

- 2 “Block model”:

$$P_{i,j} := \rho_n P_{c_i, c_j}$$

- Stochastic or not doesn't matter so much in practice

Stochastic block model (SBM)

Arguably the “most famous” network model

- Simple, yet expressive
- Communities are very interpretable

Examples:

- social circles (friendship network) (Yang et al. 2013)
 - functional regions (brain image network) (Wu et al. 2022)
 - research areas (citation network) (Ji et al. 2022)
- Towards general models: SBM with growing K

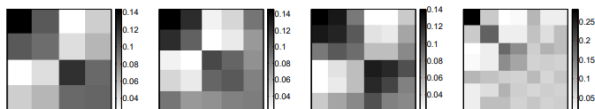
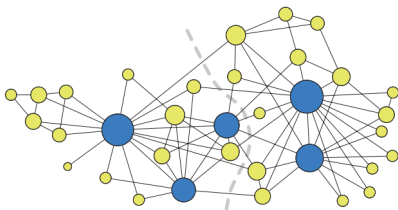


Figure: Choi et al. (2012): as K increases, SBM captures more structural details

Degree-corrected SBM



- What may go wrong with SBM?
- Why? (Recall spectral clustering ...)
- In community detection, we want to **adjust for degree heterogeneity**

Degree-corrected SBM

- **Formula:**

$$P_{i,j} = \rho_n \theta_i \theta_j B_{c_i, c_j}$$

where θ_i : degree correction

- **Matrix form:**

$$P = \rho_n \Theta Z B Z^T \Theta$$

where $\Theta = \text{diag}(\theta_1, \dots, \theta_n)$

- **Identifiability:** require $\sum_{i:c_i=k} \theta_i = 1$ for each community k

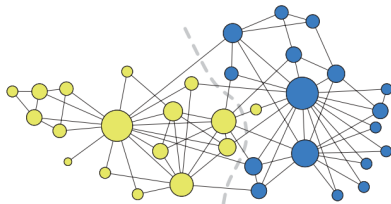


Figure: Karrer & Newman (2011): DCSBM adjusts for degrees

Degree-corrected SBM

- **Spectral clustering** (Von Luxburg 2007):

- 1 Compute Laplacian:

$$L := D^{-1/2}AD^{-1/2}$$

where $D = \text{diag}(d_1, \dots, d_n)$

Interpretation: reweighted edges

$$L_{i,j} = \frac{A_{i,j}}{\sqrt{d_i d_j}}$$

(how much attention do we pay to our relationship?)

- 2 Leading- K SVD of L :

$$L \approx \widehat{U}\widehat{S}\widehat{U}^T$$

- 3 Cluster rows of \widehat{U}

Degree-corrected SBM

The Spectral Embedding Algorithm

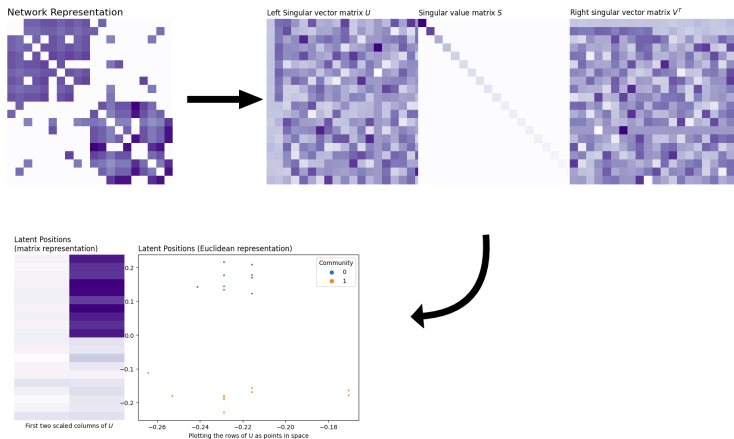


Figure: Illustration of spectral community detection for SBM (Bridgeford et al. 2022)

Quick recap

Which methods have we seen so far?

- **Moment estimator** for ER model
- **MLE** for β -model and SBM
- **Spectral method** for SBM

More about SBM/DCSBM

Overlapping communities:

- **Airoldi et al. (2008) Mixed-Membership Stochastic Blockmodel (MMSB):**

- 1 Prior step:

$$\vec{\pi}_i \stackrel{\text{i.i.d.}}{\sim} \text{Dirichlet}$$

- 2 For each (i, j) , draw $c_i \sim \text{Multinom.}(\vec{\pi}_i)$, do the same for j .
Then generate $P_{i,j}$:

$$P_{i,j} = B_{c_i, c_j}$$

* **Parameter estimation:** EM algorithm

- **Zhang et al. (2020) Overlapping Continuous Community Assignment Model (OCCAM):**

- **Matrix form:**

$$P = \rho_n \Theta Z B Z^T \Theta$$

but now Z may have continuous rows, satisfying $\|Z_{i,\cdot}\|_2 = 1$

Similarly, (Jin et al. 2024): $\|Z_{i,\cdot}\|_1 = 1$

- **Parameter estimation:** spectral clustering

More about SBM/DCSBM

Other estimation methods:

- Modularity (Girvan & Newman 2002, Newman 2006)

$$\frac{1}{2m} \sum_{i < j} \left(A_{i,j} - \frac{d_i d_j}{2m} \right) \mathbb{1}_{[c_i = c_j]}$$

m : total number of edges

- Analytical approximate EM algorithm (Ball et al. 2011)
- Semi-definite programming (SDP)

$$\arg \max_X \langle A, X \rangle, \quad \text{s.t., } X_{ii} \equiv 1, X \succeq 0$$

Then spectral cluster X (Cai & Li 2015, Amini & Levina 2018)

- “SCORE” (Jin 2015)
- Spectral clustering + majority vote refinement (Gao et al. 2018)

More about SBM/DCSBM

Extension: popularity-adjusted SBM (Sengupta & Chen 2018):

- **Motivation:** overall sparse but locally dense networks
DCSBM cannot describe this scenario:
 - $i, j \in$ same community k
 - i is globally more popular than j
 - ... but not among members of some community ℓ
- **Formulation:** for $i \in$ community k , $j \in$ community ℓ :

$$P_{i,j} = \Lambda_i^{(\ell)} \Lambda_j^{(k)}$$

where Λ encodes:

- community-level edge probabilities
- node's popularity for each (k, ℓ)

You can think: $\Lambda_i^{(\ell)} = \theta_i^{(\ell)} \sqrt{\rho_n B_{k,\ell}}$ (recall $c_i = k$)

- **Reduce to DCSBM:**

$$\Lambda_i^{(\ell)} = \theta_i \sqrt{\rho_n B_{k,\ell}} \quad \Rightarrow \quad P_{i,j} = \rho_n \theta_i \theta_j B_{k,\ell}$$

- MLE for Λ 's \Rightarrow community detection

More about SBM/DCSBM

popularity-adjusted SBM (Cont'd)

$$P_{i,j} = \Lambda_i^{(\ell)} \Lambda_j^{(k)}$$

This is subtle: MLE for Λ 's \Rightarrow community detection

- MLE gives you all $\Lambda_i^{(\ell)}$'s ..
- ... but that's insufficient for determining $P_{i,j}$
- For each (i, j) :

$$P_{i,j} = \Lambda_i^{(?)} \Lambda_j^{(?)}$$

Random dot-product model (RDPG)

- Generalization of SBM/DCSBM (Young & Scheinerman 2007)
- **Formula:** recall DCSBM (no ρ_n):

$$P = \Theta Z B Z^T \Theta$$

Suppose $B \succ 0$ (diagonal all positive), then

$$P = (\Theta Z B^{1/2}) \cdot (\Theta Z B^{1/2}) =: X X^T$$

where $X \in \mathbb{R}^{n \times K}$: latent space matrix

- In other words:
 - Each node has an **embedding** $X_{i,\cdot}$.
 - Edge probability: $P_{i,j} = \langle X_{i,\cdot}, X_{j,\cdot} \rangle$
- **Estimation:** can estimate X up to a rotation/reflection O :
Under mild conditions, \hat{X} by SVD: $\|\hat{X} - X O\|_{2 \rightarrow \infty} \xrightarrow{P} 0$ (Cape et al. 2019); CLT (Athreya et al. 2022)

Random dot-product model (RDPG)

Approximately low-rank structure is ubiquitous

- Udell & Townsend (2019): for any A

$$\inf_{\text{rank}(\hat{A}) \leq C \log n / \varepsilon^2} \|A - \hat{A}\|_{\max} \leq \varepsilon \|A\|$$

- Xu (2018): Holder smoothness in graphon (will explain later) \Rightarrow polynomial eigenvalue decay

$$\sum_{i>r} \lambda_i^2 = O(r^{-\beta} + n^{-1})$$

Random dot-product model (RDPG)

$$P = XX^T, \quad A \approx \hat{X}\hat{X}^T, \quad \hat{X} \approx XO$$

Assortativity constraint?

- Generalized RDPG ([Rubin-Delanchy et al. 2022](#))

$$P = X \begin{pmatrix} I_p & 0 \\ 0 & -I_q \end{pmatrix} X^T$$

Estimation: still SVD A :

$$A = \hat{U}\hat{S}\hat{V}^T$$

compare the signs of columns: $\hat{U}_{\cdot,i}$ vs $\hat{V}_{\cdot,i}$ to know sign

Random dot-product model (RDPG)

$$P = XX^T, \quad A \approx \hat{X}\hat{X}^T, \quad \hat{X} \approx XO$$

Interpretability

- Rows i of \hat{X} : embedding of node i

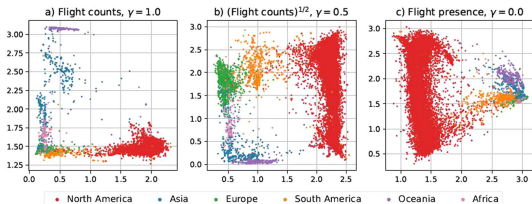


Figure: Gallagher et al. (2024): embedding of airports in flight network

- However, this embedding is limited to that particular network, due to the unknown O
Cannot be directly used for comparing nodes' roles **across** networks

Latent space models

Distance model (Hoff et al. 2002)

- **Formulation:**

$$\text{Logit}(P_{i,j}) := \alpha + \beta x_{i,j} + \|z_i - z_j\|$$

where

- $x_{i,j}$: observable edge covariate
(two node covariates \rightarrow edge covariate)
- z_i : latent space position
- **Identifiability:** $\sum_{i=1}^n z_i = \vec{0}$
- **Estimation:** Variational EM

Multiplicative Z term (Ma et al. 2020)

- **Formulation:**

$$\text{Logit}(P_{i,j}) := \alpha + \beta x_{i,j} + \langle z_i, z_j \rangle$$

Identifiability condition omitted here

- **Estimation:** (regularized) MLE by (projected) gradient

Exponential random graph models (ERGM) (Hunter et al. 2008)

- Directly model the distributions of subgraphs (motifs)
- Dependent edge generation
- Not a convenient generative model (model $\xrightarrow{?}$ data)

Formula:

- Set of motifs (subgraphs) $\mathcal{R}_1, \dots, \mathcal{R}_m$

Examples: edge, triangle, three-star

- Given an adjacency matrix a , let $R(a)$ count the frequency of \mathcal{R} in a
- Network likelihood (Hunter & Handcock 2006):

$$\mathbb{P}(A = a) \propto \exp \left\{ \sum_{i=1}^m \theta_i R_i(a) \right\}$$

To fit an ERGM to data:

- Choose the set of motifs, compute $R_i(a)$'s
- Estimate θ_i 's

$$\mathbb{P}(A = a) \propto \exp \left\{ \sum_{i=1}^m \theta_i R_i(a) \right\}$$

Given θ , how to generate data?

- Start with an initial adjacency
- Each time, decide whether to flip an edge (i, j) , resulting A (before flip) $\rightarrow A'$ (after flip)
- **Metropolis-Hastings** (Liu & Liu 2001):
accept the flip w.p.

$$\min \left(1, \exp \left[\sum_i \theta_i \{ R_i(A') - R_i(A) \} \right] \right)$$

- Iterate many rounds (with burn-in and thinning)
 - Burn-in: discard the first few rounds (e.g., first 1000 rounds)
 - Thinning: take data from every few rounds, as if they were i.i.d.

$$\mathbb{P}(A = a) \propto \exp \left\{ \sum_{i=1}^m \theta_i R_i(a) \right\}$$

Given $A = a$, how to estimate θ ?

- MLE is hard, don't know normalizing constant
- Score equation:

$$\nabla \ell(\theta) = 0,$$

that is, for each i ,

$$R_i(a) = \mathbb{E}[R_i(A)]$$

- Robbins–Monro update (**Snijders 2002**):

$$\theta_i^{(n+1)} = \theta_i^{(n)} + \alpha_n \{ R_i(a) - \widehat{\mathbb{E}}^{\theta^{(n)}} [R_i(A)] \},$$

where: $\widehat{\mathbb{E}}[R_i(A)]$ evaluated by MH-MC

step size α_i : $\sum_i \alpha_i = \infty$, $\sum_i \alpha_i^2 < \infty$

Graphon model

Overview:

- A very general framework
- Contains the aforementioned models as special cases

Two-stage network generation:

- ① Latent node positions: $X_1, \dots, X_n \stackrel{\text{i.i.d.}}{\sim} \text{Uniform}[0, 1]$
- ② Latent graphon function:

$$P_{i,j} = f(X_i, X_j) \tag{1}$$

Understandings:

- X_i describes “what kind of node”
- f captures all structural information in the network

Graphon model

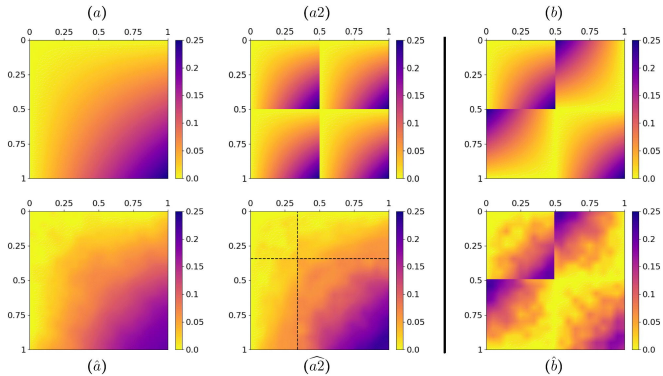


Figure: Illustration of some graphons (Sischka & Kauermann 2025)

Examples:

- ER model: $f(x, y) \equiv p$
- β -model:

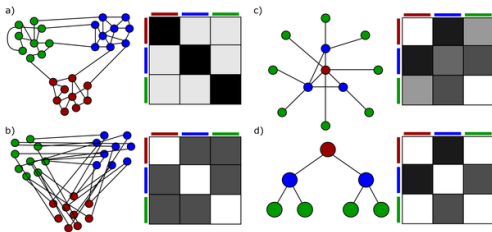
$$\beta_i = g(X_i)$$
$$f(x, y) = \text{Logit}^{-1}(g(X_i) + g(X_j))$$

for proper $g(\cdot)$

Graphon model

Examples: SBM

- K communities \Leftrightarrow partition $[0, 1]$ into K sub-intervals
- $X_i \in k\text{th interval} \Leftrightarrow i \in \text{community } k$
- Graphon f : piece-wise constant



Aldous-Hoover theorem (Zhao 2023)

- **Exchangeable network**: for any permutation $\pi : \mathbb{N} \rightarrow \mathbb{N}$,

$$(A_{i,j}) \stackrel{d}{=} (A_{\pi(i),\pi(j)})$$

- Aldous-Hoover: Exchangeable network has the representation

$$A_{i,j} \sim \text{Bernoulli}(f(X_i, X_j)),$$

for $X_i \stackrel{\text{i.i.d.}}{\sim} \text{Uniform}[0, 1]$

- Exchangeability \Rightarrow graphon model
- All nodes i.i.d. choose their roles \Rightarrow exchangeability

Examples:

- RDPG with i.i.d. latent positions \Rightarrow graphon model
- Latent space model with i.i.d. latent positions \Rightarrow graphon model
- Recall ERGM:

$$\mathbb{P}(A = a) \propto \exp \left\{ \sum_{i=1}^m \theta_i R_i(a) \right\}$$

where each $R_i(a)$: some motif count

Chatterjee & Diaconis (2013): ERGM \approx graphon model (as $n \rightarrow \infty$)

Graphon model

$$X_1, \dots, X_n \stackrel{\text{i.i.d.}}{\sim} \text{Uniform}[0, 1]$$
$$P_{i,j} = f(X_i, X_j)$$

Graphon estimation:

- **Cannot** estimate f or X_i 's (not identifiable)
- May estimate $P_{i,j}$'s
- Smoothness assumption on f [in equiv. class]
- How to estimate $P_{i,j}$'s?

Graphon model

$$X_1, \dots, X_n \stackrel{\text{i.i.d.}}{\sim} \text{Uniform}[0, 1]$$
$$P_{i,j} = f(X_i, X_j)$$

Graphon estimation:

- **Method 1:** universal singular value thresholding (USVT) (Chatterjee 2015)

① SVD A :

$$A = \widehat{U} \widehat{S} \widehat{V}^T, \quad (S \text{ diag. sorted})$$

② Keep singular values above $C\sqrt{n}$ for $C > 2$

Say, keep the first $K \Rightarrow (\widetilde{U}, \widetilde{S}, \widetilde{V}) = (\widehat{U}_{\cdot, 1:K}, \widehat{S}_{1:K, 1:K}, \widehat{V}_{\cdot, 1:K})$

③ Estimator:

$$\widehat{P} := \widetilde{U} \widetilde{S} \widetilde{V}^T$$

Achieves SOTA MSE (Xu 2018, Luo & Gao 2024)

Graphon model

$$X_1, \dots, X_n \stackrel{\text{i.i.d.}}{\sim} \text{Uniform}[0, 1]$$

$$P_{i,j} = f(X_i, X_j)$$

Graphon estimation:

- **Method 2: Neighborhood smoothing (NS)** (Zhang et al. 2017)

- 1 Dissimilarity measure of graphon slices $f(\cdot, X_i)$ and $f(\cdot, X_j)$:

$$d(i, j) := \max_{i' \neq i, j} |\langle A_{i, \cdot} - A_{j, \cdot}, A_{i', \cdot} \rangle| / n$$

- 2 For each node i , set neighborhood $\mathcal{N}_i := \{j : \text{smallest } h = \log n / n \text{ proportion of } d(i, j) \text{'s}\}$
- 3 Estimator:

$$\hat{P}_{i,j} := \frac{1}{|\mathcal{N}_i|} \sum_{i' \in \mathcal{N}_i} A_{i',j}$$

Achieves state-of-the-art “ $2 \rightarrow \infty$ error”; explicit formula

Selected research topics

Overview

We will cover these selected topics

- Method-of-moments
- Goodness-of-fit tests
- Cross-validation
- Graph matching
- Extensions to weighted, directed, bipartite graphs
- Other topics (briefly mention)

Method-of-moments

Network moments

- **Definition:** a **network moment** indexed by a motif R is

$$\hat{U}_n := \binom{n}{r}^{-1} \sum_{i_1 < \dots < i_r} \mathbb{1}_{[A_{i_1, \dots, i_r} \cong R]}$$

- **Examples:** triangle count

$$\hat{U}_n = \binom{n}{3}^{-1} \sum_{i < j < k} A_{i,j} A_{j,k} A_{k,i}$$

- **CLT:**

$$\frac{\hat{U}_n - \mathbb{E}[U_n]}{\hat{S}_n} \xrightarrow{p} N(0, 1),$$

where S_n is some variance estimator (Bickel et al. 2011, Zhang & Xia 2022)

- Moments can be used to identify network models
Application: network two-sample test

Goodness-of-fit tests

Is the network “structureless”? (Gao & Lafferty 2017b)

- $H_0 : P_{i,j} \equiv p$ (ER as null model)
- H_a : non-constant graphon model
- **Method-of-moment test**: null distribution:

$$n^{3/2}g(p) \cdot (T_2, T_3)^T \xrightarrow{d} N(0, I_2), \quad (2)$$

where $T_2 = E^3 - T$ and $T_2 = 3E^2(1 - E) - V$, here:

- E counts edges
- T counts triangles
- V counts V-shapes

$g(\cdot)$ is known function; easy to estimate p

Goodness-of-fit tests

GoF test for SBM (Lei 2016)

- $H_0 : K = K_0$ vs $H_a : K > K_0$
- Based on observed A , fit SBM with $K = K_0$, get \hat{P}
- **Idea:** is the residual network “pure noise”?
- Test statistic:

$$\tilde{A}_{i,j} := \frac{A_{i,j} - \hat{P}_{i,j}}{\sqrt{(n-1)\hat{P}_{i,j}(1-\hat{P}_{i,j})}}, \quad i \neq j$$

Then under H_0 ,

$$n^{2/3} \cdot \{ \pm \lambda_1(\tilde{A}) - 2 \} \xrightarrow{d} \text{Tracy-Widom}(1),$$

where $\lambda_1 \geq \dots \geq \lambda_n$ are sorted eigenvalues

- **Application: decide K**

Goodness-of-fit tests

Key to build valid GoF method:

- Number of continuous parameters is small under H_0
- **Example:** ER model has 1 parameter
- **Example:** SBM has $O(n + K^2)$, but the n memberships can be perfectly recovered for not-very-sparse networks, so we just deal with $O(K^2)$ parameters
- **Example:** DCSBM (Zhang & Amini 2023, Jin et al. 2025)
 H_0 : model is an SBM (no degree corrections)

Goodness-of-fit tests

No known method to test GoF for:

- β -model
- Latent space models, including RDPG
- General graphon model

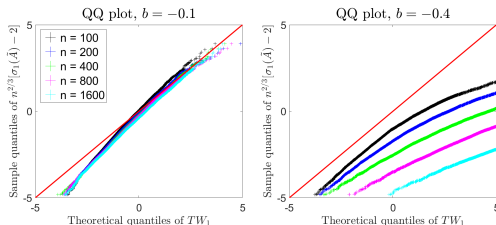


Figure: Shao et al. (2023): exactly mimicking Lei (2016) for β -model won't work (each curve)

Cross-validation

CV for network data?

- **Motivation:** select tuning parameter(s)
Example: number of communities K in SBM
Example: bandwidth h in neighborhood smoothing
- **Challenge:** we may observe only one A , how to split data?

Method 1 (node-split CV) (Chen & Lei 2018)

1 Write

$$A = \begin{pmatrix} A^{(11)} & A^{(12)} \\ A^{(21)} & A^{(22)} \end{pmatrix}$$

- 2 Use $[A^{(11)}, A^{(12)}]$ to **train** model (say, SBM)
- 3 **Validation:** look at how well fitted model predicts $A^{(22)}$
(beware $A^{(21)} = \{A^{(12)}\}^T$ already used for training)

Method 2 (edge-splitting CV): (Li et al. 2020)

- 1 Randomly sample $M_{i,j} \stackrel{\text{i.i.d.}}{\sim} \text{Bernoulli}(q)$; set $M_{j,i} = M_{i,j}$
Training data: $\{(i, j) : M_{i,j} = 1\}$
Validation data: $\{(i, j) : M_{i,j} = 0\}$
- 2 Fit a model to training data using, say, low-rank method
May fill in zeros to validation data spots
This outputs \tilde{A} , then $\hat{A} = \tilde{A}/q$ is link predictor
- 3 See how well \hat{A} predicts A entries on validation data

Graph matching

- **Data:** two adjacency matrices $A^{(1)}, A^{(2)} \in \{0, 1\}^{n \times n}$
- **Problem:** estimate a permutation matrix Π minimizing

$$\|A^{(1)} - \Pi A^{(2)} \Pi^T\|_F^2 \quad (3)$$

- **Applications** (Conte et al. 2004, Lyzinski et al. 2015):
 - Account matching in different social networks
 - Connectome alignment in brain imaging
 - Structurally similar chemical particles
 - Network two-sample test
- **Complexity:** (3) is NP (not known to be P)
- Statistical question: if A 's are structured, would the problem be more solvable?

Graph matching

Approach 1: low-rank approximation

- Assume model is low-rank: $A \sim P = XX^T$
- Leading- k SVD:

$$A \approx \hat{X}\hat{X}^T$$

- **Davis-Kahan theorem:**

$$\hat{X} \approx XO$$

for (inestimable) orthonormal $O : O^T O = I$

- Minimize:

$$\|\hat{X}^{(1)} - \Pi \hat{X}^{(2)} O\|_F$$

over Π and O

$$\|\widehat{X}^{(1)} - \Pi \widehat{X}^{(2)} O\|_F \quad (4)$$

Approach 1: low-rank approximation (cont'd)

- joint optimization of (4) is still difficult
- [Zhang \(2018\)](#): first estimate O , then Π (no iteration!)
- **Key idea**: match row distributions, get \widehat{O} ; then easy to estimate Π
- Alternatively, [Tang et al. \(2014\)](#) uses a rotation-invariant kernel to run MMD ([Gretton et al. 2012](#)) on X -rows

Approach 2: local network moments

- **Correlated ER model:**

$$\text{Cor}(A_{i,j}^{(1)}, A_{\pi^*(i), \pi^*(j)}^{(2)}) = \rho$$

for some true permutation π^*

- **Method 1: degree profile** (Ding et al. 2021):
 - 1 For each node i , compute degree distribution of its neighbors: F_i
 - 2 Similarity matrix:

$$S_{i,j} := \text{distance}(F_i, F_j)$$

- 3 Estimate π by Hungarian algorithm

Graph matching

Approach 2: local network moments (cont'd)

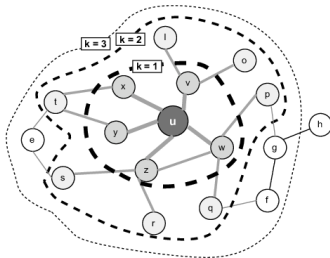
- **Method 2: large neighborhood statistics** (Mossel & Xu 2020):

- 1 Seed I_0 : a small set of correctly matched nodes
- 2 For each k and each $j_1 \in \text{net 1}$, $j_2 \in \text{net 2}$, compute

$$w_{k;j_1,j_2} := \{i : i \in I_0, i \in \{k\text{-hop neighborhoods of both } i \text{ and } j\}\}$$

k : tuning parameter; if $|w_{k;j_1,j_2}|$ is large, match (j_1, j_2)

- Many recent (strong) results (Ding & Du 2023, Mao et al. 2024) and more ...



Extensions to weighted, directed, bipartite graphs

Weighted networks:

- $A_{i,j} \sim \text{Bernoulli} \Rightarrow A_{i,j} \sim \mathcal{F}_\theta$
- **Example: Poisson β -model** (Yan et al. 2016)

$$\mathbb{P}(A_{i,j} = x) = \frac{e^{(\beta_i + \beta_j)x}}{x!} e^{-e^{\beta_i + \beta_j}}$$

- **Example: weighted SBM** (Xu et al. 2020)

$$A_{i,j} \sim \mathcal{F}_{c_i, c_j}$$

There are $\binom{K}{2}$ different \mathcal{F} 's

- Typically the same methods (MLE, spectral, ...) work for the weighted version

Extensions to weighted, directed, bipartite graphs

Directed networks:

- Each node may have two roles: **sender** and **receiver**
- **Example: directed SBM** (Wang & Wong 1987)

$$\mathbb{P}(A_{i,j} = 1) = P_{c_i, d_j},$$

where

- c_i : community label as **sender**
- d_j : community label as **receiver**
- Spectral clustering:

$$A \approx \hat{U} S \hat{V}^T,$$

then:

- Cluster rows of \hat{U} to estimate \vec{c}
- Cluster rows of \hat{V} to estimate \vec{d}

Extensions to weighted, directed, bipartite graphs

Bipartite networks:

- Two node sets of distinct meanings
Example: recommender system: buyers \leftrightarrow goods
Example: drug-target network
- A is $m \times n$
- Method and theory: mostly similar to directed networks
Some differences:
 - Unlike directed networks, no self-loops
 - Asymptotics: m, n may grow at different rates

Other topics

Resampling networks

Problem description:

- **Observe:** one A
- **Goal:** sampling distribution of some model functional (e.g., triangle count)

Recall i.i.d. case: observe X_1, \dots, X_n , goal: sample mean

- **Parametric bootstrap:** use X to fit a model $F_{\hat{\theta}}$, resample from F , repeatedly get X_1^*, \dots, X_n^* and bootstrapped \bar{X}^*
- **Non-parametric bootstrap:** resample from X , repeatedly get X_1^*, \dots, X_n^* and bootstrapped \bar{X}^*
- These two methods produce similar \bar{X}^* distributions

Resampling networks

Network? ... observe A , goal: triangle count

- **Method 1:** use A to fit \hat{P} , then repeatedly generate A^* from \hat{P} and compute triangle count
- **Method 2** (Green & Shalizi 2017): repeatedly resample nodes, each time induce A^* , count triangles
Specifically, k_1^*, \dots, k_n^* resampled from $[n]$, then

$$A_{i,j}^* := A_{k_i, k_j}$$

- Methods 1 and 2 may give you **very different** triangle count sampling distributions!

What happened?

- **Method 1.5:** use A to fit \hat{P} , then still resample nodes, each time induce \hat{P}^* , then sample A^* from \hat{P}^* , count triangles
- Method 1.5 \approx Method 2 \neq Method 1
- Reading: Zhang & Xia (2022)

Dynamic networks

What do data look like?

- **Scenario 1: network snapshots:** $\{A^{(t)}\}_{t=1}^T$ (Sewell & Chen 2015, Pensky 2019)

Examples: friendship, international trade

- **Scenario 2: stream of edges:** (i, j, t) , where t : time stamp (Less studied) (Perry & Wolfe 2013, Fang et al. 2024)

Examples: emails, contact (epidemiology)

Network snapshots:

- Build model for each layer
- Parameters evolve over time / across layers
- Different layers may have similar/shared parameters
- Tensor analysis (Dr. Xia's tutorial tomorrow)
- Some works: convert edge stream to snapshots

Nodal covariates

Sometimes you observe more than a network ...

- **Data:** $A \in \mathbb{R}^{n \times n}$ and $X \in \mathbb{R}^{n \times p}$
- **Examples:** individual covariates (gender, age, job, ...)
- Structure of X may relate to structure of A

Use X to help modeling A

Example: covariate-assisted community detection
(Binkiewicz et al. 2017)

- **Method:**

Apply spectral clustering to: $A + \lambda XX^T$

λ : tuning parameter

- Edge covariates can be utilized similarly

Nodal covariates

Use A to model/predict X

Example: network regression (Le & Li 2022)

- **Data:** on each node, observe (X_i, Y_i)
- **Goal:** use X and A to predict Y
- **Model:** $C_X := \text{ColumnSpace}(X)$, $C_P := \text{ColumnSpace}(P)$,

$$\mathbb{E}[Y|X] = \underbrace{X\beta}_{\in C_X} + \underbrace{X\theta}_{\in C_X \cap C_P} + \underbrace{\alpha}_{\in C_P}$$

and $(X\beta) \perp (X\theta)$ and $(X\theta) \perp \alpha$

- **Estimation strategy:** first estimate θ

Nodal covariates

Some understandings

- Beware of differences in X and A structures (considering X might or might not help A 's model fitting) (Zhang et al. 2016, Deshpande et al. 2018)
- Be careful when building joint models
Example: shall we worry about this model for joint community detection?

$$P_{i,j} \propto \exp \left\{ B_{c_i, c_j} + C_{X_i, X_j} \right\}$$

where suppose each $X_i \in [K]$, here

- c_i : network community label
- X_i : covariate community label

Sampling network

What if we have to explore the network?

- We might not be able to sample nodes i.i.d. from {all kinds of nodes}
- **Example (snowball):** start from an initial set of people, ask them to refer friends to our study, and friends ask their further friends, ...
- Study strategy:
 - 1 Model the true (complete) network
 - 2 Specify a sampling scheme
 - 3 Analyze what the sampling scheme will produce

Sampling network

Example: random walk (Athreya & Röllin 2016)

- Mechanism. Start from i_0 ; randomly pick a neighbor, walk to i_1 ; randomly pick a neighbor of i_1 , walk to i_2 , ...
Obtain the induced network observation
- **Theorem.** Network \sim graphon $f(u, v)$, then the network surveyed by random walk is approximately generated by

degree-weighted graphon: $f(X_i, X_j)$,

but now X_i is not Uniform $[0, 1]$, instead, has pdf:

$$p_X(u) := \frac{\int_0^1 f(u, v) dv}{\int_0^1 \int_0^1 f(u', v) du' dv} \propto \mathbb{E}[\text{degree}] \text{ at } u$$

- Many more other sampling schemes (non-backtracking, snowball, ...) (Rohe 2019)

Core periphery structure

- Aforementioned models attempt to model **all** nodes
- But often there are seclusive nodes with few connections (Rombach et al. 2014)
- **Core:** dense and structured subnetwork
- **Periphery:** the rest of network

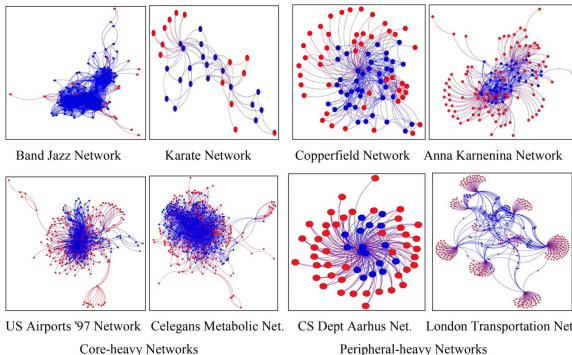


Figure: Blue: core; red: periphery (Meghanathan 2024)

Core periphery structure

Miao & Li (2023)

- Informative model for core part
- Non/Less-informative model for periphery part

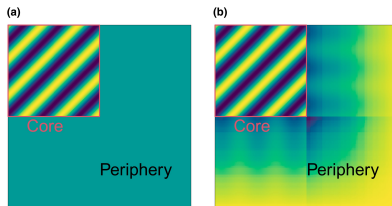


Figure: Two periphery models: left: ER; right: configuration (Miao & Li 2023)

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