

Latent variable models in biology and ecology

Chapter 4: Stochastic Block Models and Latent Block Models

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Introduction

Descriptive statistics

Probabilistic model

Inference

To go further



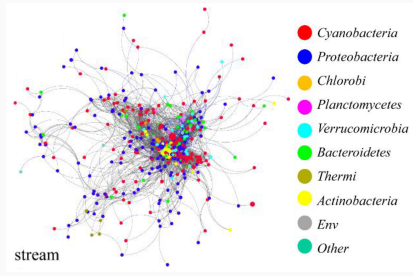
Networks can account for

- **Ecological networks** : Food web, Co-existence networks, Host-parasite interactions, Plant-pollinator interactions,
- **Social networks**
- **Inventory datasets**
- ...

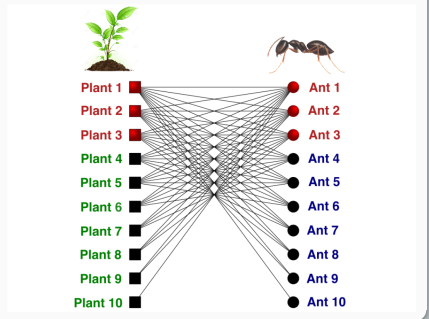
Bipartite / simple network

Networks may be or not bipartite: Interactions between nodes belonging to the same or to different functional group(s).

Simple network



Bipartite network



Terminology

A network consists in:

- nodes/vertices which represent individuals / species /ships which may interact or not,
- links/edges/connections which stand for an interaction between a pair of nodes / dyads.

A network may be

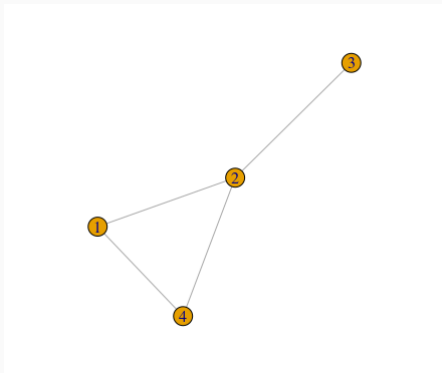
- directed / oriented (e.g. food web...),
- symmetric / undirected (e.g. coexistence network),
- with or without loops.

This distinction only makes sense for simple networks (not bipartite).

Network representation and adjacency matrix

For a non-directed network

$$Y = \begin{pmatrix} 0 & 1 & 0 & 1 \\ 1 & 0 & 1 & 1 \\ 0 & 1 & 0 & 0 \\ 1 & 1 & 0 & 0 \end{pmatrix}$$

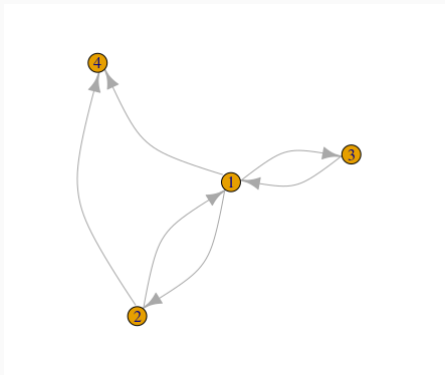


- n rows and n columns,
- symmetric matrix

Network representation and adjacency matrix

For a directed network

$$Y = \begin{pmatrix} 0 & 1 & 1 & 1 \\ 1 & 0 & 0 & 1 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \end{pmatrix}$$

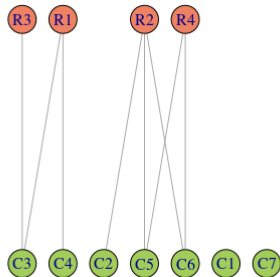


- n rows and n columns,
- non symmetric matrix

Bipartite network and incidence matrix

$$Y = \begin{pmatrix} 0 & 0 & 1 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 1 & 1 & 0 \\ 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 1 & 0 \end{pmatrix}$$

- n rows and m columns,
rectangular matrix.





- the network provided as:
 - an adjacency matrix (for simple network) or an incidence matrix (for bipartite network),
 - a list of pair of nodes / dyads which are linked.
- some additional covariates on nodes, dyads which can account for sampling effort.



- Unraveling / describing / modeling the network topology.
- Discovering particular structure of interaction between some subsets of nodes.
- Understanding network heterogeneity.
- Not inferring the network !

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Aim : give a short description of the network, give a hint about its structure, look for heterogeneity in the connections

- Many metrics supplied for simple networks
- Have been extended to bipartite networks

Example : Chilean foodweb

[Kéfi et al., 2016]



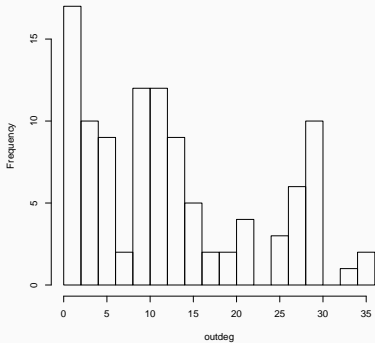
$$\begin{aligned} \deg(u) &= \sum_{v \in V} (u \leftrightarrow v), & \deg(v) &= \sum_{u \in U} (u \leftrightarrow v) \\ \deg_i &= \sum_{j=1}^{|V|} Y_{ij} & \deg_j &= \sum_{i=1}^{|U|} Y_{ij} \end{aligned}$$

- Nodes with high degree are **hubs**
- Nodes with null degree are **isolated**
- If edges are oriented : in- and out- degrees can be computed.

Degrees on Chilean foodweb

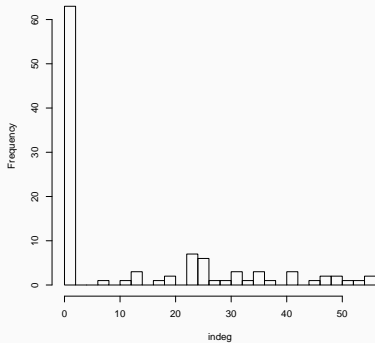
Out degree distribution

Histogram of outdeg



In degree distribution

Histogram of indeg



Closeness centrality

Property on a node

Definition

Determine whether a node can communicate with other nodes of the network directly or through the short paths.

$$C(u) = \frac{1}{\sum_{w \in U \cup V} d(u, w)}$$

where $d(u, w)$ is the length of the shortest path between u and w (through the network).

Note that, for bipartite networks

- A node $u \in U$ can have a minimum distance of 1 with $v \in V$.
- A node $u \in U$ can have a minimum distance of 2 with $u' \in U$.
- All paths between nodes of the same set are of even length.

Betweenness centrality

Property on a node

Definition

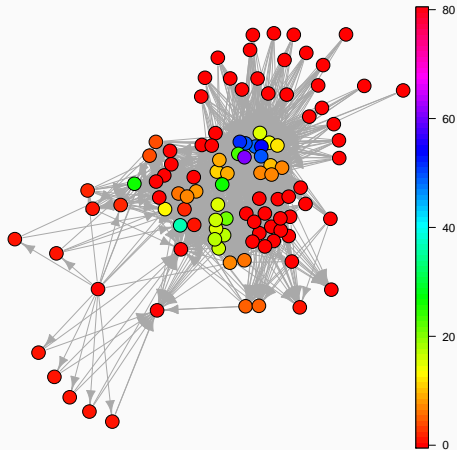
Betweenness centrality quantifies the number of times a node acts as a bridge along the shortest path between two other nodes.

The betweenness of a vertex v is computed as follows.

- For each pair of vertices (w, w') , compute the shortest paths between them. $\delta_{w,w'}$ is the number of shortest paths between (w, w')
- For each pair of vertices (w, w') , determine the fraction of shortest paths that pass through v : $\frac{\delta_{w,w'}(v)}{\delta_{w,w'}}$
- Sum this fraction over all pairs of vertices (w, w') .

$$B(v) = \sum_{w \neq w' \neq v} \frac{\delta_{w,w'}(v)}{\delta_{w,w'}}$$

Betweenness centrality



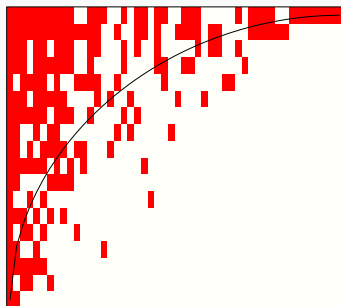
Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.000	0.000	0.000	6.604	6.929	59.570

Property on the network

Definition

- Important property in ecology
- Defined as a pattern of interactions in which specialists (e.g. pollinators that visit few plant species) interact with plants that are visited by generalists.
- Mathematically, looking for a reordering of rows and columns such that Y is nested

Nestedness



- more generally used on incidence matrices,
- significance of the nestedness index computed by random permutations of the matrix,
- this food web is found to be nested.

Property on the network

Definition

Existence of clusters (blocks, module, communities) where nodes are much more connected than with other clusters

Modularity



- | 1 | 2 | 3 | 4 |
|----|----|---|----|
| 69 | 17 | 7 | 13 |
- very low modularity.

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- Latent block models

- Some possible extensions

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Probabilistic approach

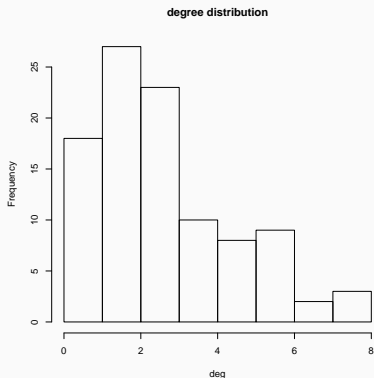
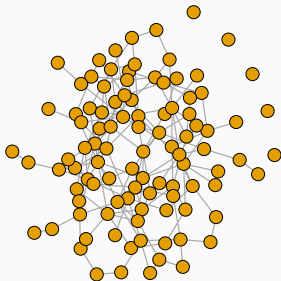
- **Context:** our matrix Y is the realization of a stochastic process.
- **Aim:** Propose a stochastic process is able to mimic heterogeneity in the connections.
- **Advantage:** benefit from the statistical tools (tests, model selection, etc...)

A first random graph model for network: null model

[Erdős and Rényi, 1959] Model for n nodes

$$\forall 1 \leq i, j \leq n, \quad Y_{ij} \overset{i.i.d.}{\sim} b(p),$$

where b is the Bernoulli distribution and $p \in [0, 1]$ a probability for a link to exist.



Limitations of an ER graph to describe real networks

- Homogeneity of the connections
- Degree distribution too concentrated, no high degree nodes,
- All nodes are equivalent (no nestedness...),
- No modularity, no hubs

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Stochastic Block Model

[Nowicki and Snijders, 2001] Let (Y_{ij}) be an adjacency matrix

Latent variables

- The nodes $i = 1, \dots, n$ are partitionned into K clusters
- $Z_i = k$ if node i belongs to cluster (block) k
- Z_i independent variables

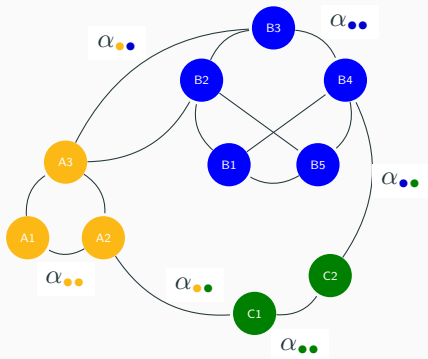
$$\mathbb{P}(Z_i = k) = \pi_k$$

Conditionally to $(Z_i)_{i=1, \dots, n} \dots$

(Y_{ij}) independent and

$$Y_{ij} | Z_i, Z_j \sim \text{Bern}(\alpha_{Z_i, Z_j}) \quad \Leftrightarrow \quad P(Y_{ij} = 1 | Z_i = k, Z_j = \ell) = \alpha_{k\ell}$$

Stochastic Block Model : illustration



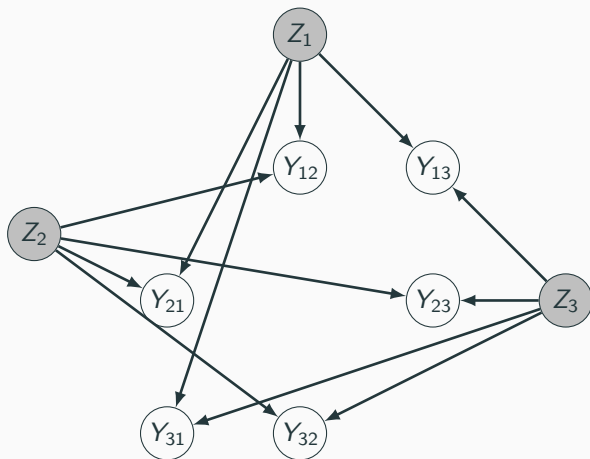
Parameters

Let n nodes divided into 3 clusters

- $\mathcal{K} = \{\bullet, \bullet, \bullet\}$ clusters
- $\pi_{\bullet} = \mathbb{P}(i \in \bullet), \bullet \in \mathcal{K}, i = 1, \dots, n$
- $\alpha_{\bullet, \bullet} = \mathbb{P}(i \leftrightarrow j | i \in \bullet, j \in \bullet)$

$$Z_i = \mathbf{1}_{\{i \in \bullet\}} \sim^{\text{iid}} \mathcal{M}(1, \pi), \quad \forall \bullet \in \mathcal{K},$$
$$Y_{ij} | \{i \in \bullet, j \in \bullet\} \sim^{\text{ind}} \mathcal{B}(\alpha_{\bullet, \bullet})$$

DAG of the model



SBM : A great generative model

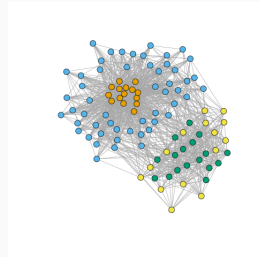
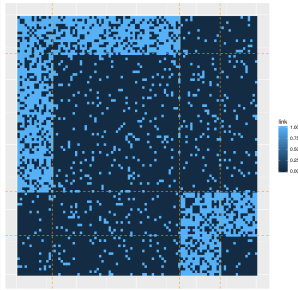
- Generative model : easy to simulate
- No a priori on the type of structure
- Combination of modularity, nestedness, etc...

Networks with hubs generated by SBM

- $\pi = c(.15, .35, .15, .35)$

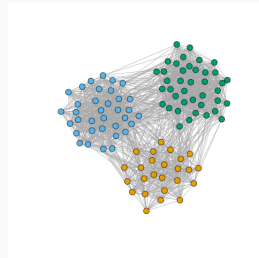
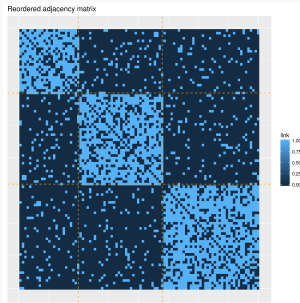
- $\alpha = \begin{pmatrix} 0.80 & 0.80 & 0.20 & 0.20 \\ 0.80 & 0.20 & 0.20 & 0.20 \\ 0.20 & 0.20 & 0.80 & 0.80 \\ 0.20 & 0.20 & 0.80 & 0.20 \end{pmatrix}$

Reordered adjacency matrix



Community network generated by SBM

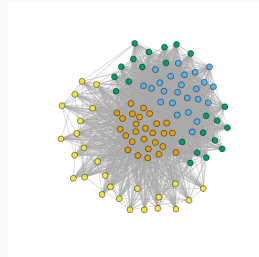
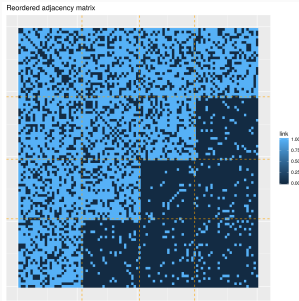
- $\pi = c(0.25, 0.35, 0.40)$
- $\alpha = \begin{pmatrix} 0.80 & 0.20 & 0.20 \\ 0.20 & 0.80 & 0.20 \\ 0.20 & 0.20 & 0.80 \end{pmatrix}$

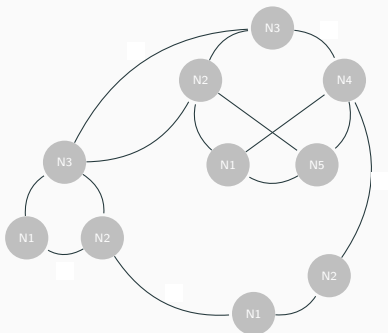


Nestedness generated by SBM

- $\pi = c(.15, .35, .15, .35)$

- $\alpha = \begin{pmatrix} 0.80 & 0.80 & 0.80 & 0.80 \\ 0.80 & 0.80 & 0.80 & 0.20 \\ 0.20 & 0.80 & 0.20 & 0.80 \\ 0.80 & 0.20 & 0.20 & 0.20 \end{pmatrix}$





Stochastic Block Model

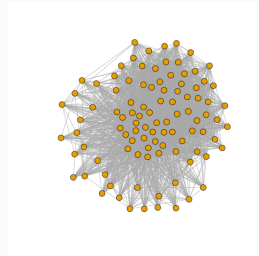
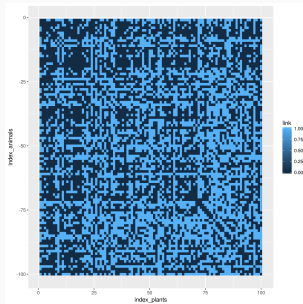
Let n nodes divided into

- $\mathcal{K} = \{\bullet, \bullet, \bullet\}$, $\text{card}(\mathcal{K})$ known
- $\pi_{\bullet} = ?$,
- $\alpha_{\bullet\bullet} = ?$

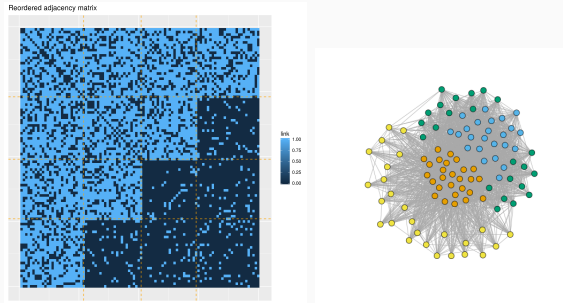
[Nowicki and Snijders, 2001], [Daudin et al., 2008]

R package: blockmodels, sbm

From....



... to



Tasks

- Find the clusters
- Find the number of clusters
- Practical implementation
- Theoretical results

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- Latent block models

- Some possible extensions

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Probabilistic model for binary bipartite networks

Let Y_{ij} be a bi-partite network. Individuals in row and cols are not the same.

Latent variables : bi-clustering

- Nodes $i = 1, \dots, n_1$ partitionned into K_1 clusters, nodes $j = 1, \dots, n_2$ partitionned into K_2 clusters
- $$\begin{aligned} Z_i^1 &= k && \text{if node } i \text{ belongs to cluster (block) } k \\ Z_j^2 &= \ell && \text{if node } j \text{ belongs to cluster (block) } \ell \end{aligned}$$
- Z_i^1, Z_j^2 independent variables

$$\mathbb{P}(Z_i^1 = k) = \pi_k^1, \quad \mathbb{P}(Z_j^2 = \ell) = \pi_\ell^2$$

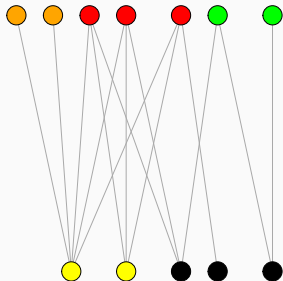
Conditionally to $(Z_i^1)_{i=1,\dots,n_1}, (Z_j^2)_{j=1,\dots,n_2} \dots$

(Y_{ij}) independent and

$$Y_{ij}|Z_i^1, Z_j^2 \sim \text{Bern}(\alpha_{Z_i^1, Z_j^2}) \quad \Leftrightarrow \quad \mathbb{P}(Y_{ij} = 1 | Z_i^1 = k, Z_j^2 = \ell) = \alpha_{k\ell}$$

[Govaert and Nadif, 2008]

Latent Block Model : illustration



Latent Block Model

- n_1 row nodes $\mathcal{K}_1 = \{\bullet, \bullet, \bullet\}$ classes
- $\pi_{\bullet}^1 = \mathbb{P}(i \in \bullet), \bullet \in \mathcal{K}_1, i = 1, \dots, n$
- n_2 column nodes $\mathcal{K}_2 = \{\bullet, \bullet\}$ classes
- $\pi_{\bullet}^2 = \mathbb{P}(j \in \bullet), \bullet \in \mathcal{K}_2, j = 1, \dots, m$
- $\alpha_{\bullet\bullet} = \mathbb{P}(i \leftrightarrow j | i \in \bullet, j \in \bullet)$

$$\begin{aligned} Z_i^1 &= \mathbf{1}_{\{i \in \bullet\}} \sim^{\text{iid}} \mathcal{M}(1, \pi^1), \quad \forall \bullet \in \mathcal{Q}_1, \\ Z_j^2 &= \mathbf{1}_{\{j \in \bullet\}} \sim^{\text{iid}} \mathcal{M}(1, \pi^2), \quad \forall \bullet \in \mathcal{Q}_2, \\ Y_{ij} \mid \{i \in \bullet, j \in \bullet\} &\sim^{\text{ind}} \text{Bern}(\alpha_{\bullet\bullet}) \end{aligned}$$

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Valued-edge networks

Values-edges networks

Information on edges can be something different from presence/absence. It can be:

1. a count of the number of observed interactions,
2. a quantity interpreted as the interaction strength,

Natural extensions of SBM and LBM

1. Poisson distribution: $Y_{ij} \mid \{i \in \bullet, j \in \bullet\} \sim^{\text{ind}} \mathcal{P}(\lambda_{\bullet\bullet})$,
2. Gaussian distribution: $Y_{ij} \mid \{i \in \bullet, j \in \bullet\} \sim^{\text{ind}} \mathcal{N}(\mu_{\bullet\bullet}, \sigma^2)$,
[Mariadassou et al., 2010]
3. More generally,

$$Y_{ij} \mid \{i \in \bullet, j \in \bullet\} \sim^{\text{ind}} \mathcal{F}(\theta_{\bullet\bullet})$$

Multiplex networks

Several kind of interactions between nodes . For instance :

- Love and friendship
- Working relations and friendship
- In ecology : mutualistic and competition

Block model for multiplex networks

$$Y_{ij} \in \{0, 1\}^Q = (Y_{ij}^a, Y_{ij}^b), \forall w \in \{0, 1\}^2$$

$$\mathbb{P}(Y_{ij}^a, Y_{ij}^b = w | Z_i = k, Z_j = \ell) = \alpha_{k\ell}^w$$

[Kéfi et al., 2016], [Barbillon et al., 2017]

In R package: `blockmodels`, `sbm` when two relations are at stake.

Remark: a particular case of multiplex network is dynamic network, [Matias and Miele, 2017].

Taking into account covariates

Sometimes covariates are available. They may be on:

- nodes,
- edges,
- both.

1. They can be used a posteriori to explain blocks inferred by SBM.
2. Extension of the SBM which takes into account covariates. Blocks are structure of interaction which is not explained by covariates !

If covariates are sampling conditions, case 2 be may more interesting.

SBM with covariates

- As before : (Y_{ij}) be an adjacency matrix
- Let $x^{ij} \in \mathbb{R}^p$ denote covariates describing the pair (i, j)

Latent variables : as before

- The nodes $i = 1, \dots, n$ are partitioned into K clusters
- Z_i independent variables

$$\mathbb{P}(Z_i = k) = \pi_k$$

Conditionally to $(Z_i)_{i=1, \dots, n} \dots$

(Y_{ij}) independent and

$$Y_{ij}|Z_i, Z_j \sim \text{Bern}(\text{logit}(\alpha_{Z_i, Z_j} + \beta \cdot x_{ij})) \quad \text{if binary data}$$

$$Y_{ij}|Z_i, Z_j \sim \mathcal{P}(\exp(\alpha_{Z_i, Z_j} + \beta \cdot x_{ij})) \quad \text{if counting data}$$

If $K = 1$: all the connection heterogeneity is explained by the covariates.

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- Parameters estimation

- Model selection

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- Selection of the number of clusters K for SBM or K_1, K_2 for LBM
- Estimation of the parameters π, θ for a given number of clusters
- Clustering $\hat{\mathbf{Z}}$

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Likelihood for SBM

For directed network .

Complete likelihood (\mathbf{Y}) et (\mathbf{Z})

$$\begin{aligned}\ell_c(\mathbf{Y}, \mathbf{Z}; \theta) &= p(\mathbf{Y}|\mathbf{Z}; \alpha)p(\mathbf{Z}; \pi) \\ &= \prod_{(i \neq j)=1}^n f_{\alpha_{Z_i, Z_j}}(Y_{ij}) \times \prod_{i=1}^n \pi_{Z_i} \\ &= \prod_{(i \neq j)=1}^n \prod_{k=1}^K \prod_{\ell=1}^K (f_{\alpha_{k, \ell}}(Y_{ij}))^{1_{Z_i=k} 1_{Z_j=\ell}} \times \prod_{i=1}^n \prod_{k=1}^K (\pi_k)^{1_{Z_i=k}}\end{aligned}$$

Marginal likelihood (\mathbf{Y})

$$\log \ell(\mathbf{Y}; \theta) = \log \sum_{\mathbf{Z} \in \mathcal{Z}} \ell_c(\mathbf{Y}, \mathbf{Z}; \theta). \quad (1)$$

$$\log \ell(\mathbf{Y}; \theta) = \log \sum_{\mathbf{Z} \in \mathcal{Z}} \ell_c(\mathbf{Y}, \mathbf{Z}; \theta).$$

Remark

$\mathcal{Z} = \{1, \dots, K\}^n \Rightarrow$ when K and n increase, impossible to compute.

Standard tool to maximize the likelihood when latent variables involved : EM algorithm.

From EM to variational EM

Standard EM

At iteration (t) :

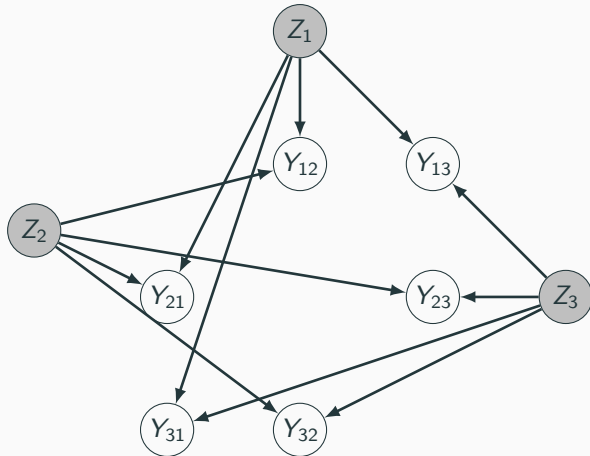
- **Step E:** compute

$$Q(\theta|\theta^{(t-1)}) = \mathbb{E}_{\mathbf{Z}|\mathbf{Y},\theta^{(t-1)}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)]$$

- **Step M:**

$$\theta^{(t)} = \arg \max_{\theta} Q(\theta|\theta^{(t-1)})$$

Limitations of standard EM



- Step E requires the computation of $\mathbb{E}_{\mathbf{Z}|\mathbf{Y}, \theta^{(t-1)}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)]$
- However, once conditioned by par \mathbf{Y} , the \mathbf{Z} are not independent anymore: complex distribution if K and n big.

Variational EM : maximization of a lower bound

Idea : replace the complicated distribution $p(\cdot|\mathbf{Y};\theta) = [\mathbf{Z}|\mathbf{Y},\theta]$ by a simpler one.

Let $\mathcal{R}_{\mathbf{Y},\tau}$ be any distribution on \mathbf{Z}

Central identity

$$\begin{aligned}\mathcal{I}_{\theta}(\mathcal{R}_{\mathbf{Y},\tau}) &= \log \ell(\mathbf{Y};\theta) - \text{KL}[\mathcal{R}_{\mathbf{Y},\tau}, p(\cdot|\mathbf{Y};\theta)] \leq \log \ell(\mathbf{Y};\theta) \\ &= \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] - \sum_{\mathbf{Z}} \mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}) \log \mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}) \\ &= \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] + \mathcal{H}(\mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}))\end{aligned}$$

Note that:

$$\mathcal{I}_{\theta}(\mathcal{R}_{\mathbf{Y},\tau}) = \log \ell(\mathbf{Y};\theta) \Leftrightarrow \mathcal{R}_{\mathbf{Y},\tau} = p(\cdot|\mathbf{Y};\theta)$$

By Bayes

$$\begin{aligned}\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta) &= \log p(\mathbf{Z}|\mathbf{Y}; \theta) + \log \ell(\mathbf{Y}; \theta) \\ \log \ell(\mathbf{Y}; \theta) &= \log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta) - \log p(\mathbf{Z}|\mathbf{Y}; \theta)\end{aligned}$$

By integration against $\mathcal{R}_{\mathbf{Y}, \tau}$:

$$\begin{aligned}\mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau}}[\log \ell(\mathbf{Y}; \theta)] &= \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau}}[\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] - \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau}}[\log p(\mathbf{Z}|\mathbf{Y}; \theta)] \\ \log \ell(\mathbf{Y}; \theta) &= \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau}}[\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] - \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau}}[\log p(\cdot|\mathbf{Y}; \theta)]\end{aligned}$$

As a consequence:

$$\begin{aligned}
 \mathcal{I}_\theta(\mathcal{R}_{\mathbf{Y},\tau}) &= \log \ell(\mathbf{Y}; \theta) - \mathbf{KL}[\mathcal{R}_{\mathbf{Y},\tau}, p(\cdot|\mathbf{Y}; \theta)] \\
 &= \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] - \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log p(\mathbf{Z}|\mathbf{Y}; \theta)] \\
 &\quad - \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} \left[\log \frac{\mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z})}{p(\mathbf{Z}|\mathbf{Y}; \theta)} \right] \\
 &= \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] - \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log p(\mathbf{Z}|\mathbf{Y}; \theta)] \\
 &\quad - \underbrace{\mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log \mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z})]}_{\mathcal{H}(\mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}))} + \mathbb{E}_{\mathcal{R}_{\mathbf{Y},\tau}} [\log p(\mathbf{Z}|\mathbf{Y}; \theta)]
 \end{aligned}$$

- Maximization of $\log \ell(\mathbf{Y}; \theta)$ w.r.t. θ replaced by maximization of the lower bound $\mathcal{I}_\theta(\mathcal{R}_{\mathbf{Y}, \tau})$ w.r.t. τ and θ .
- **Benefit** : we choose $\mathcal{R}_{\mathbf{Y}, \tau}$ such that the maximization calculus can be done explicitly
 - In our case: mean field approximation : neglect dependencies between the (Z_i)

$$P_{\mathcal{R}_{\mathbf{Y}, \tau}}(Z_i = k) = \tau_{ik}$$

Algorithm

At iteration (t), given the current value $(\theta^{(t-1)}, \mathcal{R}_{\mathbf{Y}, \tau^{(t-1)}})$,

- **Step VE** Maximization w.r.t. τ

$$\begin{aligned}\tau^{(t)} &= \arg \max_{\tau \in \mathcal{T}} \mathcal{I}_{\theta^{(t-1)}}(\mathcal{R}_{\mathbf{Y}, \tau}) \\ &= \arg \max_{\tau \in \mathcal{T}} \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau}} \left[\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta^{(t-1)}) \right] + \mathcal{H}(\mathcal{R}_{\mathbf{Y}, \tau}(\mathbf{Z})) \\ &= \arg \max_{\tau \in \mathcal{T}} \log \ell(\mathbf{Y}; \theta^{(t-1)}) - \mathbf{KL}[\mathcal{R}_{\mathbf{Y}, \tau}, p(\cdot | \mathbf{Y}; \theta^{(t-1)})] \\ &= \arg \min_{\tau \in \mathcal{T}} \mathbf{KL}[\mathcal{R}_{\mathbf{Y}, \tau}, p(\cdot | \mathbf{Y}; \theta^{(t-1)})]\end{aligned}$$

Algorithm

- **Step M** Maximization w.r.t. θ

$$\begin{aligned}\theta^{(t)} &= \arg \max_{\theta} \mathcal{I}_{\theta}(\mathcal{R}_{\mathbf{Y}, \tau^{(t)}}) \\ &= \arg \max_{\theta} \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau^{(t)}}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)] + \mathcal{H}(\mathcal{R}_{\mathbf{Y}, \tau^{(t)}}(\mathbf{Z})) \\ &= \arg \max_{\theta} \mathbb{E}_{\mathcal{R}_{\mathbf{Y}, \tau^{(t)}}} [\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta)]\end{aligned}$$

$$\tau^{(t)} = \arg \min_{\tau} \mathbf{KL}[\mathcal{R}_{\mathbf{Y},\tau}, p(\cdot|\mathbf{Y}; \theta^{(t-1)})] = \arg \max_{\tau} \mathcal{I}_{\theta^{(t-1)}}(\mathcal{R}_{\mathbf{Y},\tau}).$$

(we drop out the index $^{(t-1)}$ on θ)

$$\mathcal{I}_{\theta}(\mathcal{R}_{\mathbf{Y},\tau}) = \sum_{\mathbf{Z}} \mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}) \log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta) - \sum_{\mathbf{Z}} \mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}) \log \mathcal{R}_{\mathbf{Y},\tau}(\mathbf{Z}),$$

VE-step for SBM ii

with

$$\begin{aligned}\log \ell_c(\mathbf{Y}, \mathbf{Z}; \theta) &= \log p(\mathbf{Y}|\mathbf{Z}; \theta) + \log p(\mathbf{Z}; \theta), \\ &= \sum_{i,j=1, i \neq j}^n \log p(Y_{ij}|Z_i, Z_j; \theta) + \sum_{i=1}^n \log \pi_{Z_i}, \\ &= \sum_{i,j=1, i \neq j}^n \sum_{k,\ell=1}^K Z_{ik} Z_{j\ell} \log p(Y_{ij}|\alpha_{k\ell}) + \sum_{i=1}^n \sum_{k=1}^K Z_{ik} \log \pi_k\end{aligned}$$

Integration of the \mathbf{Z} where $\mathbf{Z} \sim \mathcal{R}_{\mathbf{Y},\tau}$

$$\mathcal{I}_\theta(\mathcal{R}_{\mathbf{Y},\tau}) = \sum_{i,j=1, i \neq j}^n \sum_{k,\ell=1}^K \tau_{iq} \tau_{j\ell} \log p(Y_{ij}|\alpha_{k\ell}) + \sum_{i=1}^n \sum_{k=1}^K \tau_{ik} \log \pi_k$$

VE-step for SBM iii

Maximization under the constraint: $\forall i = 1 \dots n, \sum_{k=1}^K \tau_{ik} = 1$.

- Derivatives of

$$\mathcal{I}_{\theta}(\mathcal{R}_{\mathbf{Y}, \tau}) + \sum_{i=1}^n \lambda_i \left[\sum_{k=1}^K \tau_{ik} - 1 \right]$$

with respect to $(\lambda_i)_{i=1 \dots n}$ and $(\tau_{ik})_{i=1 \dots n, k=1 \dots K}$ where λ_i are the Lagrange multipliers,

- Leads to collection of equations: for $i = 1 \dots n$ and $k = 1 \dots K$,

$$\sum_{\ell=1}^K \sum_{j=1, j \neq i}^n \log p(Y_{ij} | \alpha_{k\ell}) \tau_{j\ell} + \log \pi_k - \log \tau_{ik} + 1 + \lambda_i = 0,$$

- Leads to the following fixed point problem:

$$\hat{\tau}_{ik} = e^{1+\lambda_i} \alpha_k \prod_{j=1, j \neq i}^n \prod_{\ell=1}^K p(Y_{ij} | \alpha_{k\ell})^{\hat{\tau}_{j\ell}}, \quad \forall i = 1 \dots n, \forall k = 1 \dots K,$$

which has to be solved under the constraints $\forall i = 1 \dots n$, $\sum_{k=1}^K \tau_{ik} = 1$. This optimization problem is solved using a standard fixed point algorithm.

$$\theta^{(t)} = \arg \max_{\theta} \mathcal{I}_{\theta^{(t)}}(\mathcal{R}_{\mathbf{Y}, \tau^{(t)}})$$

under the constraints: $\sum_{k=1}^k \pi_k = 1$.

Maximization with respect to $\boldsymbol{\pi}$ is quite direct:

$$\hat{\pi}_q = \frac{1}{n} \sum_{i=1}^n \hat{\tau}_{ik}$$

For the Bernoulli SBM:

$$\hat{\alpha}_{k\ell} = \frac{\sum_{i,j=1, i \neq j}^n \hat{\tau}_{ik} \hat{\tau}_{j\ell} Y_{ij}}{\sum_{i,j=1, i \neq j}^n \hat{\tau}_{ik} \hat{\tau}_{j\ell}}$$

If the edge probabilities depend on covariates:

$$\text{logit}(p_{kl}) = \alpha_{kl} + \beta \cdot x_{ij},$$

then the optimization of (α_{kl}) and (β) at step M of the VEM is not explicit anymore and one should resort to optimization algorithms such as Newton-Raphson algorithm.

- Really fast
- Strongly depend on the initial values

Introduction

Descriptive statistics

Probabilistic model

Inference

Parameters estimation

Model selection

To go further

Penalized likelihood criterion

- Selection of the number of clusters K (or K_1, K_2 in the LBM)
- Integrated Classification Likelihood (ICL) [Biernacki et al., 2000]

$$ICL(\mathcal{M}_{\mathbf{K}}) = \log \ell_c(\mathbf{Y}, \hat{\mathbf{Z}}; \hat{\theta}_{\mathbf{K}}) - \text{pen}(\mathcal{M}_{\mathbf{K}}) \quad (2)$$

where

$$\hat{Z}_i = \arg \max_{k \in \{1, \dots, K\}} \hat{\tau}_{ik}. \quad (3)$$

- Integrated Complete Likelihood (ICL)

$$ICL(\mathcal{M}_{\mathbf{K}}) = \mathbb{E}_{p(\cdot | \mathbf{Y}, \hat{\theta}_{\mathbf{K}})} [\log \ell_c(\mathbf{Y}, \hat{\mathbf{Z}}; \hat{\theta}_{\mathbf{K}}) - \text{pen}(\mathcal{M}_{\mathbf{K}})] \quad (4)$$

Expression of the penalization

For SBM

$$pen_{\mathcal{M}} = \begin{cases} -\frac{1}{2} \{ (K-1) \log(n) + K^2 \log(n^2 - n) \} & \text{for directed network} \\ -\frac{1}{2} \left\{ \underbrace{(K-1) \log(n)}_{\text{Clust.}} + \frac{K(K+1)}{2} \log\left(\frac{n^2-n}{2}\right) \right\} & \text{for undirected network} \end{cases}$$

For LBM

$$pen_{\mathcal{M}} = -\frac{1}{2} \left\{ \underbrace{(K_1-1) \log(n_1) + (K_2-1) \log(n_2)}_{\text{Bi-Clust.}} + \underbrace{(K_1 K_2) \log(n_1 n_2)}_{\text{Connection}} \right\}$$

Advantages of ICL

- its capacity to outline the clustering structure in networks
- Involves a trade-off between goodness of fit and model complexity
- ICL values : goodness of fit AND clustering sharpness.

Comments on the ICL versus BIC

Conjecture

$$BIC(\mathcal{M}) = \log \ell(\mathbf{Y}; \hat{\theta}, \mathcal{M}) - \text{pen}(\mathcal{M})$$

with the same penalty

- Under this conjecture

$$\begin{aligned} ICL(\mathcal{M}) &= BIC(\mathcal{M}) + \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{Y}; \hat{\theta}_{\mathbf{K}}) \log p(\mathbf{Z}|\mathbf{Y}; \hat{\theta}_{\mathbf{K}}) \\ &= BIC(\mathcal{M}) - \mathcal{H}(p(\cdot|\mathbf{Y}; \theta)) \end{aligned}$$

- As a consequence, because of the entropy, ICL will encourage clustering with well-separated groups
-

$$\widehat{ICL}(\mathcal{M}) = BIC(\mathcal{M}) + \sum_{\mathbf{Z}} \mathcal{R}_{\mathbf{Y}}(\mathbf{Z}, \hat{\tau}) \log \mathcal{R}_{\mathbf{Y}, \hat{\tau}}(\mathbf{Z}) - \mathbf{KL}[\mathcal{R}_{\mathbf{Y}, \hat{\tau}}, p(\cdot|\mathbf{Y}; \hat{\theta})].$$

Algorithm in practice

- Going through the models and initiate VEM at the same time
- Bounds on K : $\{K_{\min}, \dots, K_{\max}\}$

Stepwise procedure

Starting from K

- **Split** : if $K < K_{\max}$
 - Maximize the likelihood (lower bound) of \mathcal{M}_{K+1}
 - K initializations of the VEM are proposed : split each cluster into 2 clusters
- **Merge** : If $K > K_{\min}$
 - Maximize the likelihood (lower bound) of model \mathcal{M}_{K-1}
 - $\frac{K(K-1)}{2}$ initializations of the VEM are proposed : merging all the possible pairs of clusters

Theoretical properties for SBM

- Identifiability and a first consistency result by [Celisse et al., 2012]
- Consistency of the posterior distribution of the latent variables [Mariadassou and Matias, 2015]
- Consistency and properties of the variational estimators [Bickel et al., 2013]

- Time evolving networks **Matias**
- Multipartite, Multiplexe networks (**R-package sbm, Bar-Hen, Barbillon, Donnet**)
- Multilevel networks (individuals and organizations) (**Chabbert-Liddell**)
- Missing data in the network,

SBM/LBM

- generative models,
- flexible,
- comprehensive models which can be linked to a lot of classical descriptors.

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