

Mixture tree model for network inference

JOBIM

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1 Motivation

2 Network inference

- General Framework
- Using trees

3 With count data

- Model
- Illustration

Context

Rising interest in **jointly analysed** species abundances:

- Metagenomics
- Microbiologie
- Ecology

Ecological network

Tool to better understand species interactions (direct/indirect, nature), eco-systems organizations (clusters ?)

Allows for resilience analyses, pathogens control, ecosystem comparison, reaction prediction...

Example

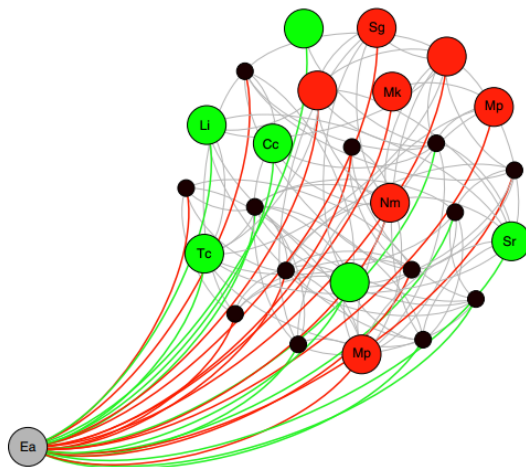


Figure: *Erysiphe althitoides* pathobiom on oak tree leaves.

Data

- **Species** : animal, bacteria, gene ...
- **Abundances** : ecological counts, Next-Generation Sequencing technologies...
- **Covariates** : coverage, temperature, water depth ...

Repeated signal : n samples of p abundances.

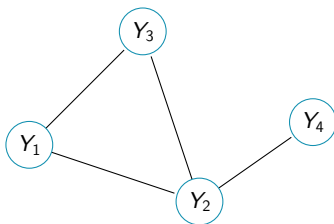
Notations

$$Y = [Y_{ij}]_{(i,j) \in \{1, \dots, n\} \times \{1, \dots, p\}}$$

- Y_{ij} : abundance of the j^{th} species in the i^{th} sample

Infer the species interaction network from Y

Graphical models



- All variables are dependant
- Some are **conditionnally independant** (i.e. indirectly dependant)

$$Y_4 \perp\!\!\!\perp (Y_1, Y_3) | Y_2$$

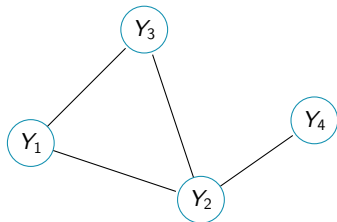
Graphical models

Factorization propriety

The joint distribution P is faithful to the graph G iff

$$P(Y_1, \dots, Y_p) \propto \prod_{C \in \mathcal{C}_G} \psi_C(Y_C)$$

where \mathcal{C}_G = set of cliques of G .



$$P(Y_1, Y_2, Y_3, Y_4) \propto \psi_1(Y_1, Y_2, Y_3) \times \psi_2(Y_2, Y_4)$$

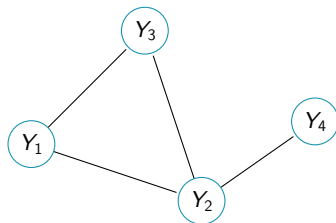
Gaussian Graphical Models (GGM)

Gaussian distribution.

$$Y_r \sim \mathcal{N}_p(\mu, \Sigma)$$

μ = vector of means, Σ = covariance matrix.

A nice property.



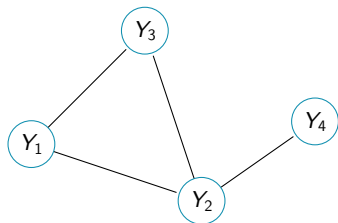
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Adjacency matrix

$$A = \begin{bmatrix} 0 & 1 & 1 & 0 \\ 1 & 0 & 1 & 0 \\ 1 & 1 & 0 & 1 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

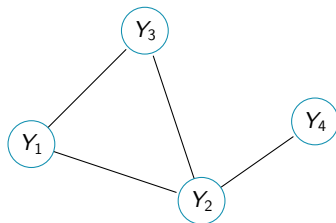
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Covariance matrix

$$\Sigma \propto \begin{bmatrix} 1 & -.25 & -.41 & .25 \\ -.25 & 1 & -.41 & .25 \\ -.41 & -.41 & 1 & -.61 \\ .25 & .25 & -.61 & 1 \end{bmatrix}$$

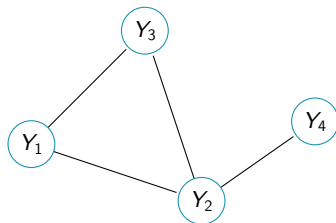
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Inverse covariance matrix

$$\Sigma^{-1} \propto \begin{bmatrix} 1 & .5 & .5 & 0 \\ .5 & 1 & .5 & 0 \\ .5 & .5 & 1 & .5 \\ 0 & 0 & .5 & 1 \end{bmatrix}$$

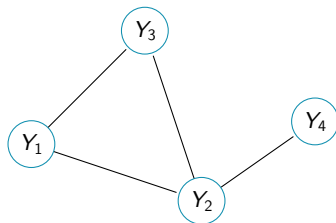
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Estimated inverse covariance matrix

$$\hat{\Sigma}^{-1} \propto \begin{bmatrix} 1 & .48 & .61 & .09 \\ .48 & 1 & .67 & .06 \\ .61 & .67 & 1 & .46 \\ .09 & .06 & .46 & 1 \end{bmatrix}$$

($n = 100$)

Sparse with Glasso

Sparsity assumption \Rightarrow find a **sparse estimate** for $\Omega = \Sigma^{-1}$

Negative log-likelihood :

$$L(Y, \Omega) \propto \frac{n}{2} \log(\det(\Omega)) - \frac{n}{2} Y^T \Omega Y$$

Graphical LASSO (glasso) :

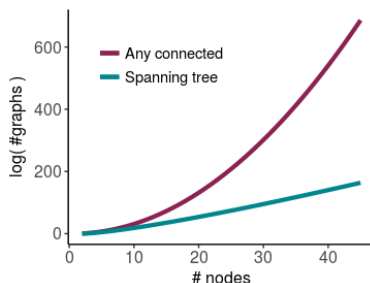
Glasso penalises the L_1 norm of the concentration matrix:

$$\hat{\Omega}_\lambda = \arg \min_{\Omega \in \mathcal{S}_d^+} \left\{ L(Y, \Omega) + \lambda \sum_{i \neq j} |w_{ij}| \right\}$$

Spanning trees

Spanning trees are another **sparse** solution :

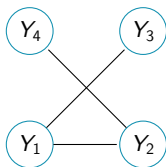
$$\left. \begin{array}{l} G \text{ is connected} \\ G \text{ has no cycle} \end{array} \right\} G \text{ has } (p - 1) \text{ edges}$$



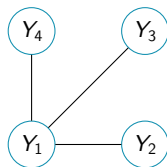
Much **smaller space** to explore:

$$\#\mathcal{G} = 2^{\frac{p(p-1)}{2}} \text{ vs. } \#\mathcal{T} = p^{(p-2)}$$

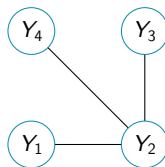
Tree averaging



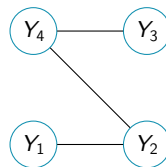
$$P\{T = T_1|Y\}$$



$$P\{T = T_2|Y\}$$



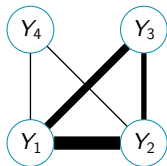
$$P\{T = T_3|Y\}$$



$$P\{T = T_4|Y\}$$

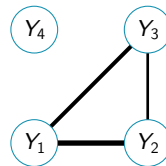
...

Compute edge
probabilities:



$$P\{(j, k) \in T|Y\}$$

Thresholding
probabilities:



$$P\{(j, k) \in T|Y\}$$

PLN model

Poisson log-Normal distribution

$$\left. \begin{array}{l} Z_i \text{ iid} \sim \mathcal{N}_d(\mu, \Sigma) \\ (Y_{ij})_j \perp\!\!\!\perp | Z_i \\ Y_{ij} | Z_{ij} \sim \mathcal{P}(e^{Z_{ij}}) \end{array} \right\} Y \sim \mathcal{PLN}(\mu, \Sigma)$$

- Gaussian latent layer
- Easily generalized to multi-variate data (contrary to Negative binomial distribution)

PLN model

Poisson log-Normal distribution

$$\left. \begin{aligned} Z_i \text{ iid} &\sim \mathcal{N}_d(\mu, \Sigma) \\ (Y_{ij})_j &\perp\!\!\!\perp Z_i \\ Y_{ij}|Z_{ij} &\sim \mathcal{P}(e^{x_i^T \Theta_j + Z_{ij}}) \end{aligned} \right\} Y \sim \mathcal{PLN}(x^T \Theta + \mu, \Sigma)$$

- Gaussian latent layer
- Easily generalized to multi-variate data (contrary to Negative binomial distribution)
- Allow adjustment for covariates

PLN model

Poisson log-Normal distribution

$$\left. \begin{aligned} Z_i \text{ iid} &\sim \mathcal{N}_d(\mu, \Sigma) \\ (Y_{ij})_j &\perp\!\!\!\perp Z_i \\ Y_{ij}|Z_{ij} &\sim \mathcal{P}(e^{\mathbf{x}_i^T \Theta_j + Z_{ij}}) \end{aligned} \right\} Y \sim \mathcal{PLN}(\mathbf{x}^T \Theta + \mu, \Sigma)$$

- Gaussian latent layer
- Easily generalized to multi-variate data (contrary to Negative binomial distribution)

Idea: Infer the **latent Gaussian network**.

Oak Mildew