Econometrics May 27, 2023

Topic 16: Graphical Network Inference

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Key points:

Disclaimer: The note is built on Prof. Jinchi Lv's lectures of the course at USC, DSO 607, High-Dimensional Statistics and Big Data Problems.

16.1 Motivation

Consider a classic question: For n observations of dimension p, how can we capture the statistical relationships between the variables of interest? Consider the example of the multivariate Gaussian distribution:

Example 16.1.1: Multivariate Gaussian Distribution

Suppose we have n observations of dimension p, $\mathbf{x} \sim \mathcal{N}(\mu, \Sigma)$. let \mathbf{S} be the empirical covariance matrix. Then the probability density

$$f(\mathbf{x}) = \frac{1}{(2\pi)^{p/2} \det(\mathbf{\Sigma})^{1/2}} \exp\left\{-\frac{1}{2}(\mathbf{x} - \boldsymbol{\mu})' \mathbf{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})\right\}$$

define the **inverse covariance matrix** or **precision matrix** as $\Omega = \Sigma^{-1}$, then we have

$$f_{\mu,\Omega} = \exp\left\{\mu'\Omega x - \left(\Omega, \frac{1}{2}xx'\right) - \frac{p}{2}\log(2\pi) + \frac{1}{2}\log\det(\Omega) - \frac{1}{2}\mu'\Omega\mu\right\}$$

where $\langle \mathbf{A}, \mathbf{B} \rangle = \operatorname{tr}(\mathbf{A}\mathbf{B})$.

In this example, we know that **every** multivariate Gaussian distribution can be represented by a pairwise **Gaussian Markov Random Field (GMRF)**, which an **undirected graph** G = (V, E)

- representing the collection of variables **x** by a vertex set $\mathcal{V} = \{1, \dots, p\}$
- encoding correlations between variables by a set of edges $\mathcal{E} = \{(i,j) \in \mathcal{V} \mid i = \neq j, \Omega_{ij} \neq 0\}$

For simplicity, we normalize $\mu = 0$. If we draw n i.i.d. samples $\mathbf{x}_1, \dots, \mathbf{x}_n \sim \mathcal{N}(\mathbf{0}, \Sigma)$, then the log-likelihood is

$$\mathcal{L}(\mathbf{\Omega}) = \frac{1}{n} \sum_{i=1}^{n} \log f(\mathbf{x}_i) = \frac{1}{2} \log \det(\mathbf{\Omega}) - \frac{1}{2n} \sum_{i=1}^{n} \mathbf{x}_1' \mathbf{\Theta} \mathbf{x}_i$$
$$= \frac{1}{2} \log \det(\mathbf{\Omega}) - \frac{1}{2} \left\langle \mathbf{\Omega}, \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i' \mathbf{x}_i' \right\rangle$$

What's the goal? We want to estimate a **sparse** graph structure given $n \ll p$ i.i.d. observations. But what does sparsity means in this context? A sparse graph is **equivalent** to a sparse precision matrix: the precision

matrix should have many 0s.

Sparse precision matrix for the Gaussian vector mentioned above $\mathbf{x} \sim \mathcal{N}(\mathbf{0}, \mathbf{\Sigma})$, we have $\forall u, v$

$$x_u \perp x_v \mid \mathbf{x}_{V \setminus \{u,v\}} \Leftrightarrow \Omega_{u,v} = 0$$

that is, sparsity of the precision matrix is equivalent to **conditional independence**¹. Consider a graph, where x_1 and x_4 are only connected through other nodes, that is x_1 and x_4 are conditional independent, then we can have the precision matrix be something like:

$$\mathbf{\Theta} = \begin{bmatrix} * & * & 0 & 0 & * & 0 & 0 & 0 \\ * & * & 0 & 0 & 0 & * & * & 0 \\ 0 & 0 & * & 0 & * & 0 & 0 & * \\ 0 & 0 & 0 & * & 0 & 0 & * & 0 \\ * & 0 & * & 0 & * & 0 & 0 & * \\ 0 & * & 0 & 0 & 0 & * & 0 & 0 \\ 0 & * & 0 & * & 0 & 0 & * & 0 \\ 0 & 0 & * & 0 & * & 0 & 0 & * \end{bmatrix}$$

where 0 captures precisely the conditional independence.



 x_1 and x_4 are connected



 x_1 and x_4 are NOT connected, conditionally

Intuitively, a sparse graph is much simpler, which is why conditional independence is desired. So how to achieve sparsity? We can again use a L-1 regularization when maximizing the log-likelihood $\mathcal{L}(\Omega)$. Denote the sample covariance matrix as $\mathbf{S} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_{i} \mathbf{x}'_{i}$, then the problem becomes the so-called **Graphical Lasso**

$$\max_{\boldsymbol{\Omega} \geq \mathbf{0}} \log \det(\boldsymbol{\Omega}) - \operatorname{tr}(\mathbf{S}\boldsymbol{\Omega}) - \rho \; \|\boldsymbol{\Omega}\|_1$$

which is equivalent to

$$\min_{\Omega \geq 0} - \log \det(\Omega) + \operatorname{tr}(\mathbf{S}\Omega) + \rho \ \|\Omega\|_1$$

16.2 Graphical Lasso

The graphical lasso method is developed by (Friedman et al., 2008). For the optimization problem

$$\min_{\Omega \ge 0} -\log \det(\Omega) + \operatorname{tr}(\mathbf{S}\Omega) + \rho \|\Omega\|_{1}$$
(16.1)

¹Meanwhile, for independence: $\Sigma_{u,v} = 0 \Leftrightarrow x_u \perp x_v$

The first-order optimality condition gives

$$\mathbf{0} \in \mathbf{\Omega}^{-1} - \mathbf{S} - \lambda \mathbf{\Gamma}$$

where Γ is a matrix of component-wise signs of Ω

$$\Gamma = \partial \|\mathbf{\Omega}\|_1 \Rightarrow \gamma_{jk} \begin{cases} = \operatorname{sign}(\omega_{jk}), & \omega_{jk} \neq 0 \\ \in [-1, 1], & \omega_{jk} = 0 \end{cases}$$

since in a graph, we always have that, following the global stationary conditions, $\omega_{jj} > 0$, which implies that

$$w_{ii} = s_{ii} + \lambda \qquad \qquad i = 1, \cdots, p \tag{16.2}$$

where we denote a working version of Ω^{-1} as **W**.

The idea is to repeatedly cycle through all columns-rows and in each step optimize only a single column-row. Consider the following partition where all matrices are partitioned into one column/row versus the rest

$$\mathbf{\Omega} = \begin{pmatrix} \mathbf{\Omega}_{11} & \boldsymbol{\omega}_{12} \\ \boldsymbol{\omega}_{12}' & \boldsymbol{\omega}_{22} \end{pmatrix} \qquad \mathbf{S} = \begin{pmatrix} \mathbf{S}_{11} & \mathbf{s}_{12} \\ \mathbf{s}_{12}' & \boldsymbol{s}_{22} \end{pmatrix} \qquad \mathbf{W} = \begin{pmatrix} \mathbf{W}_{11} & \mathbf{w}_{12} \\ \mathbf{w}_{12}' & \boldsymbol{w}_{22} \end{pmatrix} \qquad \boldsymbol{\Gamma} = \begin{pmatrix} \boldsymbol{\Gamma}_{11} & \boldsymbol{\gamma}_{12} \\ \boldsymbol{\gamma}_{12}' & \boldsymbol{\gamma}_{22} \end{pmatrix}$$

apply this partition to the optimality condition, get

$$\mathbf{\Omega}^{-1} = \mathbf{S} - \lambda \mathbf{\Gamma}$$

$$\begin{pmatrix} \mathbf{W}_{11} & \mathbf{w}_{12} \\ \mathbf{w}'_{12} & w_{22} \end{pmatrix} = \begin{pmatrix} \mathbf{S}_{11} & \mathbf{s}_{12} \\ \mathbf{s}'_{12} & s_{22} \end{pmatrix} + \lambda \begin{pmatrix} \mathbf{\Gamma}_{11} & \mathbf{\gamma}_{12} \\ \mathbf{\gamma}'_{12} & \mathbf{\gamma}_{22} \end{pmatrix}$$

where Ω_{11} is $(p-1) \times (p-1)$, ω_{12} is $(p-1) \times 1$, ω_{22} is a scalar.

Consider a **blockwise** step: suppose we fix all but the last row/column, then using properties of inverses of block-partitioned matrices, we have

$$\begin{pmatrix} \mathbf{W}_{11} & \mathbf{w}_{12} \\ \mathbf{w}_{12}' & w_{22} \end{pmatrix} = \begin{pmatrix} \left(\mathbf{\Omega}_{11} - \frac{\omega_{12}\omega_{12}'}{\omega_{22}} \right)^{-1} & -\mathbf{W}_{11} \frac{\omega_{12}}{\omega_{22}} \\ & \frac{1}{\omega_{22}} - \frac{\omega_{12}'\mathbf{W}_{11}\omega_{12}}{\omega_{22}^{2}} \end{pmatrix}$$

$$= \begin{pmatrix} \mathbf{\Omega}_{11}^{-1} + \frac{\mathbf{\Omega}_{11}^{-1}\omega_{12}\omega_{12}'\mathbf{\Omega}_{11}^{-1}}{\omega_{22} - \omega_{12}'\mathbf{\Omega}_{11}^{-1}\omega_{12}} & -\frac{\mathbf{\Omega}_{11}^{-1}\omega_{12}}{\omega_{22} - \omega_{12}'\mathbf{\Omega}_{11}^{-1}\omega_{12}} \\ & \frac{1}{\omega_{22} - \omega_{12}'\mathbf{\Omega}_{11}^{-1}\omega_{12}} \end{pmatrix}$$

then, by the partitioned optimality condition, we have:

$$\mathbf{0} = -\mathbf{w}_{12} + \mathbf{s}_{12} + \lambda \boldsymbol{\gamma}_{12} = \mathbf{W}_{11} \frac{\boldsymbol{\omega}_{12}}{\boldsymbol{\omega}_{22}} + \mathbf{s}_{12} + \lambda \boldsymbol{\gamma}_{12}$$

$$= \frac{\boldsymbol{\Omega}_{11}^{-1} \boldsymbol{\omega}_{12}}{\boldsymbol{\omega}_{22} - \boldsymbol{\omega}_{12}' \boldsymbol{\Omega}_{11}^{-1} \boldsymbol{\omega}_{12}} + \mathbf{s}_{12} + \lambda \boldsymbol{\gamma}_{12}$$

$$= \boldsymbol{\omega}_{22} \boldsymbol{\Omega}_{11}^{-1} \boldsymbol{\omega}_{12} + \mathbf{s}_{12} + \lambda \boldsymbol{\gamma}_{12}$$

here, by Eq.16.2, we know that $w_{22} = s_{22} + \lambda$, which is fixed.

References

Jerome Friedman, Trevor Hastie, and Robert Tibshirani. Sparse inverse covariance estimation with the graphical lasso. *Biostatistics*, 9(3):432–441, 2008.