Joint Estimation and Inference for Multiple Multi-layered Gaussian Graphical Models

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Abstract: The rapid development of high-throughput technologies has enabled generation of data from biological processes that span multiple layers, like genomic, proteomic or metabolomic data; and pertain to multiple sources, like disease subtypes or experimental conditions. In this work we propose a general statistical framework based on graphical models for horizontal (i.e. across conditions or subtypes) and vertical (i.e. across different layers containing data on molecular compartments) integration of information in such datasets. We start with decomposing the multi-layer problem into a series of two-layer problems. For each two-layer problem, we model the outcomes at a node in the lower layer as dependent on those of other nodes in that layer, as well as all nodes in the upper layer. Following the biconvexity of our objective function, this estimation problem decomposes into two parts, where we use neighborhood selection and subsequent refitting of the precision matrix to quantify the dependency of two nodes in a single layer, and use group-penalized least square estimation to quantify the directional dependency of two nodes in different layers. Finally, to test for differences in these directional dependencies across multiple sources, we devise a hypothesis testing procedure that utilizes already computed neighborhood selection coefficients for nodes in the upper layer. We establish theoretical results for the validity of this testing procedure and the consistency of our estimates, and also evaluate their performance through simulations and a real data application.

Keywords: Data integration; Gaussian Graphical Models; Neighborhood selection; Group lasso

1 Notations

We shall denote scalars by small letters, vectors by bold small letters and matrices by bold capital letters. For any matrix $\mathbf{A}$, $(\mathbf{A})_{ij}$ denote its element in the $(i,j)^{\text{th}}$ position. For $a, b \in \mathbb{N}$, we denote the set of all $a \times b$ real matrices by $\mathbb{M}(a,b)$.

2 Model

Consider the two -layered setup:

$$\mathbb{X}^k = (X_1^k, \ldots, X_p^k)^T \sim \mathcal{N}(0, \Sigma_x^k) \tag{2.1}$$

$$\mathbb{Y}^k = \mathbb{X}^k \mathbf{B}^k + \mathbf{E}^k; \quad \mathbf{E}^k = (E_1^k, \ldots, E_p^k)^T \sim \mathcal{N}(0, \Sigma_y^k) \tag{2.2}$$

$$\mathbf{B}^k \in \mathbb{M}(p,q); \quad \Omega_x^k = (\Sigma_x^k)^{-1}; \quad \Omega_y^k = (\Sigma_y^k)^{-1} \tag{2.3}$$

Want to estimate $\{(\Omega_x^k, \Omega_y^k, \mathbf{B}^k); 1 \le k \le K\}$ from data $\mathcal{Z}^k = \{(\mathbf{Y}^k, \mathbf{X}^k); \mathbf{Y}^k \in \mathbb{M}(n,q), \mathbf{X}^k \in \mathbb{M}(n,p), k = 1, \ldots, K\}$. in presence of known grouping structures $\mathcal{G}_x, \mathcal{G}_y, \mathcal{H}$ respectively.

Estimation of $\{\Omega_x^k\}$ done using JSEM. For the other part, we use the following two-step procedure:

1. Run neighborhood selection on y-network incorporating effects of x-data and an additional block-wise group penalty:

$$\min_{\mathcal{B}, \Theta} \left\{ \sum_{j=1}^q \frac{1}{n_k} \left[\sum_{k=1}^K \|\mathbf{y}_j^k - (\mathbf{Y}_{-j}^k - \mathbf{X}^k \mathbf{B}_{-j}^k)\boldsymbol{\theta}_j^k - \mathbf{X}^k \mathbf{b}_j^k\|^2 + \sum_{j \ne i} \sum_{g \in \mathcal{G}_y^{ij}} \lambda_{ij}^g \|\boldsymbol{\theta}_{ij}^{[g]}\| \right] + \sum_{b \in \mathcal{G}_x \times \mathcal{G}_y \times \mathcal{H}} \eta^b \|\mathbf{B}^{[b]}\| \right\} \tag{2.4}$$

$$= \min \left\{ f(\mathcal{Y}, \mathcal{X}, \mathcal{B}, \Theta) + P(\Theta) + Q(\mathcal{B}) \right\} \tag{2.5}$$

where $\Theta = \{\Theta_i\}, \mathcal{B} = \{\mathbf{B}^k\}, \mathcal{Y} = \{\mathbf{Y}^k\}, \mathcal{X} = \{\mathbf{X}^k\}, \mathcal{E} = \{\mathbf{E}^k\}$.

This estimates $\mathcal{B}$ **(possibly refit and/or within-group threshold)** .

2. Step I part 2 and step II of JSEM (see 15-656 pg 6) follows to estimate $\{\Omega_y^k\}$.

The objective function is bi-convex, so we are going to do the following in step 1-

- Start with initial estimates of $\mathcal{B}$ and Θ, say $\mathcal{B}^{(0)}, \Theta^{(0)}$.

- Iterate:

$$\Theta^{(t+1)} = \arg\min \left\{ f(\mathcal{Y}, \mathcal{X}, \mathcal{B}^{(t)}, \Theta^{(t)}) + P(\Theta^{(t)}) \right\} \tag{2.6}$$

$$\mathcal{B}^{(t+1)} = \arg\min \left\{ f(\mathcal{Y}, \mathcal{X}, \mathcal{B}^{(t)}, \Theta^{(t+1)}) + Q(\mathcal{B}^{(t)}) \right\} \tag{2.7}$$

- Continue till convergence.

3 Two-sample testing

Suppose there are two disease subtypes: $k = 1, 2$, and we are interested in testing whether the downstream effect of a predictor is X-data is same across both subtypes, i.e. if $\mathbf{b}_i^1 = \mathbf{b}_i^2$ for some $i \in \{1, \ldots, p\}$. For this we consider the modified optimization problem:

$$\min_{\mathcal{B}, \Theta} \frac{1}{n} \left\{ \sum_{j=1}^{q} \sum_{k=1}^{2} \|\mathbf{Y}_j^k - \mathbf{Y}_{-j}^k \boldsymbol{\theta}_j^k - \mathbf{X}^k \mathbf{b}_j^k\|^2 + \sum_{j \neq j'} \lambda_{jj'} \|\boldsymbol{\theta}_{jj'}^*\| + \sum_{i=1}^{p} \eta_i \|\mathbf{B}_{i*}^*\| \right\} \tag{3.1}$$

$$= \min \left\{ f(\mathcal{Y}, \mathcal{X}, \mathcal{B}, \Theta) + P(\Theta) + Q(\mathcal{B}) \right\} \tag{3.2}$$

with $n_1 = n_2 = n$ for simplicity; and $\mathbf{B}^k = (\mathbf{b}_1^k, \ldots, \mathbf{b}_q^k), (\mathbf{B}_{i*}^*) \in \mathbb{R}^{q \times K}$

4 Conditions

Conditions A1, A2, A3 from JSEM paper.

5 Results

Define

$$\hat{\Theta}_j = \arg\min_{\Theta_j} \left\{ \frac{1}{n} \sum_{k=1}^{K} \|\mathbf{y}_j^k - (\mathbf{Y}_{-j}^k - \mathbf{X}^k \hat{\mathbf{B}}_{-j}^k) \boldsymbol{\theta}_j^k - \mathbf{X}^k \hat{\mathbf{b}}_j^k\|^2 + \sum_{j \neq j'} \sum_{g \in \mathcal{G}_y^{jj'}} \lambda_{jj'}^g \|\boldsymbol{\theta}_{jj'}^{[g]}\| \right\} \tag{5.1}$$

Theorem 5.1. *Assume fixed $\mathcal{X}, \mathcal{E}$ and deterministic $\hat{\mathcal{B}} = \{\hat{\mathbf{B}}^k\}$. Also for $k = 1, \ldots, K$,*

(T1) $\|\hat{\mathbf{B}}^k - \mathbf{B}_0^k\|_F \leq v_\beta$;

(T2) $\|\mathbf{X}^k(\hat{\mathbf{B}}^k - \hat{\mathbf{B}}_0^k)\|_F \leq c(v_\beta)$ *for some non-negative function $c(.)$;*

Group uniform IC.

Then

(I) Estimation consistency

(II) Direction consistency

Proof of Theorem 5.1. Part I. Follows proof of thm 1 in 15-656. The proof has 3 parts: consistency of neighborhood regression, selection of edge sets, and finally the refitting step.

For any $g \in \mathcal{G}^{jj'}, k \in g$, and $j \neq j'$, let

$$\hat{\boldsymbol{\epsilon}}_j^k = \mathbf{y}_j^k - (\mathbf{Y}_{-j}^k - \mathbf{X}^k\hat{\mathbf{B}}_{-j}^k)\boldsymbol{\theta}_{0,j}^k - \mathbf{X}^k\hat{\mathbf{b}}_j^k; \quad \hat{\zeta}_{jj'}^k = \frac{(\hat{\boldsymbol{\epsilon}}_j^k)^T \mathbf{Y}_{j'}^k}{n}; \quad \hat{\boldsymbol{\zeta}}_{jj'}^{[g]} = (\hat{\zeta}_{jj'}^k)_{k \in g}$$

Consider the random event $\mathcal{A} = \bigcap_{j,j' \neq j,g} \mathcal{A}_{jj'}^g$ with $\mathcal{A}_{jj'}^g = \{2\|\hat{\boldsymbol{\zeta}}_{jj'}^{[g]}\| \leq \lambda_{jj'}^g\}$.

Proposition 5.2. *Given that λ_{ij}^g are chosen as*

$$\lambda_{ij}^g \geq \max_{k \in g} \frac{2}{\sqrt{n\omega_{ii}^k}} \left(\sqrt{|g|} + \frac{\pi}{\sqrt{2}}\sqrt{q \log G_0} + \sqrt{c(v_\beta)} \right)$$

we shall have $\mathbb{P}(\mathcal{A}) \geq 1 - 2pG_0^{1-q}$ for some $q > 1$.

Proof of Proposition 5.2. We follow the proof of Lemma E.2 in 15-656, with $\mathbf{Y}_j^k, \hat{\boldsymbol{\epsilon}}_i^k, \hat{\zeta}_{ij}^k, \hat{\boldsymbol{\zeta}}_{ij}^{[g]}$ in place of $\mathbf{X}_j^k, \boldsymbol{\epsilon}_i^k, \zeta_{ij}^k, \boldsymbol{\zeta}_{ij}^{[g]}$ respectively. Proceeding in a similar fashion we get

$$\|\hat{\boldsymbol{\zeta}}_{ij}^{[g]}\|^2 = \frac{1}{n}(\|\mathbf{Z}^{[g]}\|^2 + 2\sum_{k \in g} Z^k(\mathbf{Q}_j^k)^T\boldsymbol{\delta}_i^k + \|(\mathbf{Q}_j^k)^T\boldsymbol{\delta}_i^k\|^2)$$

where $\mathbf{Z}^{[g]} = (Z^k)_{k \in g}; Z^k = (\mathbf{Q}_j^k)^T\boldsymbol{\epsilon}_i^k$ with $\boldsymbol{\epsilon}_i^k := \mathbf{Y}_i^k - \mathbf{Y}_{-i}^k\boldsymbol{\theta}_{0,i}^k - \mathbf{X}^k\mathbf{B}_{0,i}^k$, $\mathbf{Q}_j^k$ is the first eigenvector of $\mathbf{Y}_j^k(\mathbf{Y}_j^k)^T/n$, and $\boldsymbol{\delta}_i^k := \mathbf{X}^k(\mathbf{B}_{0,i}^k - \hat{\mathbf{B}}_i^k)$. Applying Cauchy-schwarz inequality to right side and by assumption (T2),

$$\|\hat{\boldsymbol{\zeta}}_{ij}^{[g]}\| \leq \frac{1}{\sqrt{n}}(\|\mathbf{Z}^{[g]} + \sqrt{c(v_\beta)})$$

thus

$$\mathbb{P}(\{\mathcal{A}_{ij}^g\}^c) = \mathbb{P}\left(\|\hat{\boldsymbol{\zeta}}_{ij}^{[g]}\| > \frac{\lambda_{ij}^g}{2} \right) \leq \mathbb{P}\left(\|\mathbf{Z}^{[g]}\| > \frac{\sqrt{n}\lambda_{ij}^g}{2} - \sqrt{c(v_\beta)} \right)$$

We now proceed through the proof of Lemma E.2 in 15-656 to end up with the choice of λ_{ij}^g. $\square$

All subsequent derivations in the theorem go through with the new choice of λ_{ij}^g.

Part II. Proof of Thm 2 in 15-656 follows. We only need a new bound for $Var(\mathbf{Y}_i^k|\mathbf{Y}_{-i}^k, \mathbf{X}^k, \hat{\mathbf{B}}_i^k)$. For this we have

$$Var(\mathbf{Y}_i^k|\mathbf{Y}_{-i}^k, \mathbf{X}^k, \hat{\mathbf{B}}_i^k) = \mathbb{E}(\hat{\epsilon}_i^k)^2 = \mathbb{E}(\boldsymbol{\epsilon}_i^k + \boldsymbol{\delta}_i^k)^2 \leq \left(\frac{1}{d_0} + \frac{c(v_\beta)}{n}\right)^2$$

applying cauchy-schwarz inequality followed by assumption (A2). Now Replace $1/\sqrt{nd_0}$ in choice of λ, α_n in Thm 2 statement with $1/\sqrt{n}(\sqrt{1/d_0} + \sqrt{c(v_\beta)/n})$.

$\square$

Proposition 5.3. *Given fixed $\hat{\mathcal{B}}$, prediction errors follow bound in T2 with high enough probability.*

Now concentrate on the k-population estimation problem. We want to obtain

$$\hat{\boldsymbol{\beta}} = \underset{\boldsymbol{\beta} \in \mathbb{R}^{pqK}}{\arg\min}\{-2\boldsymbol{\beta}\hat{\boldsymbol{\gamma}} + \boldsymbol{\beta}^T \boldsymbol{\Gamma} \boldsymbol{\beta} + \|\boldsymbol{\beta}\|_{2,g}\}$$

with

$$\boldsymbol{\beta} = \begin{bmatrix} \text{vec}(\mathbf{B}^1) \\ \vdots \\ \text{vec}(\mathbf{B}^K) \end{bmatrix}; \quad \boldsymbol{\Gamma} = \begin{bmatrix} I_q \otimes (\mathbf{X}^1)TX^1/n & & \\ & \ddots & \\ & & I_q \otimes (\mathbf{X}^K)^T X^K/n \end{bmatrix}$$

Theorem 5.4.