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

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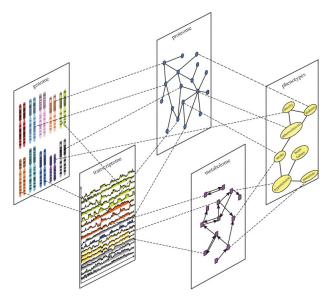
University of Florida Informatics Institute

IISA-2017 Conference, Hyderabad, India December 28, 2017

December 26, 2017

- Biological processes in the body have a natural hierarchical structure,
   e.g. Gene > Protein > Metabolite;
- There are within layer and between-layer connections in this structure;
- These connections can be different inside different organs, experimental conditions, or for different subtypes of the same disease;
- We design a framework for joint estimation and hypothesis testing for all the connections in these complex biological structure.

## Schematic of data integration

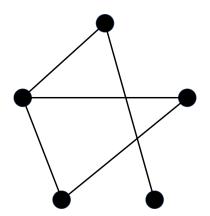


(Source: Gligorijević and Pržulj (2015))

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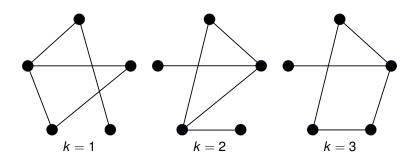
- Formulation of multiple multi-level graphical models
- Model formulation, computation and theory
- 3 Hypothesis testing
- Simulation studies

$$\mathbb{X} = (X_1, \ldots, X_p)^T \sim \mathcal{N}_p(0, \Sigma_x); \quad \Omega_x = \Sigma_x^{-1}$$



Sparse estimation of  $\Omega_x$ : Meinshausen and Bühlmann (2006) Multiple testing and error control: Drton and Perlman (2007).

$$\mathbb{X}^k = (X_1^k, \dots, X_p^k)^T \sim \mathcal{N}_p(0, \Sigma_x^k); \quad \Omega_x^k = (\Sigma_x^k)^{-1}$$
  
 $k = 1, 2, \dots, K$ 

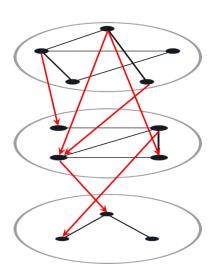


- Joint estimation of  $\{\Omega_x^k\}$ :
- Difference and similarity testing with FDR control: Liu (2017+)

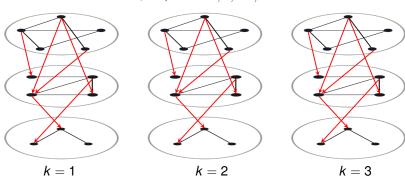
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$$\begin{split} \mathbb{E} &= (E_1, \dots, E_q)^T \sim \mathcal{N}_p(0, \Sigma_y); \\ \Omega_y &= (\Sigma_y)^{-1} \\ \mathbb{Y} &= \mathbb{X} \boldsymbol{B} + \mathbb{E} \end{split}$$

- Ω<sub>x</sub>, Ω<sub>y</sub> give undirected within-layer edges, while **B** gives directed between-layer edges.
- Sparse estimation of  $(\Omega_V, \Omega_X, \mathbf{B})$ : Lin et al. (2016).



$$\begin{split} \mathbb{E}^k &= (E_1^k, \dots, E_q^k)^T \sim \mathcal{N}_p(0, \Sigma_y^k); \quad \Omega_y^k = (\Sigma_y^k)^{-1} \\ \mathbb{Y}^k &= \mathbb{X}^k \textbf{B}^k + \mathbb{E}^k; \quad k = 1, 2, \dots, K \end{split}$$



- We estimate  $\{\Omega_x^k, \Omega_y^k, \mathbf{B}^k\}$  jointly for all k from a single model;
- For K = 2 and  $i \in \{1, 2, ..., p\}$ , we also provide a global test for  $\mathbf{b}_i^1 = \mathbf{b}_i^2$ , and do multiple testing for  $b_i^1 = b_i^2$ , j = 1, 2, ..., q.

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#### **Notations**

For single GGM: Estimate neighboring edges for each node, then refit.

$$\begin{split} \widehat{\boldsymbol{\zeta}}_i &= \operatorname*{argmin}_{\boldsymbol{\zeta}_i} \left\{ \frac{1}{n} \| \mathbf{X}_i - \mathbf{X}_{-i} \boldsymbol{\zeta}_i \|^2 + \nu_n \sum_{i' \neq i} |\zeta_{ii'}| \right\}; \\ \widehat{\boldsymbol{\Omega}}_{\boldsymbol{\chi}} &= \operatorname*{argmin}_{\boldsymbol{\Omega}_{\boldsymbol{\chi}} \in \cup, \mathrm{support}(\boldsymbol{\zeta}_i)} \{ \mathrm{Tr}(\mathbf{S}_{\boldsymbol{\chi}} \boldsymbol{\Omega}_{\boldsymbol{\chi}}) + \log \det(\boldsymbol{\Omega}_{\boldsymbol{\chi}}) \} \end{split}$$

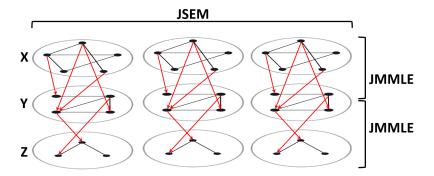
Can do this because  $\zeta_{ii'} = -\omega_{ii'}/\omega_{ii}$ .

 For multiple GGM: Incorporate penalty across different k (JSEM: Ma and Michailidis (2016)).

$$\begin{split} \widehat{\boldsymbol{\zeta}}_i &= \operatorname*{argmin}_{\boldsymbol{\zeta}_i} \left\{ \sum_{k=1}^K \frac{1}{n_k} \| \mathbf{X}_i^k - \mathbf{X}_{-i}^k \boldsymbol{\zeta}_i^k \|^2 + \nu_n \sum_{i' \neq i, g \in \mathcal{G}_{ii'}} \| \boldsymbol{\zeta}_{ii'}^{[g]} \| \right\}; \\ \widehat{\Omega}_X^k &= \operatorname*{argmin}_{\Omega_X^k \in \cup, \text{support}(\boldsymbol{\zeta}_i^k)} \left\{ \text{Tr}(\mathbf{S}_X^k \Omega_X^k) + \log \det(\Omega_X^k) \right\}; \end{split}$$

### **Joint Multiple Multi-Level Estimation (JMMLE)**

We decompose the multi-layer problem into a series of two layer problems. For within-layer connections in the topmost layer, we use JSEM. For other connections we use JMMLE.



We combine sparse neighborhood selection in the Y-network with sparse estimation of  $\{\mathbf{B}^k\}$ .

$$\begin{split} \{\widehat{\mathcal{B}}, \widehat{\Theta}\} &= \underset{\mathcal{B}, \Theta}{\operatorname{argmin}} \left\{ \sum_{j=1}^{q} \frac{1}{n_{k}} \sum_{k=1}^{K} \left\| \mathbf{Y}_{j}^{k} - (\mathbf{Y}_{-j}^{k} - \mathbf{X}^{k} \mathbf{B}_{-j}^{k}) \boldsymbol{\theta}_{j}^{k} - \mathbf{X}^{k} \mathbf{B}_{j}^{k} \right\|^{2} \right. \\ &+ \lambda_{n} \sum_{h \in \mathcal{H}} \| \mathbf{B}^{[h]} \| + \gamma_{n} \sum_{j' \neq j, g \in \mathcal{G}_{jj'}} \| \boldsymbol{\theta}_{jj'}^{[g]} \| \right\} \\ &\widehat{\Omega}_{y}^{k} = \underset{\Omega_{y}^{k} \in \cup, \text{support}(\boldsymbol{\theta}_{j}^{k})}{\operatorname{argmin}} \left\{ \operatorname{Tr}(\mathbf{S}_{y}^{k} \Omega_{y}^{k}) + \log \det(\Omega_{y}^{k}) \right\} \quad k = 1, 2, \dots, K \end{split}$$

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$$\{\widehat{\mathcal{B}}, \widehat{\Theta}\} = \operatorname*{argmin}_{\mathcal{B}, \Theta} \{f(\mathcal{Y}, \mathcal{X}, \mathcal{B}, \Theta) + P(\mathcal{B}) + Q(\Theta)\}$$

The objective function is biconvex, so we solve the above by the following alternating iterative algorithm:

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

$$\begin{split} \mathcal{B}^{(t+1)} &= \underset{\mathcal{B}}{\operatorname{argmin}} \left\{ f(\mathcal{Y}, \mathcal{X}, \mathcal{B}, \Theta^{(t)}) + Q(\mathcal{B}) \right\} \\ \Theta^{(t+1)} &= \underset{\Theta}{\operatorname{argmin}} \left\{ f(\mathcal{Y}, \mathcal{X}, \mathcal{B}^{(t+1)}, \Theta) + P(\Theta) \right\} \end{split}$$

Continue till convergence.

For  $\lambda_n \geq 4\sqrt{|h_{\max}|}\mathbb{R}_0\sqrt{\frac{\log(pq)}{n}}$ , the following hold with probability approaching 1 as  $n \to \infty$ ,

$$\|\widehat{eta} - eta_0\|_1 \leq rac{48\sqrt{|h_{\mathsf{max}}|}s_eta\lambda_n}{\psi^*}$$
  $\|\widehat{eta} - eta_0\| \leq rac{12\sqrt{s}_eta\lambda_n}{\psi^*}$   $\sum_{h \in \mathcal{H}} \|eta^{[h]} - eta_0^{[h]}\| \leq rac{48s_eta\lambda_n}{\psi^*}$ 

with  $\psi^*$ ,  $\mathbb{R}_0$  being constants, and  $\boldsymbol{\beta} = (\text{vec}(\mathbf{B}^1)^T, \dots, \text{vec}(\mathbf{B}^K)^T)^T$ ,  $|h_{\text{max}}|$  the maximum group size in  $\boldsymbol{\beta}_0$  and  $\boldsymbol{s}_{\boldsymbol{\beta}}$  the sparsity of  $\boldsymbol{\beta}_0$ .

For  $\gamma_n = 4\sqrt{|g_{\max}|}\mathbb{Q}_0\sqrt{\frac{\log(pq)}{n}}$ , the following hold with probability approaching 1 as  $n \to \infty$ ,

$$\begin{split} \|\widehat{\Theta}_{j} - \Theta_{0,j}\|_{F} &\leq \frac{12\sqrt{s_{j}}\gamma_{n}}{\psi} \\ \sum_{i \neq j', g \in \mathcal{G}_{y}^{jj'}} \|\widehat{\boldsymbol{\theta}}_{jj'}^{[g]} - \boldsymbol{\theta}_{0,jj'}^{[g]}\| &\leq \frac{48s_{j}\gamma_{n}}{\psi} \\ \left| \mathsf{support}(\widehat{\Theta}_{j}) \right| &\leq \frac{128s_{j}}{\psi} \\ \frac{1}{K} \sum_{k=1}^{K} \|\widehat{\Omega}_{y}^{k} - \Omega_{y}^{k}\|_{F} &\leq O\left(\frac{\sqrt{S}\gamma_{n}}{\sqrt{K}}\right) \end{split}$$

with  $\psi$ ,  $\mathbb{Q}_0$  being constants,  $|g_{\text{max}}|$  the maximum group size in  $\Theta_0$ ,  $s_j$  the sparsity of  $\Theta_i$  and  $S = \sum_i s_i$ .

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- Consider the case K = 2, and suppose we are interested in testing if the effect of variable i in the X-data is different across the two populations.
- For this we use the  $i^{th}$  rows of the estimates  $\hat{\mathbf{B}}^1$  and  $\hat{\mathbf{B}}^2$ .
- We debias these row vectors using the neighborhood coefficients in the X-network computed previously using JSEM: In this setup, define the desparsified estimate of b<sup>k</sup><sub>i</sub> as

$$\widehat{\mathbf{c}}_{i}^{k} = \widehat{\mathbf{b}}_{i}^{k} + \frac{1}{nt_{i}^{k}} \left( \mathbf{X}_{i}^{k} - \mathbf{X}_{-i}^{k} \widehat{\boldsymbol{\zeta}}_{i}^{k} \right)^{T} (\mathbf{Y}^{k} - \mathbf{X}^{k} \widehat{\mathbf{B}}^{k})$$

for k = 1, 2, where  $t_i^k := (\mathbf{X}_i^k - \mathbf{X}_{-i}^k \widehat{\zeta}_i^k)^T \mathbf{X}_{-i}^k / n$ .

Assume we have 'good enough' estimators:

$$\begin{split} \|\widehat{\boldsymbol{\zeta}}^k - \boldsymbol{\zeta}_0^k\|_1 &= O\left(\sqrt{\frac{\log p}{n}}\right); \quad \|\widehat{\mathbf{B}}^k - \mathbf{B}_0^k\|_1 = O\left(\sqrt{\frac{\log (pq)}{n}}\right) \\ \left\|(\widehat{\Omega}_y^k)^{1/2} - (\Omega_y^k)^{1/2}\right\|_{\infty} &= O\left(\sqrt{\frac{\log q}{n}}\right) \end{split}$$

Also define

$$\widehat{\boldsymbol{s}}_{i}^{k} := \sqrt{\|\boldsymbol{\mathsf{X}}_{i}^{k} - \boldsymbol{\mathsf{X}}_{-i}^{k} \widehat{\boldsymbol{\zeta}}_{i}^{k}\|^{2}/n}; \quad \boldsymbol{m}_{i}^{k} := \sqrt{n} t_{i}^{k} / \widehat{\boldsymbol{s}}_{i}^{k}$$

Then for sample size satisfying  $n \gtrsim \log(pq)$ ,  $\log p = o(n^{1/2})$ ,  $\log q = o(n^{1/2})$  we have

$$\begin{bmatrix} \widehat{\Omega}_y^1 & \\ & \widehat{\Omega}_y^2 \end{bmatrix}^{1/2} \begin{bmatrix} m_i^1 (\widehat{\mathbf{c}}_i^1 - \mathbf{b}_i^1) & \\ & m_i^2 (\widehat{\mathbf{c}}_i^2 - \mathbf{b}_i^2) \end{bmatrix} \sim \mathcal{N}_{2q}(\mathbf{0}, \mathbf{I}) + o_P(1)$$

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- **①** Obtain the debiased estimators  $\hat{\mathbf{c}}_i^1, \hat{\mathbf{c}}_i^2$ ;
- 2 Calculate the test statistic

$$D_i = \left\| m_i^1 (\widehat{\Omega}_y^1)^{1/2} \widehat{\mathbf{c}}_i^1 - m_i^2 (\widehat{\Omega}_y^2)^{1/2} \widehat{\mathbf{c}}_i^2 \right\|^2$$

**3** Reject  $H_0$  if  $D_i \geq \chi^2_{2q,1-\alpha}$ .

• Calculate the pairwise test statistics  $d_{ij}$  for j = 1, ..., q:

$$d_{ij} = rac{ au_{ij}^1 \hat{\mathbf{c}}_{ij}^1 - au_{ij}^2 \hat{\mathbf{c}}_{ij}^2}{1/m_i^1 + 1/m_i^2}$$

where  $\tau_{ii}^k$  is the  $(i,j)^{\text{th}}$  element of  $(\widehat{\Omega}_{v}^k)^{1/2}, k=1,2$ .

Obtain the threshold

$$\hat{ au} = \inf \left\{ au \in \mathbb{R} : 1 - \Phi( au) \leq rac{lpha}{2q} \max \left( \sum_{j \in \mathcal{I}_q} \mathbb{I}(| extbf{ extit{d}}_{ij}| \geq au), 1 
ight) 
ight\}$$

**③** For  $j \in \mathcal{I}_q$ , reject  $H_0^j$  if  $|d_{ij}| \geq \hat{\tau}$ .

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# **THANK YOU!**

**Questions?**