### Envelope method with ignorable missing data

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  - Regression with multiple responses
  - Motivation
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  - The envelope model
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  - Motivation and preliminary
  - EM envelope algorithm
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### Regression with multiple responses

Linear regression model can be written as:

$$\mathbf{Y}_{1\times r} = \mathbf{X}_{1\times p} \boldsymbol{\beta}_{p\times r} + \boldsymbol{\varepsilon}_{1\times r},$$

where the error vector  $\varepsilon_{1\times r}\in\mathbb{R}^r$  is normally distributed with mean  $\mathbf{0}$  and unknown parameter  $\Sigma$ . When  $\Sigma>0$ , the model has a total number of pr+r(r+1)/2 unknown parameters.

• Suppose we observe the data set  $((\mathbf{x}_1, \mathbf{y}_1), \dots, (\mathbf{x}_n, \mathbf{y}_n))$ , we can write them as a matrix form:

$$\mathbf{Y}_{n\times r} = \mathbf{X}_{n\times p}\beta_{p\times r} + \varepsilon_{n\times r}.$$

• The log-likelihood of the linear model:

$$I(oldsymbol{eta}, oldsymbol{\Sigma} | oldsymbol{Y}) = -rac{n}{2} \det(oldsymbol{\Sigma}) - rac{1}{2} (oldsymbol{Y} - oldsymbol{X} oldsymbol{eta}) oldsymbol{\Sigma}^{-1} (oldsymbol{Y} - oldsymbol{X} oldsymbol{eta})^T + C$$

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• Setting the partial derivative of  $\beta$  to  $\mathbf{0}$ , we have

$$\frac{\partial l}{\partial \boldsymbol{\beta}} = \mathbf{X}^T \mathbf{Y} \mathbf{\Sigma}^{-1} - \mathbf{X}^T \mathbf{X} \boldsymbol{\beta} \boldsymbol{\Sigma}^{-1} = \mathbf{0}.$$

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ullet Since  $\Sigma$  is positive definite, we can cancel it on both sides. Hence the MLE of eta is

$$\hat{\boldsymbol{\beta}}_{MLE} = (\mathbf{X}^T \mathbf{X})^{\dagger} \mathbf{X}^T \mathbf{Y},$$

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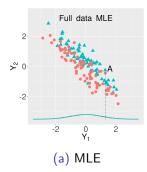
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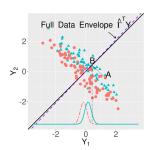
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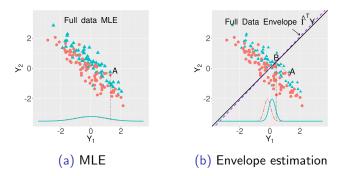
#### Remark

The estimator of  $oldsymbol{eta}$  does not depend on  $oldsymbol{\Sigma}$  at all.

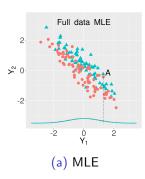


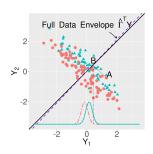


(b) Envelope estimation



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- Suppose we want to test whether  $\beta_1 = 0$ . Using the standard method, we get p-value= 0.37; Using envelope estimation, the p-value is less than 0.001.
- Actually, the true mean group difference  $\beta_1=0.32$ .  $\hat{\beta}_{1,MLE}=0.12$  with bootstrap standard error being 0.12, while  $\hat{\beta}_{1,env}=0.32$  with bootstrap standard error being 0.03.

- From the graph, we see that some characteristics of the response vector could be unaffected by changes in the predictors.
- For the ease of notation later, we slightly change the model as

$$\mathbf{Y}_{r\times 1} = \boldsymbol{\beta}_{r\times p} \mathbf{X}_{p\times 1} + \boldsymbol{\varepsilon}_{r\times 1}.$$

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• Consider a subspace  $\mathcal{E} \subseteq \mathbb{R}^r$  such that  $(\mathbf{P}_{\mathcal{E}} = \text{projection onto } \mathcal{E}, \mathbf{Q}_{\mathcal{E}} = \mathbf{I} - \mathbf{P}_{\mathcal{E}})$ :

$$\begin{split} \textbf{Q}_{\mathcal{E}}\textbf{Y}|(\textbf{X}=\textbf{x}_1) \sim \textbf{Q}_{\mathcal{E}}\textbf{Y}|(\textbf{X}=\textbf{x}_2), \ \forall \textbf{x}_1,\textbf{x}_2 &\Longleftrightarrow \operatorname{span}(\boldsymbol{\beta}) \subset \mathcal{E} \\ \textbf{P}_{\mathcal{E}}\textbf{Y} \perp \!\!\! \perp \textbf{Q}_{\mathcal{E}}\textbf{Y}|\textbf{X} &\Longleftrightarrow \boldsymbol{\Sigma} = \textbf{P}_{\mathcal{E}}\boldsymbol{\Sigma}\textbf{P}_{\mathcal{E}} + \textbf{Q}_{\mathcal{E}}\boldsymbol{\Sigma}\textbf{Q}_{\mathcal{E}} \end{split}$$

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• This implies the impact of  $\mathbf{X}$  on  $\mathbf{Y}$  is concentrated only in  $\mathbf{P}_{\mathcal{E}}\mathbf{Y}$ . We refer to  $\mathbf{P}_{\mathcal{E}}\mathbf{Y}$  and  $\mathbf{Q}_{\mathcal{E}}\mathbf{Y}$  as material and immaterial part of  $\mathbf{Y}$ .

• Suppose  $\Gamma_{r \times u}$  is a semi-orthogonal matrix  $(\Gamma^T \Gamma = \mathbf{I}_{u \times u})$  such that  $\operatorname{span}(\Gamma) = \mathcal{E}$ , then the previous two conditions are equivalent to

$$\operatorname{Span}(\boldsymbol{\beta})\subseteq\operatorname{Span}(\boldsymbol{\Gamma})$$

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#### Example

Assume  $\mathbf{Y} = (Y_1, Y_2)^T$ , where  $Y_1 = \beta \mathbf{X} + \varepsilon_1$ , and  $Y_2 = -\beta \mathbf{X} + \varepsilon_2$ . Suppose  $\varepsilon_1$  and  $\varepsilon_2$  follow two normal distributions and they are independent of each other.

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Then, the predictors  ${\bf X}$  do not affect the summation of response  $Y_1+Y_2=\varepsilon_1+\varepsilon_2$ . Also,  $Y_1+Y_2$  is independent of  $Y_1-Y_2$ , so that  $Y_1+Y_2$  can be completely discarded. That is, the regression of  ${\bf Y}$  on  ${\bf X}$  can be replaced with the regression of  $Y_1-Y_2$  on  ${\bf X}$ .

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#### Formal definition

• The semi-orthogonal matrix  $\Gamma$  is not unique. For example, identity matrix  $\mathbf{I}_r$  satisfies the two conditions trivially, i.e.,

$$\mathrm{Span}(oldsymbol{eta}) \subseteq \mathrm{Span}(I_r)$$
  
 $\Sigma = P_{I_r} \Sigma P_{I_r} + Q_{I_r} \Sigma Q_{I_r}$ 

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### Definition 1. (Cook et al., 2010)

The intersection of all subspaces  $\mathcal{E}$  with properties  $\mathrm{Span}(\beta) \subseteq \mathrm{Span}(\Gamma)$  and  $\Sigma = \mathbf{P}_{\mathcal{E}} \Sigma \mathbf{P}_{\mathcal{E}} + \mathbf{Q}_{\mathcal{E}} \Sigma \mathbf{Q}_{\mathcal{E}}$  is defined as the  $\Sigma$ -envelope of  $\beta$ , denoted by  $\mathcal{E}_{\Sigma}(\beta)$ .  $u = \dim(\mathcal{E}_{\Sigma}(\beta))$  is called the envelope dimension.

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#### Proposition 1. (Cook et al., 2010)

Assume that  $\Sigma_{r \times r}$  is symmetric and has  $q \leq r$  distinct eigenvalues. Let  $\mathbf{P}_i$ ,  $i = 1, \ldots, q$ , indicate the projection onto the corresponding eigenspaces. Then,

$$\mathcal{E}_{oldsymbol{\Sigma}}(\mathcal{B})=\oplus_{q=1}^q\mathsf{P}_i\mathcal{B}$$

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- Let the columns of the semi-orthogonal matrices  $\Gamma \in \mathbb{R}^{u \times r}$  be the base of  $\mathcal{E}_{\Sigma}(\mathcal{B})$ , and  $\Gamma_0 \in \mathbb{R}^{r \times (u-r)}$  be its orthogonal complements.

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- There exist  $\eta \in \mathbb{R}^{u \times p}$  such that  $\beta = \Gamma \eta$ , where  $\eta$  contains the coordinates of  $\beta$  relative to  $\Gamma$ .

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#### Envelope model

$$\mathbf{Y} = \mathbf{\Gamma} oldsymbol{\eta} \mathbf{X} + oldsymbol{arepsilon}, \ \ \mathbf{\Sigma} = \mathbf{\Gamma} \mathbf{\Omega} \mathbf{\Gamma}^{\mathcal{T}} + \mathbf{\Gamma}_0 \mathbf{\Omega}_0 \mathbf{\Gamma}_0^{\mathcal{T}}$$

Estimation of the parameters can be carried out by maximum likelihood, and u can be selected by AIC, BIC or other methods.



ullet The estimated envelope  $\hat{\mathcal{E}}_{\Sigma}(\mathcal{B})$  can be represented as

$$\hat{\mathcal{E}}_{\Sigma}(\mathcal{B}) = \arg\min_{\delta} (\log |\mathbf{P}_{\delta}\mathbf{S}_{\mathbf{Y}|\mathbf{X}}\mathbf{P}_{\delta}|_{0} + \log |\mathbf{Q}_{\delta}\mathbf{S}_{\mathbf{Y}}\mathbf{Q}_{\delta}|_{0}),$$

where where  $|\cdot|_0$  means the product of the non-zero eigenvalues and  $\delta$  is a u-dim subspace of  $\mathbb{R}^r$ .  $\mathbf{S}_{\mathbf{Y}|\mathbf{X}}$  and  $\mathbf{S}_{\mathbf{Y}}$  are the sample version of  $\mathbf{\Sigma}$  and  $\mathrm{Var}(\mathbf{Y})$ .

- $\hat{\mathcal{E}}_{\Sigma}(\mathcal{B})$  can be estimated through a 1-D algorithm proposed by Cook and Zhang (2016).
- Let  $\hat{\Gamma}$  denote the basis of  $\hat{\mathcal{E}}_{\Sigma}(\mathcal{B})$ , then estimators of regression parameters are
  - $\hat{eta}_{env} = \mathbf{P}_{\hat{\Gamma}} \hat{eta}_{std}$ , which is  $\sqrt{n}$ -consistent and asymptotically normal.
  - $\bullet \ \Sigma = \mathsf{P}_{\hat{\Gamma}}\mathsf{S}_{\mathsf{Y}|\mathsf{X}}\mathsf{P}_{\hat{\Gamma}} + \mathsf{Q}_{\hat{\Gamma}}\mathsf{S}_{\mathsf{Y}}\mathsf{Q}_{\hat{\Gamma}}$



# Efficiency gain

#### Proposition 2. (Cook et al., 2010)

$$\operatorname{avar}\{\sqrt{n}\mathrm{vec}(\hat{\beta}_{env})\} \leq \operatorname{avar}\{\sqrt{n}\mathrm{vec}(\hat{\beta}_{std})\}$$

where  $avar(\cdot)$  stands for the asymptotic covariance and  $vec(\cdot)$  stands for the vectorization of a matrix.

#### Remarks

- Envelope estimators are never worse than standard estimators in the sense of asymptotic variance.
- Envelope methods will provide the most gain in efficiency when  $\hat{\mathcal{E}}_{\Sigma}(\mathcal{B})$  can be constructed from eigenspaces of  $\Sigma$  with relatively small eigenvalues.

### An R Example

Illustration of an example using R:

```
require(envlp)
set.seed(0411)
num = 200
env dim <- 5
p = 5
q = 20
sq_err_env <- NULL
sq err std <- NULL
for (i in 1:10) {
  GAMMA \leftarrow matrix(runif(env dim * q), nrow = q)
  beta0 <- matrix(runif(p * q, -10, 10), nrow = p)
  beta <- beta0 %*% P(GAMMA)
  Omega <- 0.1 * diag(nrow(GAMMA))</pre>
  Omega0 <- 1000 * diag(nrow(GAMMA))</pre>
  Sigma_y <- P(GAMMA) %** Omega %** P(GAMMA) + Q(GAMMA) %** Omega0 %** Q(GAMMA)
  A \leftarrow matrix(runif(p ^ 2, -10, 10), nrow = p)
  mu_x < -runif(p, -10, 10)
  Sigma x <- A %*% t(A)
  X <- mvrnorm(num, mu_x, Sigma_x)</pre>
  Y \leftarrow X ** beta + mvrnorm(num, rep(0, g), Sigma v)
```

### An R example

```
u = u.env(X, Y)$u.bic
env_beta <- t(env(X, Y, u)$beta)
std_beta <- solve(crossprod(X)) %*% crossprod(X, Y)
sum((std_beta - beta)^2)
sq_err_env <- c(sum((env_beta - beta)^2), sq_err_env)
sq_err_std <- c(sum((std_beta - beta)^2), sq_err_std)
}
mean(sq_err_env)
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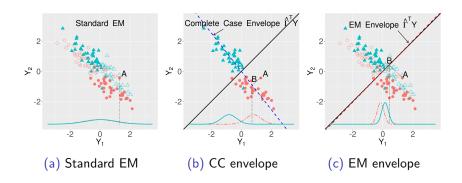
Output: 0.007500058 14.07488

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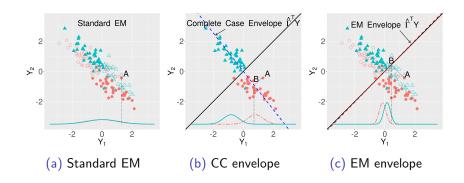
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  - MCAR: Missingness is independent both of observed and unobserved data.
  - MAR: Missingness is independent of unobserved data.
  - MNAR: is data that is not MAR, is also known as nonignorable missing.
- The former two mechanisms are addressed as ignorable missing since the missing information could be partially recovered using the observed data.

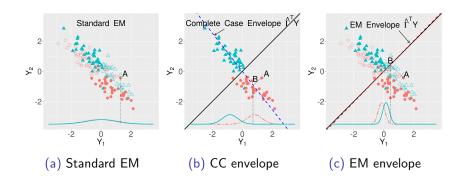
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- The former two mechanisms are addressed as ignorable missing since the missing information could be partially recovered using the observed data.
- Complete case analysis will introduce bias even if data is ignorable missing.



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- The *p*-values for each plot are 0.37, < 0.001, < 0.001.
- $\hat{\beta}_{std,EM}=0.11$  with standard error 0.12;  $\hat{\beta}_{cc,env}=-1.63$  with standard error 0.15;  $\hat{\beta}_{std,EM}=0.31$  with standard error 0.04.

# EM algorithm

$$I(\theta|\mathbf{x},\mathbf{y}) = \log(f_{\mathbf{y}|\mathbf{x}}(\mathbf{y}|\mathbf{x},\theta)) + \log(f_{\mathbf{x}}(\mathbf{x}|\theta))$$

$$= \sum_{i=1}^{n} \left( -\frac{1}{2}\log|\mathbf{\Sigma}| - \frac{1}{2}(\mathbf{y}_{i} - \mathbf{x}_{i}\beta)^{T}\mathbf{\Sigma}^{-1}(\mathbf{y}_{i} - \mathbf{x}_{i}\beta) + \log(f_{\mathbf{x}}(\mathbf{x}_{i}|\boldsymbol{\rho})) \right) + C$$

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**E-Step:** Let  $\theta=(\eta,\Gamma,\Omega_0,\Omega_1,\rho)$  and let  $\theta_t$  denote the current estimate of the parameter  $\theta$ . The E-step evaluate the expectation of full data likelihood given the current parameter estimates as

$$Q(\boldsymbol{\theta}|\boldsymbol{\theta}_t) = E[I_{full}(\boldsymbol{\theta}|\mathbf{X},\mathbf{Y})|\boldsymbol{D}_{obs},\boldsymbol{\theta}_t] = \int I_{full}(\boldsymbol{\theta}|\boldsymbol{L})f(\boldsymbol{D}_{mis}|\boldsymbol{D}_{obs},\boldsymbol{\theta}_t)d\boldsymbol{D}_{mis}.$$



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**M-Step:** The M-step computes  $\theta^{(t+1)}$  by maximizing the expected log-likelihood obtained in the E-step:

$$Q(\theta^{(t+1)}|\theta_t) \ge Q(\theta|\theta_t),$$
 for all  $\theta$ .

We iterate the E- and M-steps until convergence.

L.Ma, L.Liu, W.Yang

#### Remarks

• Since solving for envelope involves reparametrization of the covariance matrix, i.e.  $\Sigma = \mathbf{P}_{\Gamma}\Sigma\mathbf{P}_{\Gamma} + \mathbf{Q}_{\Gamma}\Sigma\mathbf{Q}_{\Gamma}$ , hence, it is non-trivial to combine EM algorithm and envelope moedels.

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- In E-step, we need to calculate the conditional expectations:

$$\begin{array}{l} \boldsymbol{A}_{i1,t} = \mathbb{E}(\boldsymbol{y_i'y_i}|\boldsymbol{\theta}_t, \boldsymbol{D}_{obs}), \ \boldsymbol{A}_{i2,t} = \mathbb{E}(\boldsymbol{y_i'x_i}|\boldsymbol{\theta}_t, \boldsymbol{D}_{obs}), \\ \boldsymbol{A}_{i3,t} = \mathbb{E}(\boldsymbol{x_i'x_i}|\boldsymbol{\theta}_t, \boldsymbol{D}_{obs}), \ \boldsymbol{A}_{i4,t} = \mathbb{E}(\boldsymbol{x_i'}|\boldsymbol{\theta}_t, \boldsymbol{D}_{obs}), \\ \text{Denote} \ \boldsymbol{A}_{j,t} = \sum_{i=1}^n \boldsymbol{A}_{ij,t}, \ j = 1, 2, 3, 4 \end{array}$$

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### Parameter updates

- Using the 1-D algorithm proposed by Cook and Zhang (2016) to estimate  $\Gamma_t$  based on  $\theta_t$ .
- $$\begin{split} \bullet \ \ & \Sigma_{1,t+1} = \frac{1}{n} P_{\Gamma_t} (\pmb{A}_{1,t} \pmb{A}_{2,t} \pmb{A}_{3,t}^{-1} \pmb{A}_{2,t}') P_{\Gamma_t}; \\ & \rho_{t+1} = \arg\max_{\rho \in \Theta} \mathbb{E}(\log(f_{\scriptscriptstyle X}(\pmb{x}_i|\rho))|\pmb{D}_{obs}, \theta_t); \\ & \beta_{t+1} = \pmb{A}_{3,t}^{-1} \pmb{A}_{2,t}' P_{\pmb{\Sigma}_{1,t+1}}; \\ & \Sigma_{t+1} = \pmb{\Sigma}_{1,t+1} + \frac{1}{n} Q_{\Gamma_t} \pmb{A}_{1,t} Q_{\Gamma_t} \end{split}$$

## **Algorithm 0:** The EM envelope algorithm

**Data:** n observations with MAR p predictors and q responses .

**Result:** Finding the estimator  $\hat{\beta}_{em\_env}$ .

for 
$$k = 1, 2, ..., q$$
 do

initialization: 
$$\Sigma_t = I_q$$
,  $\beta_t = 0$ ,  $\rho_t = \rho_0$ ,  $\theta_t = (\Sigma_t, \beta_t, \rho_t)$ ,  $\Delta = 1$ ;

while 
$$\Delta > \delta$$
 do

1. Calculate 
$$\mathbf{A}_{1,t} = \sum_{i=1}^{n} \mathbf{A}_{i1,t}$$
,  $\mathbf{A}_{2,t} = \sum_{i=1}^{n} \mathbf{A}_{i2,t}$ ,

$$\mathbf{A}_{3,t} = \sum_{i=1}^{n} \mathbf{A}_{i3,t}$$
 based on  $\mathbf{\theta}_t$ ;

2. Using 1-D algorithm to calculate  $\Gamma_t$ , then

$$\Sigma_{1,t+1} = \frac{1}{n} P_{\Gamma_t} (A_{1,t} - A_{2,t} A_{3,t}^{-1} A_{2,t}') P_{\Gamma_t};$$

3. Update:  $\rho_{t+1} = \arg\max_{\rho \in \Theta} \mathbb{E}(\log(f_x(\mathbf{x}_i|\rho))|\mathbf{D}_{obs}, \theta_t),$  $\beta_{t+1} = \mathbf{A}_{3,t}^{-1} \mathbf{A}_{2,t}' \mathbf{P}_{\Sigma_{1,t+1}}, \Sigma_{t+1} = \Sigma_{1,t+1} + \frac{1}{n} \mathbf{Q}_{\Gamma_t} \mathbf{A}_{1,t} \mathbf{Q}_{\Gamma_t};$ 

4. Set 
$$\Delta = \|\beta_{t+1} - \beta_t\|_1$$
,  $\theta_t = (\Sigma_{t+1}, \beta_{t+1}, \rho_{t+1})$ ;

end

$$\mathrm{BIC}_k = -2Q(\hat{\boldsymbol{\theta}}|\hat{\boldsymbol{\theta}}) + pu\log n, \ \hat{\boldsymbol{\beta}}_k = \boldsymbol{\beta}_{t+1}$$

#### end

Find u such that  $\mathrm{BIC}_k$  is minimum. The corresponding  $\hat{\beta}_u$  is the EM envelope estimator.

# EM envelope algorithm

### Proposition 1.

Denote  $\hat{\beta}_{env}$  as the estimator by EM envelope algorithm, and  $\hat{\beta}_{std}$  as the estimator by standard EM algorithm. Then, under some regularity conditions,

$$\sqrt{n} \{ \operatorname{vec}(\hat{\beta}_{env}) - \operatorname{vec}(\beta) \} \xrightarrow{d} N(\mathbf{0}, \mathbf{V}_{env}), 
\sqrt{n} \{ \operatorname{vec}(\hat{\beta}_{std}) - \operatorname{vec}(\beta) \} \xrightarrow{d} N(\mathbf{0}, \mathbf{V}_{std}), 
\text{where } \mathbf{V}_{env} \leq \mathbf{V}_{std}.$$

### **Simulations**

We run simulations to compare four different methods: standard complete case analysis, complete case envelope, standard EM, and EM envelope.

- Suppose  $\mathbf{X}_n \sim N(\boldsymbol{\mu}_{\scriptscriptstyle X}, \boldsymbol{\Sigma}_{\scriptscriptstyle X})$ ,  $\mathbf{Y}_n \sim N(\mathbf{X}_n \boldsymbol{\beta}, \boldsymbol{\Sigma}_{\epsilon})$
- Generate n = 500 samples, each has q = 20 responses, and p = 5 covariates with envelope dimension u = 3.
- ullet Generate  $\Gamma$ ,  $eta_0$ ,  $\mu_{\scriptscriptstyle X}$ ,  $\Sigma_{\scriptscriptstyle X}$  at random, and set  $eta=eta_0 {\sf P}_\Gamma$ .
- $\Sigma_{\epsilon} = \Gamma \Omega \Gamma' + \Gamma_0 \Omega_0 \Gamma'_0$ . Fix  $\Omega = 0.1 \mathbf{I}_q$ . We run two simulations when setting  $\Omega_0 = 1000 \mathbf{I}_q$  and  $\Omega_0 = 10 \mathbf{I}_q$ .
- $\bullet$  Each predictor suffers from about 10%-15% missingness, and each response about 5%-10%.

### Results

Table: Summary of MSE when  $\Omega_0=1000 I_a$ 

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
$\hat{eta}_{em\_env}$	1.64e-05	3.58e-05	4.44e-05	1.03e-03	5.70e-05	8.66e-02
$\hat{oldsymbol{eta}}_{ extsf{comp\_env}}$	3.28e-04	8.16e-04	1.27e-03	1.47e-02	2.52e-03	4.33
$\hat{oldsymbol{eta}}_{em\_std}$	2.37e-02	4.41e-02	5.34e-02	5.47e-02	6.38e-02	0.12
$\hat{oldsymbol{eta}}_{ extsf{comp\_std}}$	0.82	3.43	4.95	79.3	9.79	3.64e + 04

Table: Summary of MSE when  $\Omega_0=10 I_a$ 

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
$\hat{oldsymbol{eta}}_{em\_env}$	4.54e-05	9.08e-05	1.06e-04	1.36e-04	1.25e-04	1.05e-03
$\hat{oldsymbol{eta}}_{ extsf{comp\_env}}$	1.69e-03	4.50e-03	6.16e-03	0.590	1.24e-02	3.95e + 02
$\hat{oldsymbol{eta}}_{em\_std}$	2.17e-04	4.52e-04	5.42e-04	5.62e-04	6.49e-04	1.34e-03
$\hat{oldsymbol{eta}}_{ extsf{comp\_std}}$	1.14e-02	3.42e-02	5.06e-02	6.02	9.55e-02	4.60e+03

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- The CRIC study recruited 3939 participants from April 8, 2003 through September 3, 2008 and continued through March 31, 2013.
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#### More about the data set

- The CRIC study recruited 3939 participants from April 8, 2003 through September 3, 2008 and continued through March 31, 2013.
- The study cohort is racially and ethnically diverse group aged from 21 to 74 years with mild to moderate chronic kidney disease (CKD).
- It is of interest to investigate the difference in the distributions of baseline biomarkers among patients who develop end-stage renal diseases (ESRD) or not.

### In the regression setting

 $\bullet$  Predictors: ESRD status, gender, age, race, systolic and diastolic blood pressures, and hemoglobin from CBC lab data. (p =10)

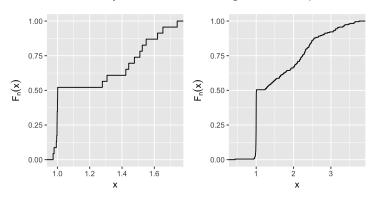
## In the regression setting

- ullet Predictors: ESRD status, gender, age, race, systolic and diastolic blood pressures, and hemoglobin from CBC lab data. (p = 10)
- Responses: Biomarkers, which are urine albumin, urine creatinine, high sensitivity C-reactive protein (HS\_CRP), brain natriuretic peptide (BNP), chemokine ligand 12 (CXCL12), fetuin A, fractalkine, myeloperoxidase (MPO), neutrophil gelatinase associated lipocalin (NGAL), fibrinogen, troponini, urine calcium, urine sodium, urine potassium, urine phosphate, high sensitive troponin T (TNTHS), aldosterone, C-peptide, insulin value, total parathyroid hormone (Total PTH), CO2, 24-hour urine protein, estimated glomerular filtration rate. (q = 23)
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- All the biomarkers have some missingness ranging from 0% to 6%.
- We compared EM envelope and the standard EM to examine their performance.

- Using BIC, we chose the envelope dimension u = 15.
- These two methods found the same set of biomarkers significant.
   However, using bootstrap method, the standard error of regression parameter are usually smaller when using EM envelope.



 It is found in the literature that although many novel biomarkers are found to be marginally significant associate with ESRD status, such association is muted after adjusting for glomerular filtration rate (GFR) and the amount of urine protein excreted in 24 hours.

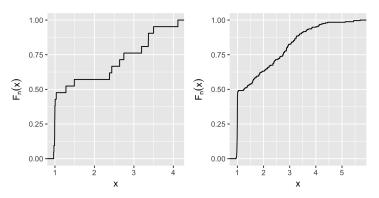
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- The estimated envelope dimension is u = 17.

Table: The point estimates, bootstrap standard errors, confidence intervals and p-values for biomarkers adjusted for the established biomarkers

	Our Method					Standard EM				
	$\hat{oldsymbol{eta}}$	SΈ	2.5%	97.5%	p-value	$\hat{oldsymbol{eta}}$	SΈ	2.5%	97.5%	<i>p</i> —value
HS_CRP	-0.04	0.02	-0.07	-2e-3	0.05	-0.12	0.07	-0.28	0.02	0.10
NGAL	-0.01	0.03	-0.07	0.04	0.69	0.18	0.07	0.06	0.31	6e-3
ALDOSTERONE	0.06	0.02	0.02	0.09	2e-3	0.04	0.04	-0.04	0.13	0.31
C_PEPTIDE	-0.10	0.04	-0.17	-0.03	9e-3	0.21	0.12	-0.02	0.44	0.08

Figure: The Empirical Cumulative Distribution of ratio between standard error of EM OLS and EM envelope for the coefficient corresponds to ESRD (left) and for all coefficients (right), adjusting for established biomarkers.



# More papers

- Scaled envelopes: scale-invariant and efficient estimation in multivariate linear regression. Biometrika, (2013).
- Envelopes and partial least squares regression. Journal of the Royal Statistical Society: Series B (Statistical Methodology), (2013).
- Foundations for envelope models and methods. Journal of the American Statistical Association, (2015).
- Simultaneous envelopes for multivariate linear regression.
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- Envelopes and reduced-rank regression. Biometrika, (2015).
- Algorithms for envelope estimation. Journal of Computational and Graphical Statistics, (2016).
- Sparse envelope model: efficient estimation and response variable selection in multivariate linear regression. Biometrika, (2016).
- Fast envelope algorithms. Statistica Sinica, (2018).

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Thank you!

- Cook, R. D., Li, B., and Chiaromonte, F. (2010). Envelope models for parsimonious and efficient multivariate linear regression. *Statistica Sinica*, pages 927–960.
- Cook, R. D. and Zhang, X. (2016). Algorithms for envelope estimation. *Journal of Computational and Graphical Statistics*, 25(1):284–300.