Sorted L-One Penalized Estimation (SLOPE)

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Outline

Genetic motivation

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Multiple Testing

Outline

- Genetic motivation
- Multiple Testing
- Model selection in multiple regression
 - Noiseless case Linear programming transition curve

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- Noisy case LASSO, SLOPE
- groupSLOPE
- further extensions



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Genetic variability

- ► About 99,9% of genetic information is the same for all people.
- A polymorphism is a difference in DNA structure, which is present in at least 1% of population
- A Single Nucleotide Polymorphism(SNP) is a polymorphism with the difference in the single base:
 - ► A typical SNP: a position in DNA in which
 - 85% of population has Cytosine(C)
 - 15% has a Thymine(T).
- ► There are usually two forms of a SNP at a given locus

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three genotypes : AA, Aa, aa.

Main purpose

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Y - quantitative trait

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Y - quantitative trait

Examples: blood pressure, cholesterol level, gene expression level, response to the treatment

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$$Y = (Y_1, \ldots, Y_n)^T$$
 - vector of trait values for n individuals

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 $Y = (Y_1, \dots, Y_n)^T$ - vector of trait values for *n* individuals $G_{n \times p}$ - matrix of SNP genotypes Usual coding for additive effects:

$$X_{ij} = \begin{cases} 0 & \text{if} \quad G_{ij} = aa \\ 1 & \text{if} \quad G_{ij} = Aa \\ 2 & \text{if} \quad G_{ij} = AA \end{cases}$$

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The dependency between SNPs is of a short range - even relatively close SNPs can be modeled as independent random variables.

Simple regression model for i-th gene:

$$Y_j = \beta_0 + \beta_i X_{ij} + \epsilon_j, \ \epsilon_j \sim N(0, \tau^2)$$

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H_0 true	U	V	p_0
H_0 false	Т	S	p_1
	W	R	m

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FWER = P(V > 0),		$FDR = E\left(\frac{1}{R}\right)$	$\left(\frac{\sqrt{2}}{\sqrt{1}}\right)$

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Simple regression model for i-th gene: $Y_i = \beta_0 + \beta_i X_{ii} + \epsilon_i, \ \epsilon_i \sim N(0, \tau^2)$ $\hat{\beta}_i$ - estimated β_i , $\hat{\beta}_i \sim N(\beta_i, \sigma_i^2)$, $i = 1, \dots, p$ $H_{0i}: \beta_i = 0$ vs $\beta_i \neq 0$ Reject H_{0i} when $|\hat{\beta}_i| > c_i$ Significance level: $\alpha = P_{H_{\alpha}}(|T_i| > c)$ H_0 accepted | H_0 rejected H_0 true U V p_0 Т S H_0 false p_1

 $FWER = P(V > 0), \quad FDR = E\left(\frac{V}{R \lor 1}\right)$

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WRmFWER = P(V > 0),FDR = E $\left(\frac{V}{R \lor 1}\right)$

$$E(V) = \alpha p_0$$

$$\alpha = 0.05, p_0 = 5000 \rightarrow E(V) = 250$$

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Multiple testing procedures

 T_1, \ldots, T_n independent $T_i \sim N(\mu_i, 1)$, $H_{0i}: \mu_i = 0$ vs $\mu_i \neq 0$ Bonferroni correction: Use significance level $\frac{\alpha}{p}$.

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Reject
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 if $|T_i| \ge \Phi^{-1} \left(1 - \frac{\alpha}{2p}\right) = \sqrt{2\log p} (1 + o(1))$

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Benjamini-Hochberg procedure:

(1)
$$|T|_{(1)} \ge |T|_{(2)} \ge \ldots \ge |T|_{(p)}$$

(2) Find the largest index *i* such that

$$|T|_{(i)} \ge \Phi^{-1}(1 - \alpha_i), \quad \alpha_i = \alpha \frac{i}{2p}, \tag{1}$$

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Call this index i_{SU} . (3) Reject all $H_{(i)}$'s for which $i \leq i_{SU}$

Bonferroni correction



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Benjamini and Hochberg correction



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For Bonferroni correction $\textit{FWER} \leq \alpha$

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$$\mathsf{FDR} = \mathbb{E}\left[\frac{V}{R \vee 1}\right] = \alpha \frac{p_0}{p},\tag{2}$$

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where p_0 is the number of true null hypotheses, $p_0 = |\{i: \mu_i = 0\}|$

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where p_0 is the number of true null hypotheses, $p_0 = |\{i : \mu_i = 0\}|$ (Benjamini, Yekutieli, 2001) If test statistics are dependent then BH controls FDR at the level $\alpha \frac{p_0}{p}$ if $|\mathcal{T}|_{(i)}$ is compared with

$$\sigma\Phi^{-1}\left(1-\frac{i\alpha}{p\sum_{i=1}^{p}\frac{1}{i}}\right).$$

Abramovich, Benjamini, Donoho and Johnstone, Ann.Statist. 2006

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 $T_i \sim N(\beta_i, 1)$, independent for $i = 1, \dots, p$

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Goal - estimation of $\beta = (\beta_1, \ldots, \beta_p)$

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 $\tilde{\beta}_i = T_i$ if BH rejects $H_{0i}: \beta_i = 0$

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 $\tilde{\beta}_i = T_i$ if BH rejects $H_{0i}: \beta_i = 0$

 $ilde{eta}_i = 0$ otherwise
$$heta = rac{p-p_0}{p}$$
 - proportion of alternative hypotheses,

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 - proportion of alternative hypotheses, $q_{
ho}$ - nominal FDR level

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$$R_{p} = \inf_{\hat{\beta}} \sup_{\beta:\theta < \eta_{p}} E||\hat{\beta} - \beta||^{r}$$

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$$R_{p} = \inf_{\hat{\beta}} \sup_{\beta:\theta < \eta_{p}} E||\hat{\beta} - \beta||^{r}$$

Theorem

Assumptions:

> q_p → c ∈ [0, 1/2] and q_p > γ/log(p) for some γ > 0
η_p ∈ [p⁻¹ log⁵ p, p^{-δ}], δ > 0
r ∈ (0, 2]
$$\sup_{\beta:\theta < \eta_p} E||\tilde{\beta} - \beta||^r = R_p(1 + o(1))$$

Bogdan, Chakrabarti, Frommlet, Ghosh, Ann.Statist. 2011 Neuvial, Roquain, Ann. Statist. 2012

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$$Y = X_{n \times p} \beta$$



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 $k = ||eta||_0 \leq \textit{min}(n,p)$ - number of nonzero elements in eta

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Theorem

If X_{ij} are iid $N(0, \sigma^2)$ then with probability converging to 1 the linear program can identify the true solution if $\rho < \rho(\delta)$. If $\rho > \rho(\delta)$ then the probability of recovering the true solution converges to 0.

Transition curve (2)

Phase Transition: (l_1, l_0) equivalence



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Noisy case - statistical problem

$$Y_{n\times 1} = X_{n\times p}\beta_{p\times 1} + z_{n\times 1}, \ z \sim N(0, \sigma I)$$



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Convex program: Minimize $||b||_1$ subject to $||Y - Xb||_2^2 \le \epsilon$ Or alternatively: $min_{b \in R^p} \frac{1}{2} ||y - Xb||_2^2 + \lambda ||b||_1$ Noisy case - statistical problem

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In statistics this procedure is called LASSO (Tibshirani, 1996)

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Selection of the tuning parameter for LASSO

- General rule: the reduction of λ_L results in identification of more elements from the true support (true discoveries) but at the same time it produces more falsely identified variables (false discoveries)
- ► The choice of \u03c6_L is challenging- e.g. crossvalidation typically leads to many false discoveries
- When $X^T X = I$ Lasso selects X_j iff $|\hat{\beta}_j^{LS}| > \lambda$
- Selection λ = σΦ⁻¹(1 − α/(2p)) ≈ σ√2 log p corresponds to Bonferroni correction and controls FWER.

SLOPE (Bogdan, van den Berg, Sabatti, Su and Candès, AOAS, 2015)

 SLOPE is an extension of LASSO aimed at control of FDR rather than FWER

SLOPE is defined as solution to

$$\beta_{SL} = \operatorname{argmin}_{b} \left\{ \frac{1}{2} \left\| y - Xb \right\|_{2}^{2} + \sum_{i=1}^{p} \lambda_{i} \left| b \right|_{(i)} \right\}, \qquad (\mathsf{SLOPE})$$

where $|b|_{(1)} \ge \ldots \ge |b|_{(\rho)}$ are ordered magnitudes of coefficients of b and $\lambda_1 \ge \ldots \ge \lambda_{\rho} \ge 0$ is the sequence of tuning parameters

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$$\beta_{SL} = \operatorname{argmin}_{b} \left\{ \frac{1}{2} \left\| y - Xb \right\|_{2}^{2} + \sum_{i=1}^{p} \lambda_{i} \left| b \right|_{(i)} \right\}, \qquad (\mathsf{SLOPE})$$

where $|b|_{(1)} \ge \ldots \ge |b|_{(p)}$ are ordered magnitudes of coefficients of b and $\lambda_1 \ge \ldots \ge \lambda_p \ge 0$ is the sequence of tuning parameters

The above optimization problem is convex and can be efficiently solved even for large design matrices SLOPE (Bogdan, van den Berg, Sabatti, Su and Candès, AOAS, 2015)

 $\ensuremath{\mathsf{SLOPE}}$ is an extension of LASSO aimed at control of FDR rather than FWER

SLOPE is defined as solution to

$$\beta_{SL} = \operatorname{argmin}_{b} \left\{ \frac{1}{2} \left\| y - Xb \right\|_{2}^{2} + \sum_{i=1}^{p} \lambda_{i} \left| b \right|_{(i)} \right\}, \qquad (\mathsf{SLOPE})$$

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- The above optimization problem is convex and can be efficiently solved even for large design matrices
- ► Sorted L-One Norm: $J_{\lambda}(b) = \sum_{i=1}^{p} \lambda_i |b|_{(i)}$ reduces to $||b||_1$ if $\lambda_1 = \ldots = \lambda_p$ and to $||b||_{\infty}$ if $\lambda_1 > \lambda_2 = \ldots = \lambda_p = 0$

Unit balls for different SLOPE sequences by D.Brzyski



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• Let $\widetilde{\beta}$ be estimate of β

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- Let $\widetilde{\beta}$ be estimate of β
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► the number of false discoveries,

 $V := \left| \left\{ i : \beta_i = 0, \quad \widehat{\beta}_i \neq 0 \right\} \right|$

- Let $\widetilde{\beta}$ be estimate of β
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- ► the number of false discoveries, $V := |\{i : \beta_i = 0, \quad \widetilde{\beta}_i \neq 0\}|$
- false discovery rate, $FDR := \mathbb{E}\left[\frac{V}{\max\{R,1\}}\right]$

FDR control with SLOPE

Theorem When $X^T X = I$ SLOPE with $\lambda_i := \sigma \Phi^{-1} \left(1 - i \cdot \frac{q}{2p} \right)$

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controls FDR at the level $q \frac{p_0}{p}$.

Asymptotic optimality, Su and Candès (Annals of Statistics, 2016)

Theorem

Let $X_{ij} \sim N(0, 1/\sqrt{n})$. Fix 0 < q < 1 and choose $\lambda = \sigma(1 + \epsilon)\lambda^{BH}(q)$ for some arbitrary constant $0 < \epsilon < 1$. Suppose $k/p \rightarrow 0$ and $\frac{k \log p}{n} \rightarrow 0$. Then

$$\begin{split} \sup_{\substack{||\beta_0|| \le k}} P\left(\frac{||\hat{\beta}_{SL} - \beta||^2}{2\sigma^2 k \log(p/k)} > 1 + 3\epsilon\right) \to 0\\ \inf_{\hat{\beta}} \sup_{||\beta_0|| \le k} P\left(\frac{||\hat{\beta} - \beta||^2}{2\sigma^2 k \log(p/k)} > 1 - \epsilon\right) \to 1 \end{split}$$

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Random designs - problems with shrinkage

$$\hat{eta} = \eta_{\lambda}(eta_i + X'_i z + v_i)$$

 $v_i = \langle X_i, \sum_{j \neq i} X_j(eta_j - \hat{eta}_j)
angle$

 $\eta_\lambda(t) = \mathrm{sgn}(t)(|t|-\lambda)_+,$ applied componentwise

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Random designs - problems with shrinkage

$$\hat{eta} = \eta_{\lambda}(eta_i + X'_i z + v_i)$$

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angle$

 $\eta_{\lambda}(t) = \operatorname{sgn}(t)(|t| - \lambda)_{+}, \text{ applied componentwise}$ If $X^{T}X = I$ then $X'_{i}z = Z_{i} \sim N(0, 1), v_{i} = 0$ and H_{0i} is rejected if $\beta_{i} + Z_{i} > \lambda$

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Random designs - problems with shrinkage

$$egin{aligned} \hat{eta} &= \eta_\lambda (eta_i + X_i'z + v_i) \ v_i &= \langle X_i, \sum_{j
eq i} X_j (eta_j - \hat{eta}_j)
angle \end{aligned}$$

 $\eta_\lambda(t) = \mathrm{sgn}(t)(|t|-\lambda)_+, \hspace{1em} \mathsf{applied} \hspace{1em} \mathsf{componentwise}$

If $X^T X = I$ then $X'_i z = Z_i \sim N(0, 1)$, $v_i = 0$ and H_{0i} is rejected if $\beta_i + Z_i > \lambda$

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When the design is not orthogonal: $v_i \neq 0$ - additional noise, dependent on λ (level of shrinkage) and the level of sparsity

Random designs - problems with shrinkage

$$\hat{eta} = \eta_{\lambda}(eta_i + X'_i z + v_i)$$

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angle$

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If $X^T X = I$ then $X'_i z = Z_i \sim N(0, 1)$, $v_i = 0$ and H_{0i} is rejected if $\beta_i + Z_i > \lambda$

When the design is not orthogonal: $v_i \neq 0$ - additional noise, dependent on λ (level of shrinkage) and the level of sparsity Quantification for LASSO using AMP theory - Bayati and Montanari (IEEE Trans. Infom.Theory, 2011)

Limits on FDR control

- Bogdan, van den Berg, Su and Candes (2013, arxive)
- Bogdan, Su and Candes (2017, to appear in Ann. Statist)



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Thresholds, q = 0.2, p = 5000 (1)



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Thresholds, q = 0.05, p = 5000 (2)



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FDR, p = n = 5000, Gaussian design



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MSE, p = 262144, n = p/2, k = 10, Gaussian design



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MSE, p = 262144, n = p/2, k = 1000, Gaussian design



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geneSLOPE - application for full GWAS data

Brzyski, Peterson, Sobczyk, Bogdan, Candés, Sabatti (Genetics, 2017)



Power



Group SLOPE

Identification of groups of predictors (Brzyski, Gossmann, Su, Bogdan, arxiv 2016, under revision for JASA, ENAR Young Researcher Award for Damian Brzyski, March 2017):

$$[[\beta]]_{I} := \left(\|X_{I_{1}}\beta_{I_{1}}\|_{2}, \dots, \|X_{I_{m}}\beta_{I_{m}}\|_{2} \right)^{\mathsf{T}} .$$

$$\beta^{gS} := \operatorname{argmin}_{b} \left\{ \frac{1}{2} \|y - Xb\|_{2}^{2} + \sigma J_{\lambda} (W[[b]]_{I}) \right\},$$

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where W is a diagonal matrix with $W_{i,i} := w_i$, for $i = 1, \ldots, m$.

Group FDR

$$Rg := \left| \left\{ i : \|X_{I_i} \beta_{I_i}^{gS}\|_2 \neq 0 \right\} \right|$$
$$Vg := \left| \left\{ i : \|X_{I_i} \beta_{I_i}\|_2 = 0, \|X_{I_i} \beta_{I_i}^{gS}\|_2 \neq 0 \right\}$$

We define the false discovery rate for groups (gFDR) as

$$gFDR := \mathbb{E}\left[rac{Vg}{\max\{Rg,1\}}
ight]$$

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gFDR control

Theorem

Let the design matrix X satisfy $X_{I_i}^{\mathsf{T}} X_{I_j} = 0$, for any $i \neq j$. Denote the number of zero coefficients in $[[\beta]]_i$ by m_0 and let w_1, \ldots, w_m be positive numbers. Moreover, define the sequence of regularizing parameters $\lambda^{\max} = (\lambda_1^{\max}, \ldots, \lambda_m^{\max})^{\mathsf{T}}$, with

$$\lambda_i^{\max} := \max_{j=1,\dots,m} \left\{ \frac{1}{w_j} F_{\chi_{l_j}}^{-1} \left(1 - \frac{q \cdot i}{m} \right) \right\},\tag{3}$$

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where $F_{\chi_{l_j}}$ is a cumulative distribution function of χ distribution with l_j degrees of freedom. Then any solution, $\beta^{\text{ES}(gS)}$, to gSLOPE generates the same vector $[[\beta^{gS}]]_l$ and it holds

$$gFDR = \mathbb{E}\left[\frac{Vg}{\max\{Rg,1\}}
ight] \leq q \cdot \frac{m_0}{m}.$$



Averaged λ

$$\lambda_r^{mean} := \overline{F}^{-1} \left(1 - \frac{qr}{m} \right) \quad \text{for} \quad \overline{F}(x) := \frac{1}{m} \sum_{i=1}^m F_{w_i^{-1} \chi_{l_i}}(x) \quad , \quad (4)$$

where $F_{w_i^{-1}\chi_{l_i}}$ is the cumulative distribution function of scaled chi distribution with l_i degrees of freedom and scale $S = w_i^{-1}$. Gossmann, Brzyski et al. (2017) - Proof of FDR control in case when averaging is with respect to the distribution generating groups of different sizes and the signal is randomly placed between groups.

Simulations under Gaussian design, n = 5000, m = 1000, p = 7917



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Applications for GWAS

n = 5402, p = 26233 - roughly independent SNPs

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Applications for GWAS

n = 5402, p = 26233 - roughly independent SNPs

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Scenario 1: $Y = X\beta + z$ - additive model

Applications for GWAS

n = 5402, p = 26233 - roughly independent SNPs Scenario 1: $Y = X\beta + z$ - additive model Scenario 2: modeling dominance

$$\widetilde{z}_{ij} = \left\{ egin{array}{ccc} -1 & ext{for} & aa, AA \ 1 & ext{for} & aA \end{array}
ight.,$$
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 $y = [X, Z][\beta'_X, \beta'_Z]' + \epsilon$.



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Genes Influencing Level of Triglicerides

5 new discoveries with group SLOPE - recessive rare genetic variants. Discovery 5 - 37 rare homozygotes



Logistic SLOPE (Michal Kos, Sangkyun Lee, M.Bogdan *I(b)* - log-likelihood

$$\min_{b} \left(-l(b) + \sum \lambda_i |b_{(i)}| \right)$$

 $\lambda_i = 0.5 \Phi^{-1} \left(1 - \frac{qi}{2p} \right)$

Asymptotic FDR control when X is a random design such that

- for all i, j: x_{ij} are independent and $Ex_{ij} = 0$
- elements in each column of design matrix X are i.i.d.

•
$$\mathbb{E}[X^T X] = I \ (\Rightarrow var(x_{ij}) = n^{-1})$$

there exist constant M that for all i and j:

$$\frac{|x_{ij}|}{\sqrt{\operatorname{var}(x_{ij})}} = x_{ij}\sqrt{n} \leqslant M$$

• p and β are fixed, $n \to \infty$

FDR



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Power



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Portfolio Optimization, (Philipp Kremmer, Sangkyun Lee, M. Bogdan, Sandra Paterlini)

$$R_{t \times k} = (R_1, \dots, R_k)$$
 - asset returns,
 $E(R) = \mu, R = F_{t \times r} B_{r \times k}, r \ll k$

$$\min_{w \in \mathbb{R}^k} \frac{\phi}{2} w' \Sigma w - \mu' w + J_{\lambda}(w)$$
 (6)

$$\text{s.t.}\sum_{i=1}^{k} w_i = 1 \tag{7}$$

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Evolution of Portfolio



Other results/current work

- 1. Brzyski, Gossmann, Su, Bogdan Group SLOPE, under revision for JASA
- Lee, Brzyski, Bogdan Ordered Dantzig Selector, AISTAT 2016
- 3. Kos, Lee, Bogdan Logistic regression asymptotic FDR control under random designs (p fixed, $n \rightarrow \infty$).
- 4. Lee, Sobczyk, Bogdan Graphical models FDR control at the block level

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4. Kremmer, Lee, Bogdan, Paterlini - Portfolio optimization - prediction under factorial (correlated) designs