

Ultrahigh dimensional variable selection: Beyond the linear model

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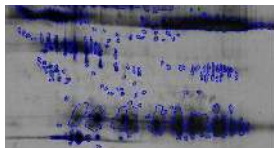
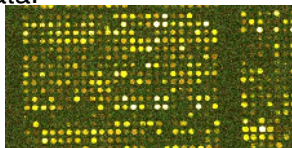
Introduction



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High-dim variable selection characterizes many contemporary statistical problems.

- Bioinformatic: disease classification using microarray, proteomics, fMRI data.



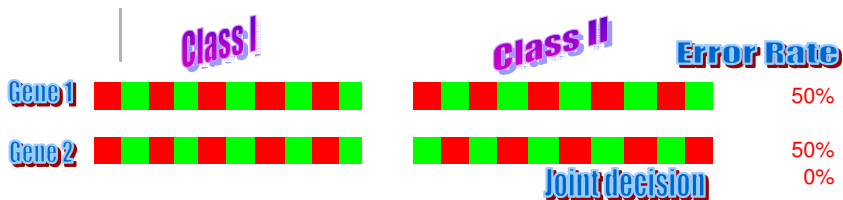
- Document or text classification: E-mail spam.
- Association studies between phenotypes and SNPs.



Growth of Dimensionality

- Dimensionality grows rapidly with interactions

Portfolio selection and network modeling: 2,000 stocks involves over 2m unknown parameters in the covariance matrix.



Gene-gene interaction: interactions of 5000 genes result in 12.5 features.



Aims of High-dimensional Regression and Classification

- To construct as effective a method as possible to predict future observations.
- To gain insight into the relationship between features and response for scientific purposes, as well as, hopefully, to construct an improved prediction method.

Bickel (2008) discussion of the SIS paper (JRSS-B).



Challenges with Ultrahigh Dimensionality

■ Computational cost

■ Estimation accuracy.

■ Stability



Key idea: **Large-scale** screening and **moderate-scale** searching.



Large-scale sreening



Independence learning

Regression: Feature ranking by **correlation learning** (Fan and Lv, 2008, JRSS-B). When $Y = \pm 1$, this implies

Classification: Feature ranking by two-sample t-tests or other tests (Tibshirani, et al, 03; Fan and Fan, 2008).

SIS: By an appropriate thresholding (e.g., n variables), **relevant features are in the selected set** (Fan and Lv, 08), relying on joint-normality assumption.

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Model setting

GLIM: $f_Y(y|X = \mathbf{x}; \theta) = \exp\{(y\theta - b(\theta))/\phi + c(y, \phi)\}$ with

$$\text{canonical link : } b'^{-1}(\mu) = \theta = \mathbf{x}^T \beta.$$

Objective: Find sparse β to minimize $Q(\beta) = \sum_{i=1}^n L(Y_i, \mathbf{x}_i^T \beta)$.

- **GLIM:** $L(Y_i, \mathbf{x}_i^T \beta) = b(\mathbf{x}_i^T \beta) - Y_i \mathbf{x}_i^T \beta$.
- **Classification:** $Y = \pm 1$.
 - ★ SVM $L(Y_i, \mathbf{x}_i^T \beta) = (1 - Y_i \mathbf{x}_i^T \beta)_+$.
 - ★ AdaBoost $L(Y_i, \mathbf{x}_i^T \beta) = \exp(-Y_i \mathbf{x}_i^T \beta)$.
- **Robustness:** $L(Y_i, \mathbf{x}_i^T \beta) = |Y_i - \mathbf{x}_i^T \beta|$.



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Questions

- 1 How to screen **discrete** variables (Genome-wide association)?
- 2 Do they have **sure screening** property?
- 3 What is the size of selected model in order to have SIS?

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$$\hat{L}_j = \hat{L}_0 - \min_{\beta_0, \beta_j} n^{-1} \sum_{i=1}^n L(Y_i, \beta_0 + X_{ij}\beta_j) \quad \text{Wilks.}$$

or $\hat{\beta}_j^M$ (**Wald**), assuming $EX_j^2 = 1$.

Feature ranking: Select features w/ **largest marginal utilities**:

$$\widehat{\mathcal{M}}_{v_n} = \{j : \hat{L}_j \geq v_n\}, \quad \widehat{\mathcal{M}}_{\gamma_n}^w = \{j : \hat{\beta}_j^M \geq \gamma_n\}$$

Dim. reduction: From $p_n = O(\exp(n^a))$ to $O(n^b)$:



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Theoretical Basis – Population Aspect I

Marginal utility: $L_j^* = E\ell(Y, \beta_0^M) - \min E\ell(Y, \beta_0 + \beta_j X_j)$.

Likelihood ratio (Fan and Song, 09)

Theorem 1: $L_j^* = 0 \iff \text{cov}(Y, X_j) = \text{cov}(b'(\mathbf{X}^T \beta^*), X_j) = 0$
 $\iff \beta_j^M = 0$.

For Gaussian covariates, conclusion holds if $|\text{cov}(\mathbf{X}^T \beta^*, X_j)| = 0$, i.e. independence.



Theoretical Basis – Population Aspect II

True model: $\mathcal{M}_\star = \{j : \beta_j^\star \neq 0\}$, where $\beta^\star = \operatorname{argmin} EL(Y, \mathbf{X}^T \beta)$.

Theorem 2: If $|\operatorname{cov}(b'(\mathbf{X}^T \beta^\star), X_j)| \geq c_1 n^{-\kappa}$ for $j \in \mathcal{M}_\star$, then

$$\min_{j \in \mathcal{M}_\star} |\beta_j^M| \geq c_1 n^{-\kappa}, \quad \min_{j \in \mathcal{M}_\star} |L_j^\star| \geq c_2 n^{-2\kappa}.$$

If $\{X_j, j \notin \mathcal{M}_\star\}$ is independent of $\{X_i, i \in \mathcal{M}_\star\}$, then $L_j^\star = 0$.

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$$|\operatorname{cov}(\mathbf{X}^T \beta^\star, X_j)| \geq c_1 n^{-\kappa}, \quad \text{min condition even for LS.}$$



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Sampling Aspect: Sure independence screening

Theorem 3: If $v_n = cn^{-2\kappa}$ for $\kappa < 1/2$, and $\log s_n = o(n^{1-2\kappa})$, then

$$P\left(\mathcal{M}_* \subset \widehat{\mathcal{M}}_{v_n}\right) \rightarrow 1 \quad \text{exponentially fast}$$

No conditions on covariance matrix!

- This is a SIS property w/ size controlled.
- Note that $\widehat{L}_j - L_j^* = O(\log p/n^{1/2})$ and minimum signal $O(n^{-2\kappa})$.

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Screening by MMLE

Let $\widehat{\mathcal{M}}_{\gamma_n}^w = \{|\widehat{\beta}_j^M| \geq \gamma_n\}$.

① $P(\max_j |\widehat{\beta}_j^M - \beta_j^M| > c_3 n^{-\kappa}) = o(1)$, if $\log p_n = o(n^{1-2\kappa})$.

② $P(\mathcal{M}_* \subset \widehat{\mathcal{M}}_{\gamma_n}^w) \rightarrow 1$, if $\gamma_n = c_0 n^{-\kappa}$, $c_0 < c_1/2$.

③ What is the selected model size? We establish

$$\|\beta^M\|^2 = o(\|\Sigma\beta^*\|^2) = O\{\lambda_{\max}(\Sigma) \beta^{*T} \Sigma \beta^*\} = O(\lambda_{\max}(\Sigma)).$$

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Sampling Aspect: Controlling number of features

Theorem 4: If $\log p_n = o(n^{1-2\kappa})$,

$$\mathbf{P}[|\widehat{\mathcal{M}}_{V_n}| \leq \mathbf{O}\{n^{2\kappa}\lambda_{\max}(\Sigma)\}] \rightarrow \mathbf{1}.$$

■ Establish $\|\mathbf{L}^*\|^2 = o(\|\beta^M\|^2) = o(\|\Sigma\beta^*\|^2)$.

■ The number of selected covariates depends on the population covariance. It is actually bounded by

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Moderate-scale selection

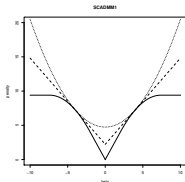


Moderate-scale of Model Selectors

■ **Penalized lik.:** $n^{-1} \sum_{i=1}^n L(Y_i, \beta_0 + \mathbf{x}_{i,d}^T \beta) + \sum_{j=1}^d \rho_\lambda(|\beta_j|)$.

Simultaneously estimate coefs and choose variables.

- Lasso (Tibshirani, 96), LARS (Efron *et al.*, 04), Adaptive Lasso (Zou, 06), Approx sparse (Huang and Zhang, 06).
- SCAD (Fan & Li, 01, 06; Fan & Peng, 04)
LQA (Fan & Li, 01), MM (Hunter & Li, 05),
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■ **Dantzig selector** (Candes & Tao, 07)

$$\min_{\beta \in \mathbb{R}^p} \|\beta\|_1 \quad \text{subject to} \quad \|\mathbf{X}^T \mathbf{r}\|_\infty \leq \lambda_{p,n} \sigma$$

with $\lambda_{p,n} > 0$, $\mathbf{r} = \mathbf{y} - \mathbf{X}\beta$ and σ noise level. \approx **Lasso** (Bickel, et al, 2008)

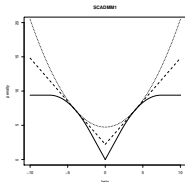


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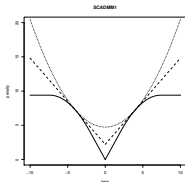


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Connections among penalized least-squares

■ **PLS**: $\|\mathbf{y} - \mathbf{X}\beta\|^2 + \sum_{i=1}^{p_n} \rho_\lambda(|\beta_i|)$.

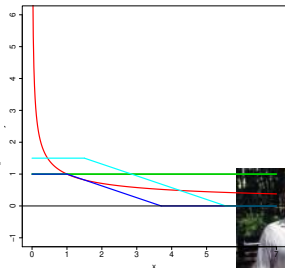
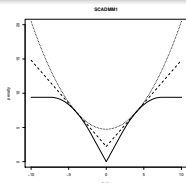
LLA: with initial value β_0 (Zou & Li, 08),

$$\|\mathbf{y} - \mathbf{X}\beta\|^2 + \sum_{i=1}^{p_n} \{\rho_\lambda(|\beta_{i,0}|) + \rho_\lambda(|\beta_{i,0}|)'(|\beta_i| - |\beta_{i,0}|)\}.$$

Weighted L_1 : $\|\mathbf{y} - \mathbf{X}\beta\|^2 + \sum_{i=1}^{p_n} \mathbf{w}(|\beta_{i,0}|)|\beta_i|$.

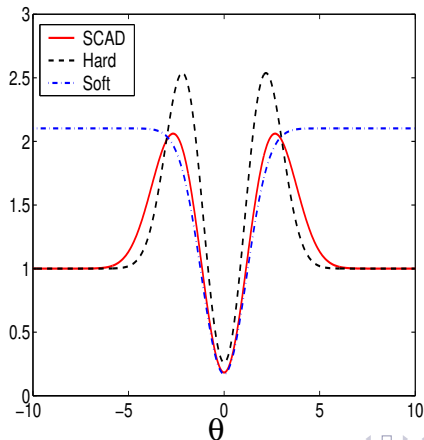
Fan and Li (01) stressed the unbiasedness.

Convergence: Objective function decreasing.



Risk Comparisons of popularized least-squares

- Penalized least-squares: $(Z - \theta)^2 + p_\lambda(|\theta|)$
- $R(\hat{\theta}, \theta) = E_\theta(\hat{\theta} - \theta)^2$ with $Z \sim N(\theta, 1)$
- $\lambda = 2$ for hard thresholding



Iterative feature selection



Drawback of Independence Screening

False negative: The features such that $\text{cov}(X_j, \mathbf{X}^T \beta^*) = 0$ can not be selected, but this can be a **signature variable**.

Example: If $\{X_j\}_{j=1}^J$ has common correlation ρ , then

$$\text{cov}(X_{J+1}, X_1 + \cdots + X_J - J\rho X_{J+1}) = 0.$$

False positive: Rank too high predictors **jointly unimportant** but marginally important:

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- 1 ■ **(Large-scale screening)**: Apply SIS to pick a set \mathcal{A}_1 ;
■ **(Moderate-scale selection)**: Employ a penalized likelihood to select a subset \mathcal{M}_1 of these indices.
- 2 **(Large-scale screening)**: Rank features according to the additional **(conditional)** contribution:

$$L_j^{(2)} = \min_{\beta_0, \beta_{\mathcal{M}_1}, \beta_j} n^{-1} \sum_{i=1}^n L(Y_i, \beta_0 + \mathbf{x}_{i, \mathcal{M}_1}^T \beta_{\mathcal{M}_1} + X_{ij} \beta_j).$$

—Resulting in new feature sets \mathcal{A}_2 .

—An improvement over Fan and Lv (08) who set $\beta_{\mathcal{M}_1} = \hat{\beta}_{\mathcal{M}_1}$ from previous fit.



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Iterative feature selection (II)

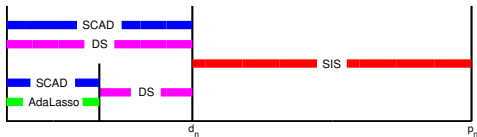
- 3 (Moderate-scale selection): Minimize wrt $\beta_{\mathcal{M}_1}, \beta_{\mathcal{A}_2}$

$$\sum_{i=1}^n L(Y_i, \beta_0 + \mathbf{x}_{i, \mathcal{M}_1}^T \beta_{\mathcal{M}_1} + \mathbf{x}_{i, \mathcal{A}_2}^T \beta_{\mathcal{A}_2}) + \sum_{j \in \mathcal{M}_1 \cup \mathcal{A}_2} \rho_\lambda(|\beta_j|).$$

—Resulting in \mathcal{M}_2

—**Allow deletion**, improvement over ISIS (Fan and Lv, 08).

- 4 Repeat Steps 1–3 until $|\mathcal{M}_\ell| = d$ (prescribed) or $\mathcal{M}_\ell = \mathcal{M}_{\ell-1}$.



Reduction of false selection rates

Variant 1: Randomly split samples to obtain $\hat{\mathcal{A}}^{(1)}$ and $\hat{\mathcal{A}}^{(2)}$.

Take $\hat{\mathcal{A}} = \hat{\mathcal{A}}^{(1)} \cap \hat{\mathcal{A}}^{(2)}$.

Intuition: If both have SIS property, so does $\hat{\mathcal{A}}$ with lower FSR.

Theorem 1: With prescribed d ,

$$P(|\hat{\mathcal{A}} \cap \mathcal{M}_*^c| \geq r) \leq \frac{\binom{d}{r}^2}{\binom{p-|\mathcal{M}_*|}{r}} \leq \frac{1}{r!} \left(\frac{d^2}{p-|\mathcal{M}_*|} \right)^r,$$

—Blessing of dimensionality!

Variant 2: Recruit as many variables into equal-sized sets $\tilde{\mathcal{A}}^{(1)}$ and $\tilde{\mathcal{A}}^{(2)}$ as required such that $|\hat{\mathcal{A}}| = d$ (prescribed).



Reduction of false selection rates

Variant 1: Randomly split samples to obtain $\hat{\mathcal{A}}^{(1)}$ and $\hat{\mathcal{A}}^{(2)}$.

Take $\hat{\mathcal{A}} = \hat{\mathcal{A}}^{(1)} \cap \hat{\mathcal{A}}^{(2)}$.

Intuition: If both have SIS property, so does $\hat{\mathcal{A}}$ with lower FSR.

Theorem 1: With prescribed d ,

$$P(|\hat{\mathcal{A}} \cap \mathcal{M}_*^c| \geq r) \leq \frac{\binom{d}{r}^2}{\binom{p-|\mathcal{M}_*|}{r}} \leq \frac{1}{r!} \left(\frac{d^2}{p-|\mathcal{M}_*|} \right)^r,$$

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Numerical Studies



Design of Simulations

Contexts: ★ Logistic ★ Poisson ★ L_1 -reg; ★ Multiclass SVM

Covariates: $p = 1000$, $X_i \sim N(0, 1)$.

- 1 $X_1, \dots, X_p \sim_{i.i.d.} N(0, 1)$
- 2 $\text{corr}(X_i, X_4) = 1/\sqrt{2}$ and otherwise $\text{corr}(X_i, X_j) = 1/2$.
- 3 The same except $\text{corr}(X_i, X_{p+1}) = 0$.



Logistic regression, independent covariate

$\beta_1 = 1.24, \beta_2 = -1.34, \beta_3 = -1.35, \beta_4 = -1.80, \beta_5 = -1.58, \beta_6 = -1.60.$

Bayes test error: **0.1368**. $n = 400, N_{\text{sim}} = 100.$

	SIS	ISIS	Var2-SIS	LASSO	NSC
$\text{med}(\ \beta - \hat{\beta}\ _1)$	1.11	1.25	1.21	8.48	N/A
$\text{med}(\ \beta - \hat{\beta}\ _2^2)$	0.49	0.52	0.52	1.70	N/A
True positive	0.99	0.84	0.91	1.00	0.34
Med. model size	6	6	6	94	3
$2Q(\hat{\beta}_0, \hat{\beta})$ (training)	237	247	243	164	N/A
AIC	250	260	256	353	N/A
BIC	278	285	282	725	N/A
$2Q(\hat{\beta}_0, \hat{\beta})$ (test)	272	273	273	319	N/A
0-1 test error	0.14	0.14	0.14	0.17	0.36



Logistic regression, difficult case — false negative

$$\beta_1 = 4, \beta_2 = 4, \beta_3 = 4, \beta_4 = -6\sqrt{2}, \text{cov}(X_4, \mathbf{X}^T \beta^*) = 0.$$

Signature variable: Bayes error: **0.107** and **.344** w/ and w/o X_4 .

	Van-SIS	ISIS	Var2-ISIS	LASSO	NSC
$\text{med}(\ \beta - \hat{\beta}\ _1)$	20.1	1.94	1.85	21.6	N/A
$\text{med}(\ \beta - \hat{\beta}\ _2^2)$	9.41	1.05	0.98	9.11	N/A
True positive	0.00	1.00	1.00	0.00	0.21
Med. model size	16	4	4	91	16.5
$2Q(\hat{\beta}_0, \hat{\beta})(\text{training})$	307	187	187	127	N/A
AIC	334	196	195	311	N/A
BIC	386	212	212	672	N/A
$2Q(\hat{\beta}_0, \hat{\beta})(\text{test})$	344	204	204	259	N/A
0-1 test error	.193	.109	.109	0.141	0.377



Logistic, the most difficult case

$\beta_1 = 4, \beta_2 = 4, \beta_3 = 4, \beta_4 = -6\sqrt{2}, \beta_{p+1} = 4/3, \text{cov}(X_4, X^T \beta^*) = 0.$

Bayes error: 0.1040.

	Van-SIS	ISIS	Var2-ISIS	LASSO	NSC
$\text{med}(\ \beta - \hat{\beta}\ _1)$	20.6	2.69	3.24	23.2	N/A
$\text{med}(\ \beta - \hat{\beta}\ _2^2)$	9.46	1.36	1.59	9.11	N/A
True Positive	0.00	0.90	0.98	0.00	0.17
Med. model size	16	5	5	102	10
$2Q(\hat{\beta}_0, \hat{\beta})$ (training)	269	188	188	109	N/A
AIC	289	198	199	311	N/A
BIC	337	218	219	714	N/A
$2Q(\hat{\beta}_0, \hat{\beta})$ (test)	361	225	226	276	N/A
0-1 test error	.193	.112	.112	.146	.387



Possion, independent covariates

$\beta_0 = 5, \beta_1 = -0.54, \beta_2 = 0.53, \beta_3 = -0.50, \beta_4 = -0.49, \beta_5 = -0.41,$
 $\beta_6 = 0.52, \quad n = 200, N_{\text{sim}} = 100.$

	SIS	ISIS	Var2-ISIS	LASSO
$\text{med}(\ \beta - \hat{\beta}\ _1)$.070	.124	.122	.197
$\text{med}(\ \beta - \hat{\beta}\ _2^2)$.023	.032	.033	.054
True Positive	.76	1.00	1.00	1.00
Med. model size	12	18	17	27
$2Q(\hat{\beta}_0, \hat{\beta})(\text{training})$	1561	1502	1510	1534
AIC	1586	1538	1542	1587
BIC	1627	1597	1595	1674
$2Q(\hat{\beta}_0, \hat{\beta})(\text{test})$	1558	1594	1589	1645



Poisson Regression, difficult case

$$\beta_0 = 5, \beta_1 = 0.6, \beta_2 = 0.6, \beta_3 = 0.6, \beta_4 = -0.9\sqrt{2}$$

$$\text{cov}(X_4, \mathbf{X}^T \beta^*) = 0.$$

	ISIS	Var2-ISIS	LASSO
med($\ \beta - \hat{\beta}\ _1$)	.271	.225	3.07
med($\ \beta - \hat{\beta}\ _2^2$)	.072	.068	1.29
True positive	1.00	.97	0.00
Median final model size	18	16	174
2Q($\hat{\beta}_0, \hat{\beta}$)(training)	1494	1509	1364
AIC	1531	1541	1718
BIC	1590	1596	2293
2Q($\hat{\beta}_0, \hat{\beta}$)(test)	1629	1615	2213



Poisson Regression, the most difficult case

$\beta_0 = 5, \beta_1 = 0.6, \beta_2 = 0.6, \beta_3 = 0.6, \beta_4 = -0.9\sqrt{2}, \beta_{p+1} = -0.15$
 $\text{cov}(X_4, \mathbf{X}^T \beta^*) = 0.$

	Van-ISIS	Var2-ISIS	LASSO
$\text{med}(\ \beta - \hat{\beta}\ _1)$.254	.232	3.09
$\text{med}(\ \beta - \hat{\beta}\ _2^2)$.068	.068	1.29
True positive	.97	.91	0.00
Median final model size	18	16	174
$2Q(\hat{\beta}_0, \hat{\beta})$ (training)	1500	1516	1367
AIC	1536	1547	1715
BIC	1595	1600	2294
$2Q(\hat{\beta}_0, \hat{\beta})$ (test)	1640	1631	2389



Neuroblastoma Data (MAQC-II)

- 1 251 patients of the German Neuroblastoma Trials NB90-NB2004, diagnosed between 1989 and 2004, aged from 0 to 296 months (median 15 months).
- 2 Neuroblastoma is a common paediatric solid cancer (15%)
- 3 251 customized oligonucleotide microarray with $p = 10,707$.
- 4 focus on “3-year Event Free Survival”, —whether each patient survived 3 years after the diagnosis of neuroblastoma ($n = 239$ w/ 49 “+” and 190 “-”).
- 5 Aims: To study which genes are responsible for neuroblastoma and its risk association.



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Results

Training set and endpoints:

- 1 **“3-y EFS”**: Random $n = 125$ subjects (25 “+” and 100 “-”).
- 2 **“Gender”**: Random 120 males and 50 females. Total: 246.

Testing set: The remainder are used as the testing set.

Object	Method	SIS	ISIS	var2-ISIS	LASSO	NSC	Total
3-y EFS	No. pred.	5	23	12	57	9413	10,707
	Test error	19	22	21	22	24	114
Gender	No. pred.	6	2	2	42	3	10,707
	Test error	4	4	4	5	4	126



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	Test error	4	4	4	5	4	



Multi-category Classification



The ISIS method

Linear classifier: $\operatorname{argmax}_k f_k(\mathbf{x})$, where $f_k(\mathbf{x}) \equiv \beta_{0k} + \mathbf{x}^T \beta_k$.

Loss: $L(Y, \mathbf{f}(\mathbf{x}; \mathbf{B})) = \sum_{j \neq Y} [1 + f_j(\mathbf{x})]_+$

Marginal utility of the j -feature (Lee et al, 2004; Liu, et al, 2007):

$$L_j = \min_{\mathbf{B}} \sum_{i=1}^n L(Y_i, \mathbf{f}(X_{ij}, \mathbf{B})) + \frac{1}{2} \sum_k \beta_{jk}^2 \text{ (identifiability)}$$



Simulation Experiments

Design: $\tilde{X}_1, \dots, \tilde{X}_4 \sim U[-\sqrt{3}, \sqrt{3}]$, and $\tilde{X}_5, \dots, \tilde{X}_p \sim N(0, 1)$.

Case 1: $X_j = \tilde{X}_j$ for $j = 1, \dots, p$

Case 2: $X_1 = \tilde{X}_1 - \sqrt{2}\tilde{X}_5$, $X_2 = \tilde{X}_2 + \sqrt{2}\tilde{X}_5$, $X_3 = \tilde{X}_3 - \sqrt{2}\tilde{X}_5$,
 $X_4 = \tilde{X}_4 + \sqrt{2}\tilde{X}_5$,
 $X_j = \sqrt{3}\tilde{X}_j$ for $j = 5, \dots, p$.

Response: 4 categories $\blacksquare P(Y = k | \tilde{\mathbf{X}} = \tilde{\mathbf{x}}) \propto \exp\{f_k(\tilde{\mathbf{x}})\}$,

$f_1(\tilde{\mathbf{x}}) = -a\tilde{x}_1 + a\tilde{x}_4$, $f_2(\tilde{\mathbf{x}}) = a\tilde{x}_1 - a\tilde{x}_2$,

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Simulation results, $n = 400$

	SIS	ISIS	Var2-ISIS	LASSO	NSC
	Case 1				
True positive	1.00	1.00	1.00	0.00	0.68
Median modal size	2.5	4	5	19	4
0-1 test error	0.306	.301	.292	.330	.452
Standard error	.007	.006	.006	.008	.021
	Case 2				
True positive	.10	1.00	1.00	.33	.30
Median modal size	4	11	9	54	9
0-1 test error	.436	.304	.298	.430	.624
Standard error	.007	.007	.006	.004	.008



Test errors: based on $200n$ cases.

Children Cancer Data

Classification: ★neuroblastoma (NB),
★rhabdomyosarcoma (RMS), ★non-Hodgkin lymphoma (NHL),
★Ewing family of tumors (EWS).

Data: cDNA microarrays with 2308 genes (from 6567).

- Training: 63 (12 NBs, 20 RMSs, 8 NHLs, and 23 EWS)
- Testing: 20 (6 NBs, 5 RMSs, 3 NHLs, and 6 EWS)

Results: All methods have zero testing errors.

Method	ISIS	var2-ISIS	LASSO	NSC
# selected genes	15	14	71	343



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- 1 Propose large scale-screening and moderate-selection
 - ▶ Use conditional independence screening.
 - ▶ Allow variable deletion in the process.
 - ▶ Estimation accuracy, comp expediency, algorithmic stability.
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Happy Birthday!

