
Topic 5. Model Selection and Regularization

Sparse Regression

- Given training set $(\mathbf{x}_i, y_i)_{i=1}^n$, with $y_i \in R$ and $\mathbf{x}_i \in R^p$, it is assumed that

$$y_i = \beta_0 + \sum_{j=1}^{p_0} \beta_j x_{ij} + \epsilon_i$$

where $p_0 \ll p$ (**sparsity**).

- $A^* = \{1, \dots, p_0\}$ indexes the informative predictors, and $\{p_0 + 1, \dots, p\}$ indexes the redundant predictors.
- The goal of variable selection is to correctly detect A^* from $\{1, \dots, p\}$.
- We focus on linear regression models, while detecting nonlinear relationship is possible and largely open.

Why Do We Care?

- Multicollinearity: masked significance, inflated variance,...
- Prediction accuracy can be deteriorated due to overfitting when p is large (*curse of dimensionality*).
- Interpretability can be unnecessarily complicated when irrelevant variables are included.

Popular Techniques

- Best subset selection
 - Various information criteria, cross validation
- Sequential variable selection
 - Forward/backward selection
- Shrinkage methods
 - Lasso and its variants
- Dimension reduction
 - Principal component analysis, sufficient dimension reduction

Best Subset Selection

Algorithm 6.1 *Best subset selection*

1. Let \mathcal{M}_0 denote the *null model*, which contains no predictors. This model simply predicts the sample mean for each observation.
 2. For $k = 1, 2, \dots, p$:
 - (a) Fit all $\binom{p}{k}$ models that contain exactly k predictors.
 - (b) Pick the best among these $\binom{p}{k}$ models, and call it \mathcal{M}_k . Here *best* is defined as having the smallest RSS, or equivalently largest R^2 .
 3. Select a single best model from among $\mathcal{M}_0, \dots, \mathcal{M}_p$ using cross-validated prediction error, C_p (AIC), BIC, or adjusted R^2 .
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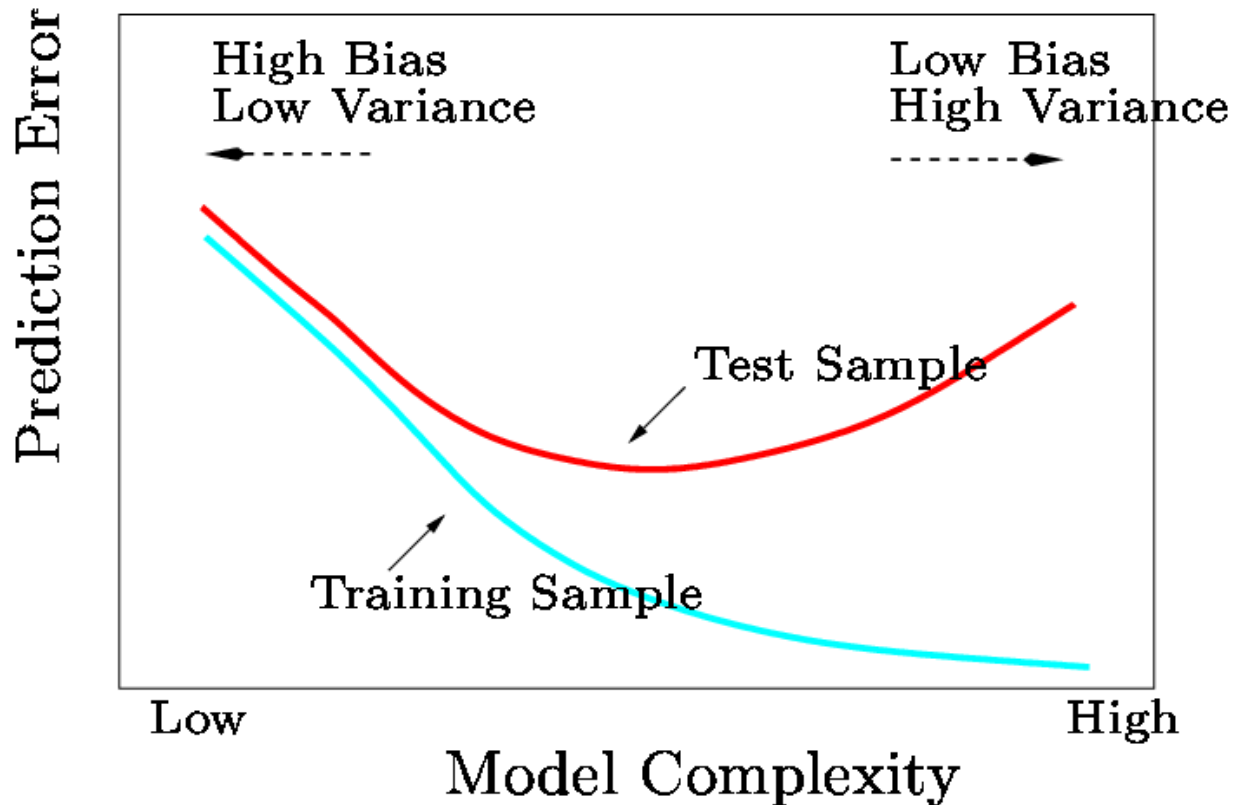
➤ Popular selection criteria

➤ Validation set

➤ Cross validation

➤ “Estimate” test error by making an adjustment to the training error to account for overfitting

Adjustment to Training Error



Model Selection Criteria

➤ For a linear model with d predictors, denote its SSE as SSE_d ,

➤ Mallows's C_p :

$$C_p = \frac{1}{n} (SSE_d + 2d\hat{\sigma}^2)$$

➤ Akaike information criterion (AIC)

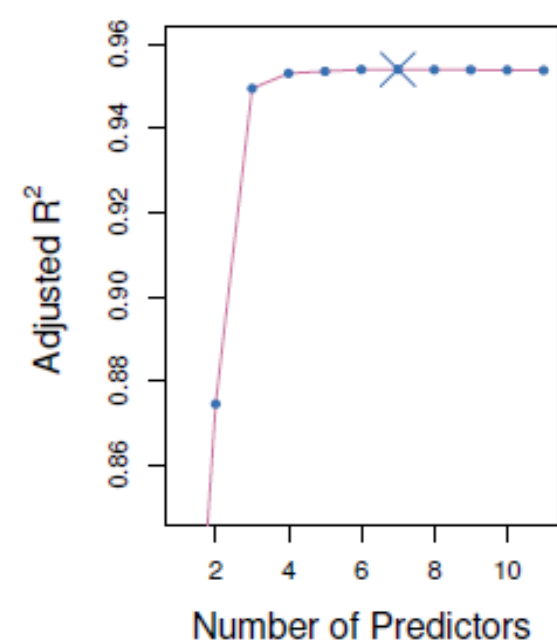
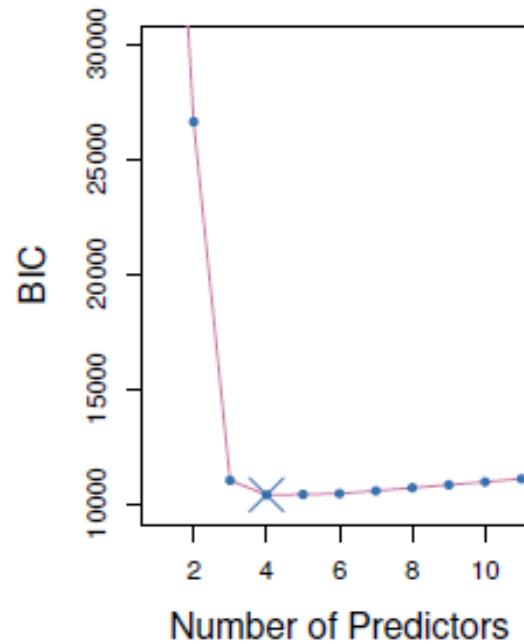
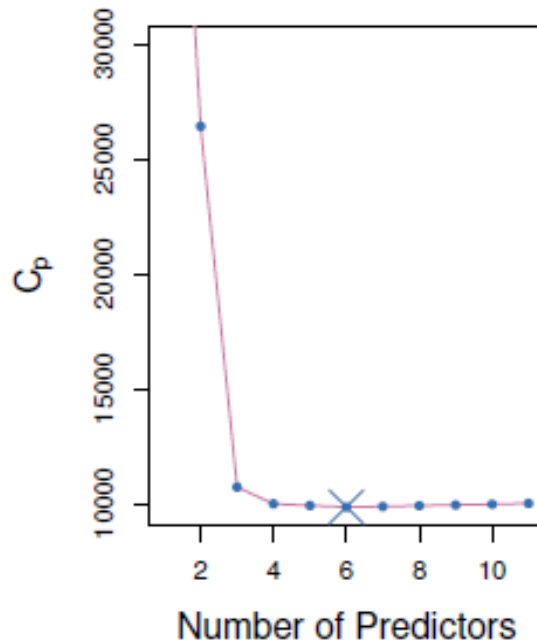
$$AIC = \frac{1}{n\hat{\sigma}^2} (SSE_d + 2d\hat{\sigma}^2)$$

➤ Bayesian information criterion (BIC)

$$BIC = \frac{1}{n\hat{\sigma}^2} (SSE_d + \log(n)d\hat{\sigma}^2)$$

➤ Other criteria: Other IC's, adjusted R^2

An Illustrative Example



Forward/Backward Selection

Algorithm 6.2 *Forward stepwise selection*

1. Let \mathcal{M}_0 denote the *null* model, which contains no predictors.
 2. For $k = 0, \dots, p - 1$:
 - (a) Consider all $p - k$ models that augment the predictors in \mathcal{M}_k with one additional predictor.
 - (b) Choose the *best* among these $p - k$ models, and call it \mathcal{M}_{k+1} . Here *best* is defined as having smallest RSS or highest R^2 .
 3. Select a single best model from among $\mathcal{M}_0, \dots, \mathcal{M}_p$ using cross-validated prediction error, C_p (AIC), BIC, or adjusted R^2 .
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- Backward selection starts with \mathcal{M}_p and iteratively deletes predictors until the best model is found.
- Stagewise selection mixes forward addition and backward deletion in each iteration.

Some Remarks

- Forward/backward selection is computationally more efficient than subset selection.
- It has no guarantee of the best possible model.
- It usually performs well in practice.
- Forward versus backward selection

Shrinkage Methods

- Shrinkage methods are formulated as

$$(\hat{\beta}_0, \hat{\beta}) = \underset{\beta}{\operatorname{argmin}} \sum_{i=1}^n (y_i - \beta_0 - \mathbf{x}_i^T \beta)^2 + \lambda J(\beta)$$

- Various choices of $J(\beta)$ lead to different shrinkage methods and possess different properties.
- After centralization, it becomes

$$\hat{\beta} = \underset{\beta}{\operatorname{argmin}} \sum_{i=1}^n (y_i - \mathbf{x}_i^T \beta)^2 + \lambda J(\beta)$$

Ridge Regression

- Ridge regression uses an L_2 -norm penalty, $\|\beta\|_2^2 = \sum_{j=1}^p \beta_j^2 = \beta^T \beta$,

$$\hat{\beta}_\lambda^{ridge} = \underset{\beta}{\operatorname{argmin}} (\mathbf{y} - \mathbf{X}\beta)^T (\mathbf{y} - \mathbf{X}\beta) + \lambda \|\beta\|_2^2$$

- The second term $\lambda \|\beta\|_2^2$ is a shrinkage penalty, which shrinks the estimates of β towards zero.
- The tuning parameter $\lambda > 0$ controls the trade-off between regression fitting and coefficient shrinkage.
- If $\lambda = 0$, ridge regression produces LSE; if $\lambda \rightarrow \infty$, the estimates of β will approach zero.

Ridge Regression (Cont.)

- Solution of the ridge regression is

$$\hat{\beta}_{\lambda}^{ridge} = (\mathbf{X}^T \mathbf{X} + \lambda I_p)^{-1} \mathbf{X}^T \mathbf{y}$$

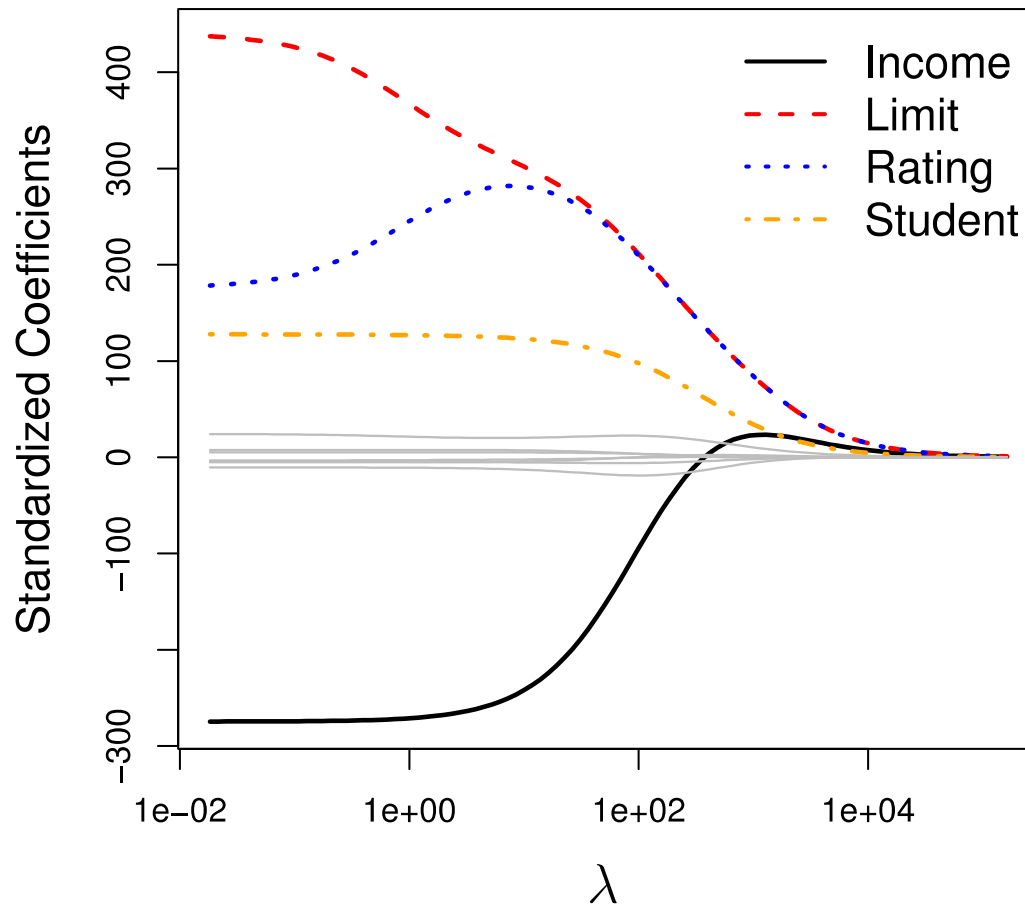
- An equivalent formulation

$$\hat{\beta}_{\lambda}^{ridge} = \underset{\beta}{\operatorname{argmin}} (\mathbf{y} - \mathbf{X}\beta)^T (\mathbf{y} - \mathbf{X}\beta)$$

$$\text{subject to } \|\beta\|^2 \leq s$$

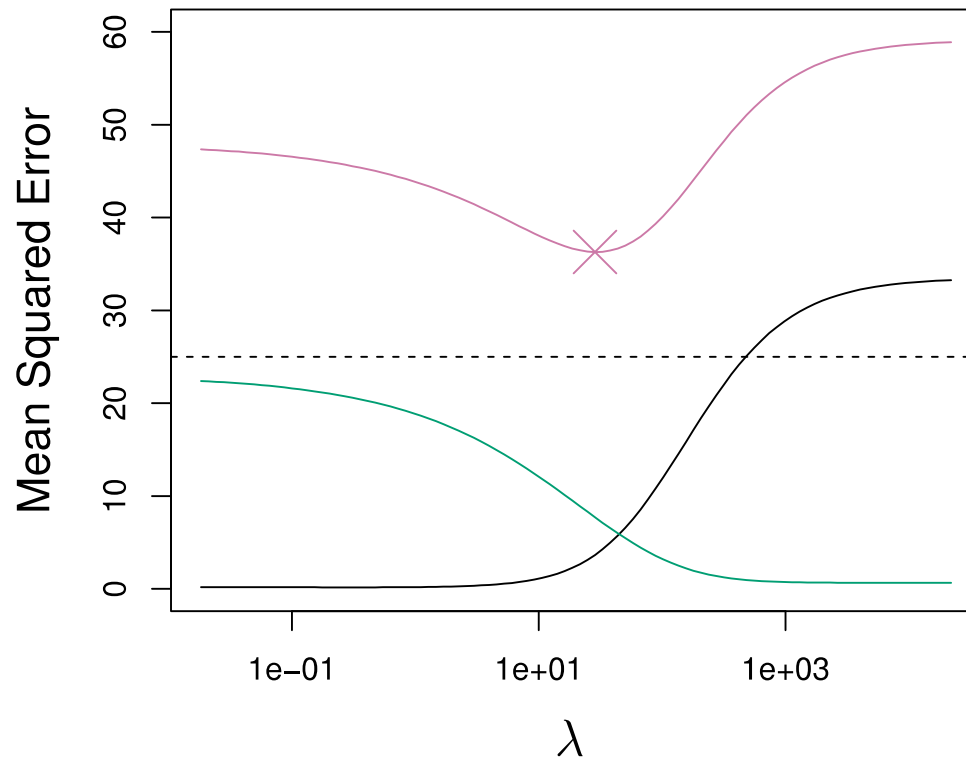
Example

- In general, $\hat{\beta}_\lambda$ is a biased estimator that may have smaller MSE than the LSE estimator.



An Example (Cont.)

- Black: Bias
- Green: Variance
- Purple: Test MSE
- Increase in λ increases bias but decreases variance



LASSO

- The lasso uses an L_1 -norm penalty, $\|\beta\|_1 = \sum_{j=1}^p |\beta_j|$,

$$\hat{\beta}_{\lambda}^{lasso} = \underset{\beta}{\operatorname{argmin}} (\mathbf{y} - \mathbf{X}\beta)^T (\mathbf{y} - \mathbf{X}\beta) + \lambda \|\beta\|_1$$

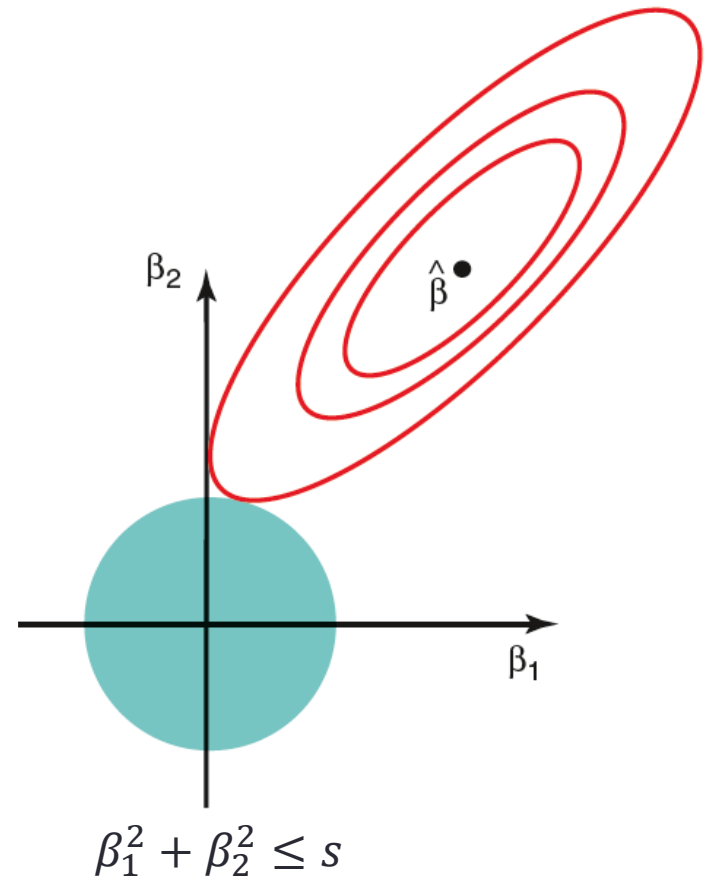
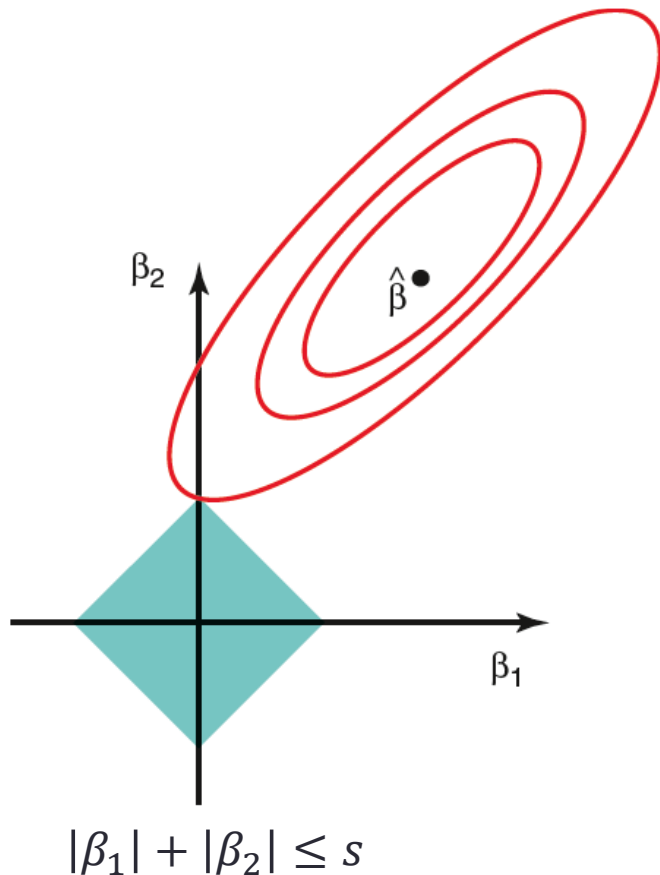
- Or equivalently,

$$\begin{aligned} \hat{\beta}_{\lambda}^{lasso} = \underset{\beta}{\operatorname{argmin}} (\mathbf{y} - \mathbf{X}\beta)^T (\mathbf{y} - \mathbf{X}\beta) \\ \text{subject to } \|\beta\|_1 \leq s \end{aligned}$$

- No explicit solution in general, and a quadratic programming (QP) algorithm can be used to solve the optimization problem.

Sparse Solution

- Some coefficients of the lasso solution will become exactly zero, and thus it does some kind of continuous variable selection.



Example: Prostate Cancer

➤ Data

Predictors (columns 1--8) **Clinical
measures**

lcavol
lweight
age
lbph
svi
lcp
gleason
pgg45

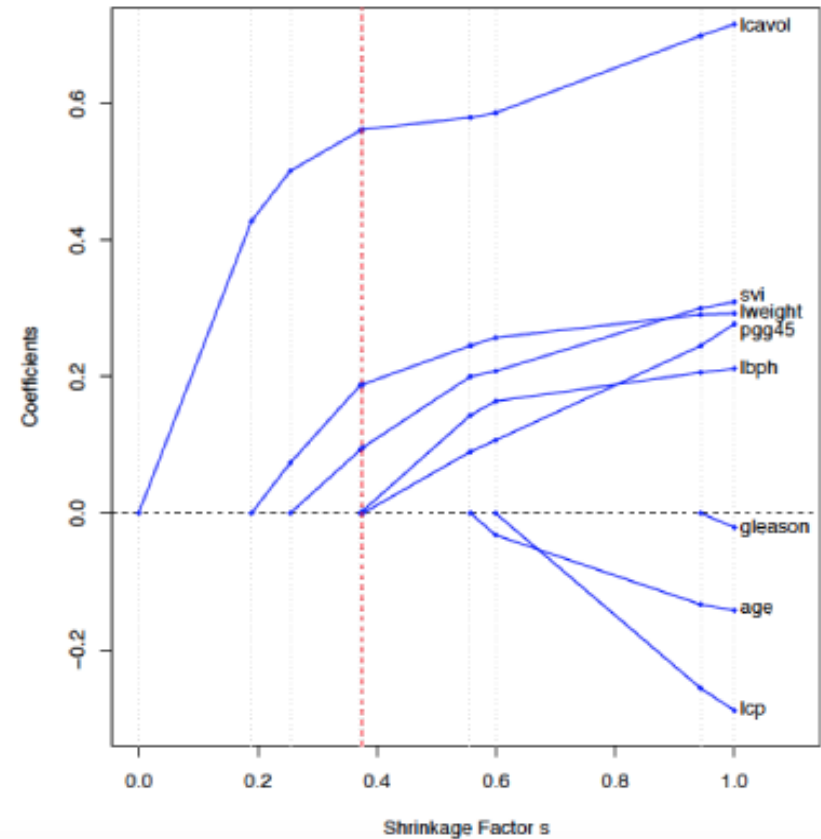
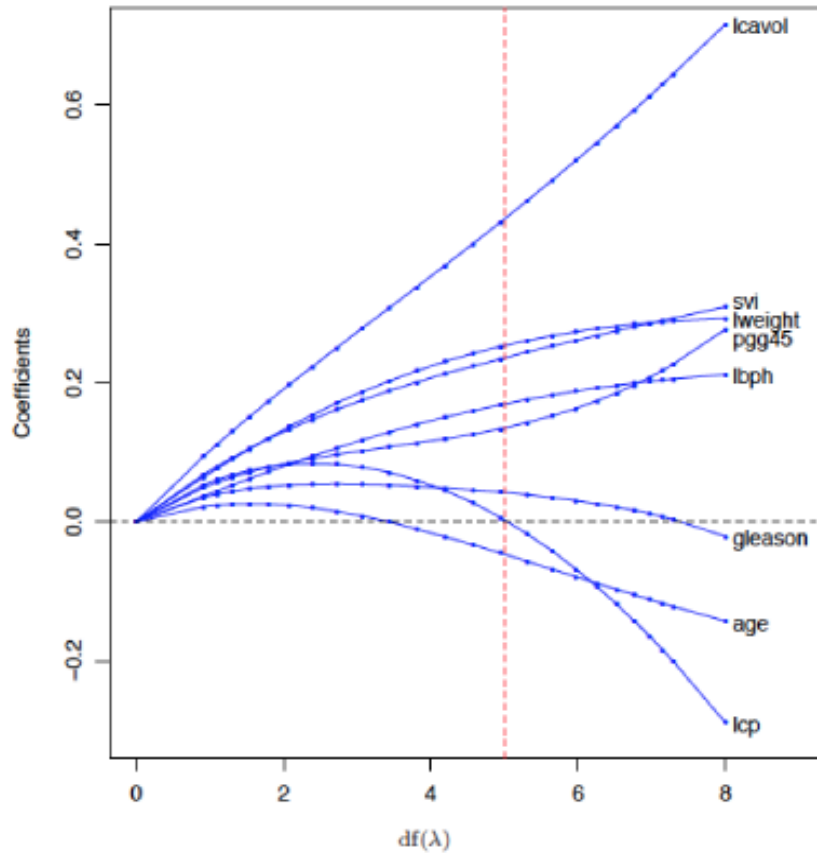
outcome (column 9)

lpsa

**Level of
prostate-
specific
antigen**

Example: Prostate Cancer (Cont.)

➤ Left: ridge regression; Right: lasso

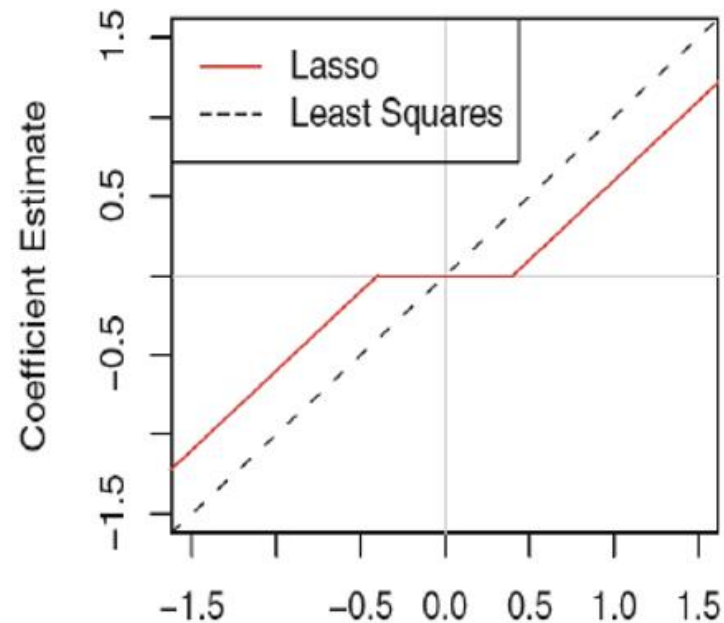
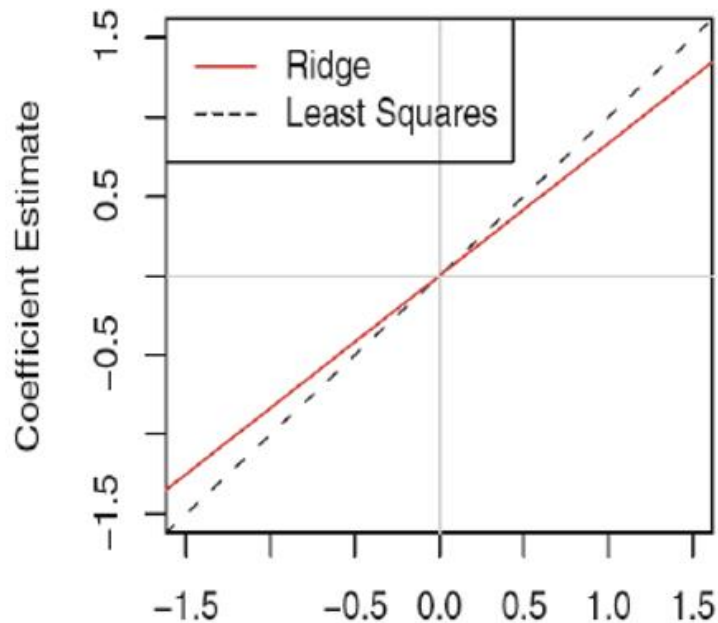


Ridge vs. Lasso

- Both lasso and ridge regression will shrink estimated coefficients while introducing some bias.
- The lasso produces simpler and more interpretable models that involve only a subset of predictors.
- It is unclear which one leads to better prediction accuracy in general though.

A Special Simple Case

- Consider a simple case with $n = p$ and $\mathbf{X} = \mathbf{I}_p$, then $\hat{\beta}_j^{ols} = y_j$,
- Ridge regression multiplies $\hat{\beta}_j^{ols}$ by a constant, $\hat{\beta}_j^{ridge} = y_j / (1 + \lambda)$.
- Lasso truncates $\hat{\beta}_j^{ols}$ towards zero by a constant,
 $\hat{\beta}_j^{lasso} = \text{sign}(y_j)(|y_j| - \lambda/2)_+$.



Bridge Estimators

➤ With $L_r(\beta) = \sum_{j=1}^p |\beta_j|^r$,

$$\hat{\beta}_\lambda^{bridge} = \underset{\beta}{\operatorname{argmin}} \|\mathbf{y} - \mathbf{X}\beta\|^2 + \lambda L_r(\beta)$$

➤ $L_0(\beta) = \sum_{j=1}^p I(\beta_j \neq 0)$ (Hard thresholding)

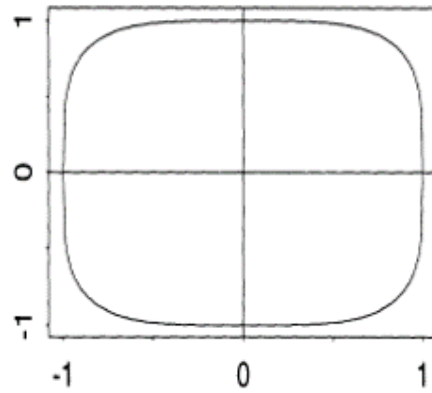
➤ $L_1(\beta) = \sum_{j=1}^p |\beta_j|$ (Lasso)

➤ $L_2(\beta) = \sum_{j=1}^p \beta_j^2$ (Ridge regression)

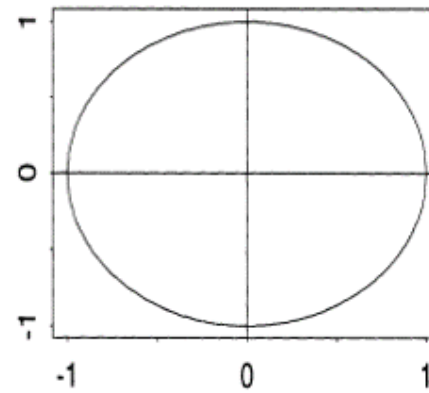
➤ $L_\infty(\beta) = \max_j \beta_j$

Constrained Areas of Bridge Regressions

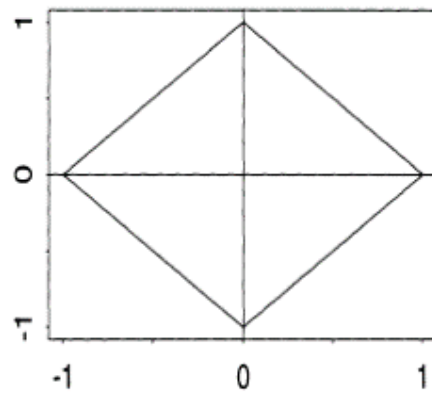
$$r > 2$$



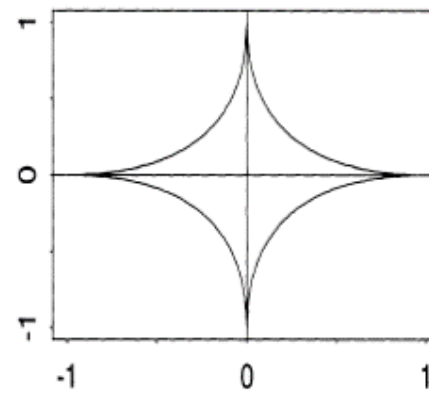
$$r = 2$$



$$r = 1$$



$$r < 1$$



Nonnegative Garrote

$$\min_c \frac{1}{2} \sum_{i=1}^n \left(y_i - \sum_{j=1}^p c_j \hat{\beta}_j x_{ij} \right)^2 + \lambda \sum_{j=1}^p c_j$$

subject to $c_j \geq 0$, and then $\hat{\beta}_j^{ng} = \hat{c}_j \hat{\beta}_j$.

➤ The resulting estimator is

$$\hat{\beta}_j^{ng} = \left(1 - \frac{\lambda}{2\hat{\beta}_j^2} \right)_+ \hat{\beta}_j$$

➤ It is almost unbiased for large $|\hat{\beta}_j|$.

➤ It shrinks small $|\hat{\beta}_j|$ to zero.

Other Extensions

- Group lasso: if the p variables are partitioned into J groups, and then it is desirable to include or exclude the whole group

$$\min_{\beta} \frac{1}{2} \|\mathbf{y} - \mathbf{X}\beta\|^2 + \lambda \sum_{j=1}^p \|\vec{\beta}_j\|_2$$

where $\vec{\beta}_j$ is a coefficient vector for the j th group.

- Elastic net:

$$\min_{\beta} \frac{1}{2} \|\mathbf{y} - \mathbf{X}\beta\|^2 + \lambda_1 \|\beta\|_1 + \lambda_2 \|\beta\|_2^2$$

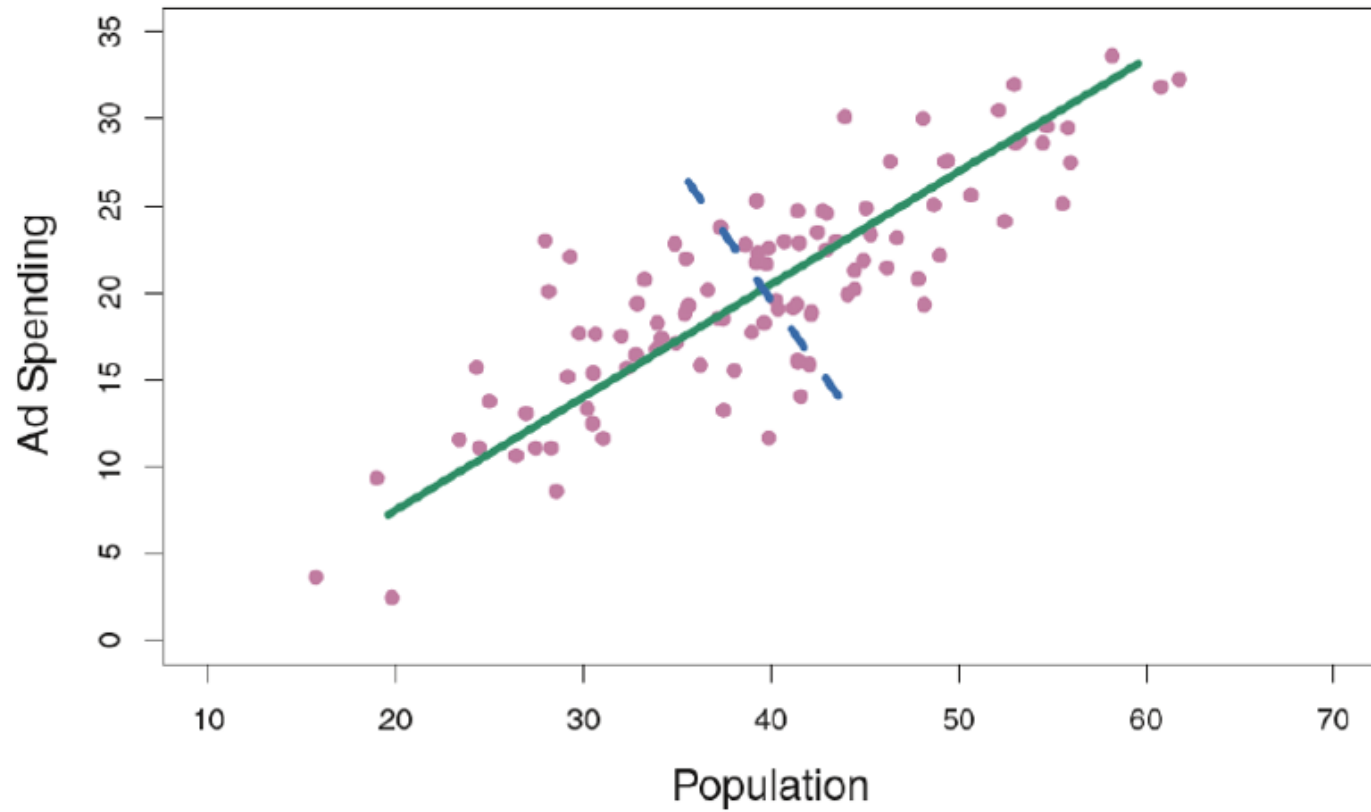
- Fused lasso:

$$\min_{\beta} \frac{1}{2} \|\mathbf{y} - \mathbf{X}\beta\|^2 + \lambda \sum_{j=2}^p \|\beta_j - \beta_{j-1}\|_1$$

Principal Component Analysis (PCA)

- Given $X = (X_1, \dots, X_p)^T$ and $\Sigma = \text{cov}(X)$, find $\{a_1, \dots, a_p\}$ with $\|a_j\| = 1, j = 1, \dots, p$ such that
 - $\text{var}(a_j^T X) = a_j^T \Sigma a_j$ is as large as possible, and
 - $\text{cov}(a_j^T X, a_l^T X) = a_j^T \Sigma a_l = 0$ when $j \neq l$.
- In general,
 - First, find $a_1 = \underset{a}{\operatorname{argmax}} a^T \Sigma a$ subject to $\|a\| = 1$,
 - Then find $a_k = \underset{a}{\operatorname{argmax}} a^T \Sigma a$ subject to $\|a\| = 1$ and $a^T \Sigma a_j = 0$ for $j = 1, \dots, k - 1$

Example



PCA and Eigen-decomposition

- Assume the eigenvalues of Σ is $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_p \geq 0$, and the associated eigenvectors are e_1, \dots, e_p , then
 - $a_j = e_j$ and the j -th PC is $U_j = e_j^T X$
 - $\text{var}(U_j) = e_j^T \Sigma e_j = \lambda_j$
 - $\text{cov}(U_j, U_l) = e_j^T \Sigma e_l = 0$
- To reduce dimension, set $0 < \alpha < 1$ and choose $k \ll p$ such that

$$\frac{\lambda_1 + \dots + \lambda_k}{\lambda_1 + \dots + \lambda_p} \geq \alpha$$

and then work on the feature space spanned by the first k PC's.

Projection Pursuit (PP)

- Imagine an example with $X_1 \sim N(0, 100)$, and $X_2 \sim N(Z, 1/100)$ with $Z = \pm 1 (\pm 1/2)$. PCA will find X_1 as the first PC, but it is less informative.
- The key idea of PP is to find direction which is non-normal.
 - Let $I(\cdot)$ be a measure of non-normality,

$$\hat{a} = \operatorname{argmax}_a I(a^T X)$$

- Popularly used $I(\cdot)$:
 - $I_1(z) = |k_m(z)|/k_2(z)^{m/2}$
 - $I_2(z) = k_3^2(z) + k_4^2(z)/4$
where $k_m(z)$ is the m -th order cumulant of z .

Projection Pursuit (PP)

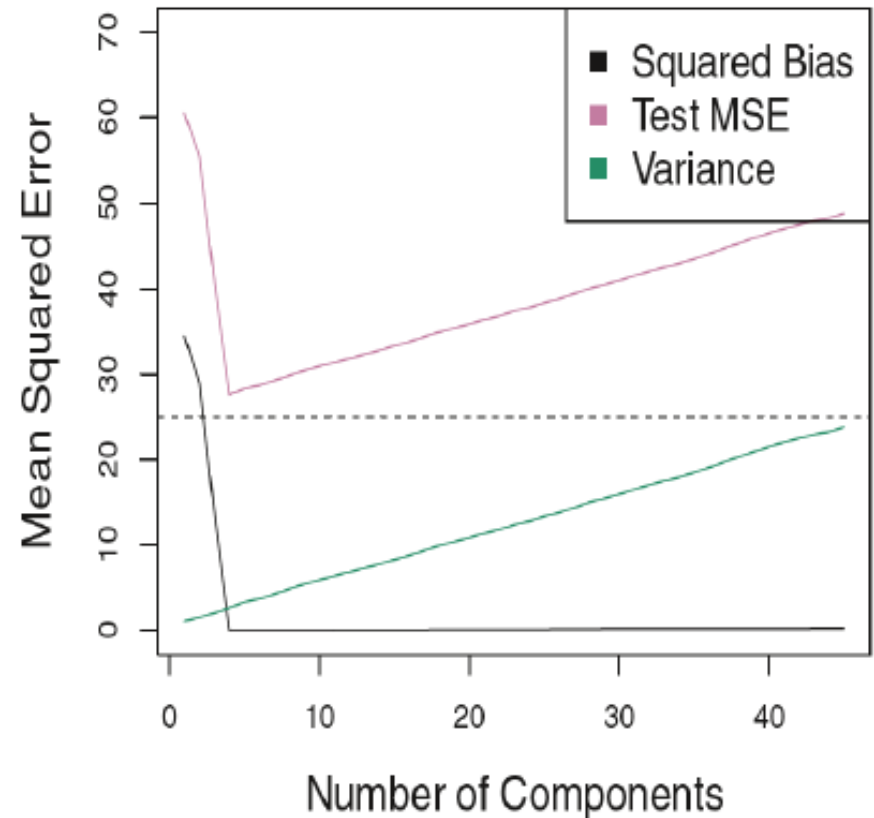
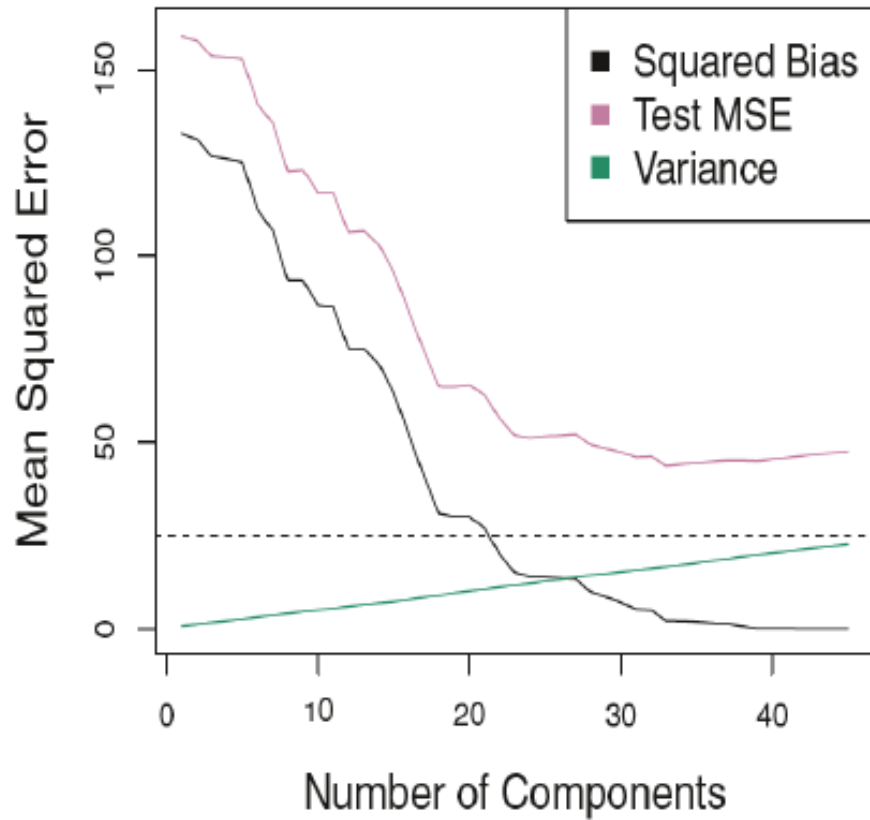
Principal Component Regression (PCR)

- Let S be the sample covariance matrix and \mathbf{q}_j , $j = 1, \dots, J$, be the PC loadings of S .
- PCR computes the derived input columns $\mathbf{z}_j = \mathbf{X}\mathbf{q}_j$ (sample principal components), and then regresses \mathbf{y} on $\mathbf{z}_1, \dots, \mathbf{z}_J$.
- Since the \mathbf{z}_j 's are orthogonal, this regression is just a sum of univariate regressions,

$$\hat{\mathbf{y}}^{pcr} = \bar{\mathbf{y}} + \sum_{j=1}^J \tilde{\gamma}_j \mathbf{z}_j$$

where $\tilde{\gamma}_j$ is the correlation coefficient of \mathbf{y} on \mathbf{z}_j .

Some Simulated Examples with PCR



Some Remarks on PCR

- PCR works well when the first few principal components are sufficient to capture most of the variation in the predictors and the relationship with the response.
- PCR does not produce variable selection, as all predictors are included in each principal component.
- The number of principal components is typically chosen by cross-validation.
- When performing PCR, it is generally recommended to first standardize each predictor.

Partial Least Squares (PLS)

- PLS also constructs a set of linear combinations of predictors for regression, but unlike PCR, it uses \mathbf{y} (and \mathbf{X}) for this construction.
- Assume \mathbf{y} is centered, and we begin by computing the univariate regression coefficient γ_j of Y on X_j .
- From this we construct the derived input $Z_1 = \sum_{j=1}^p \gamma_j \mathbf{x}_j$, which is the first PLS direction.
- Then Y is regressed on Z_1 , giving coefficient $\hat{\beta}_1$. Next we orthogonalize X_1, \dots, X_p with respect to Z_1 : $R_1 = Y - \hat{\beta}_1 Z_1$, and $X_j^* = X_j - \hat{\theta}_j Z_1$, $j = 1, \dots, p$, where $\hat{\theta}_j$ is the coefficient when X_j is regressed on Z_1 . Then find the univariate regression coefficient of R_1 on X_j^* .

Partial Least Squares (Cont.)

- We continue this process, until J directions are obtained.
- In this manner, PLS produces a sequence of derived inputs or directions Z_1, \dots, Z_J .
- As with PCR, if we continue on to construct $J = p$ directions, we get back the OLS estimates; the use of $J < p$ directions produces a reduced-dim regression.
- Notice that in the construction of each Z_j , the inputs are weighted by the strength of their univariate effect on Y .

PCR vs PLS

