Classical Statistics Biological Big Data Supervised and Unsupervised Learning

High-Dimensional Statistical Learning: Introduction

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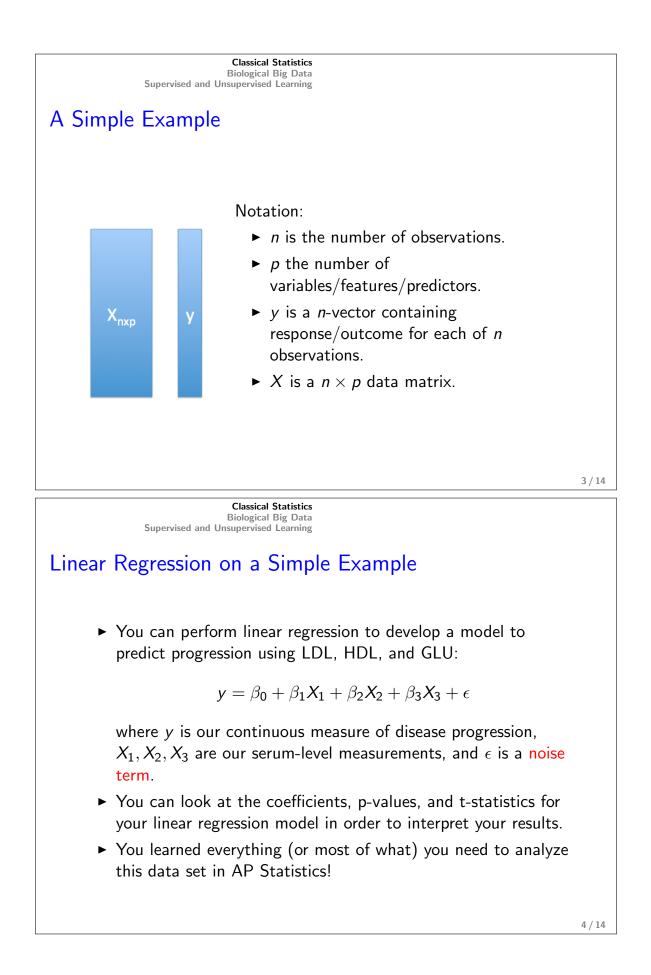
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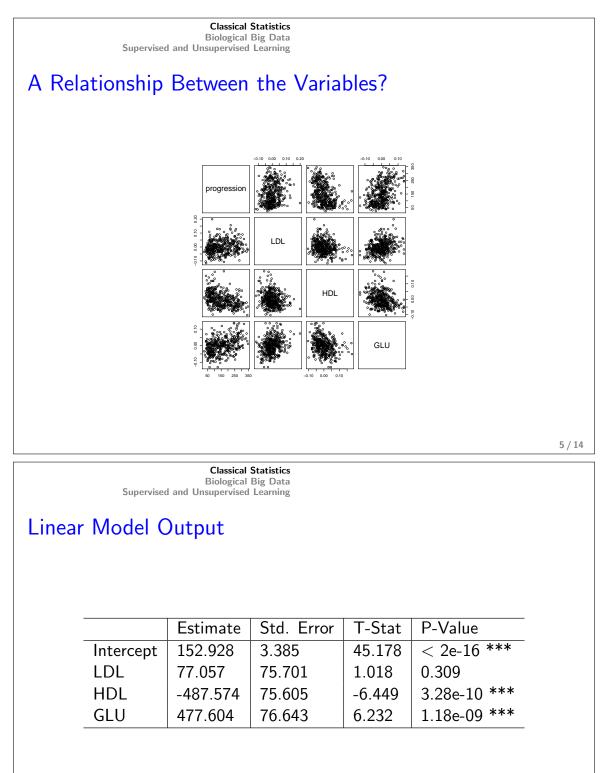
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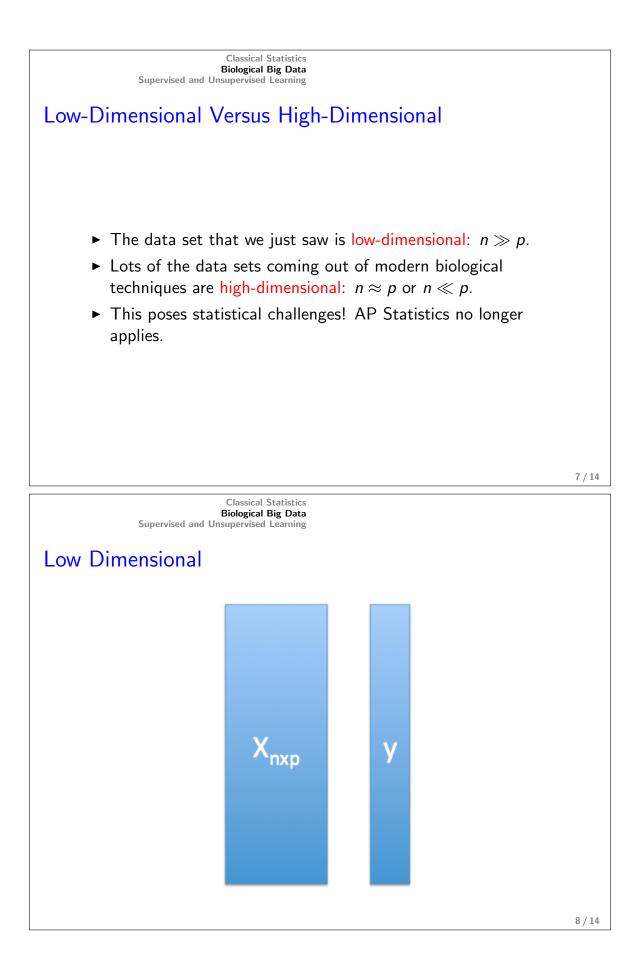
A Simple Example

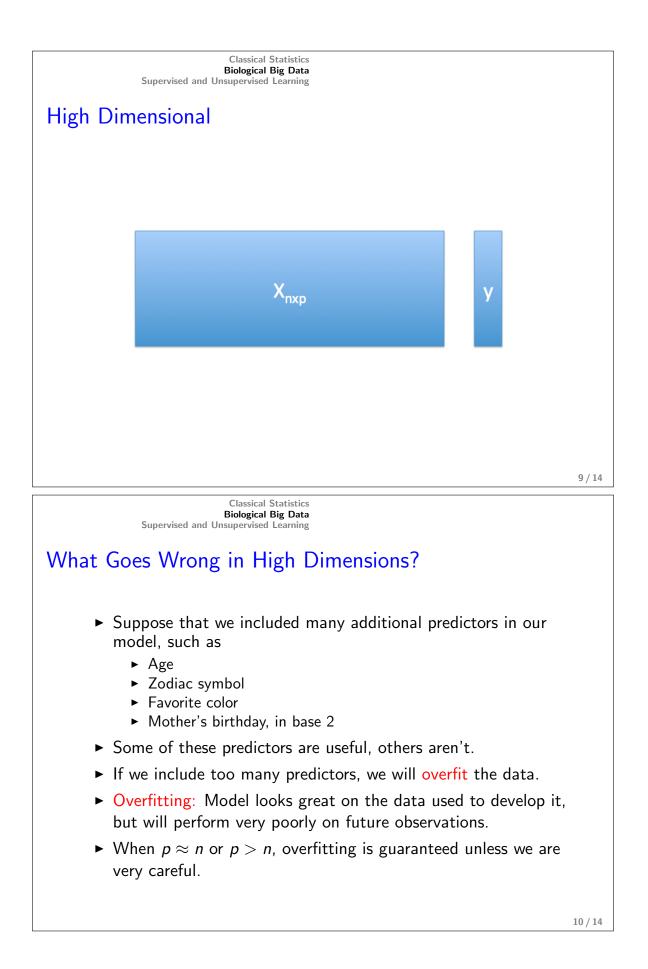
- Suppose we have n = 400 people with diabetes for whom we have p = 3 serum-level measurements (LDL, HDL, GLU).
- We wish to predict these peoples' disease progression after 1 year.

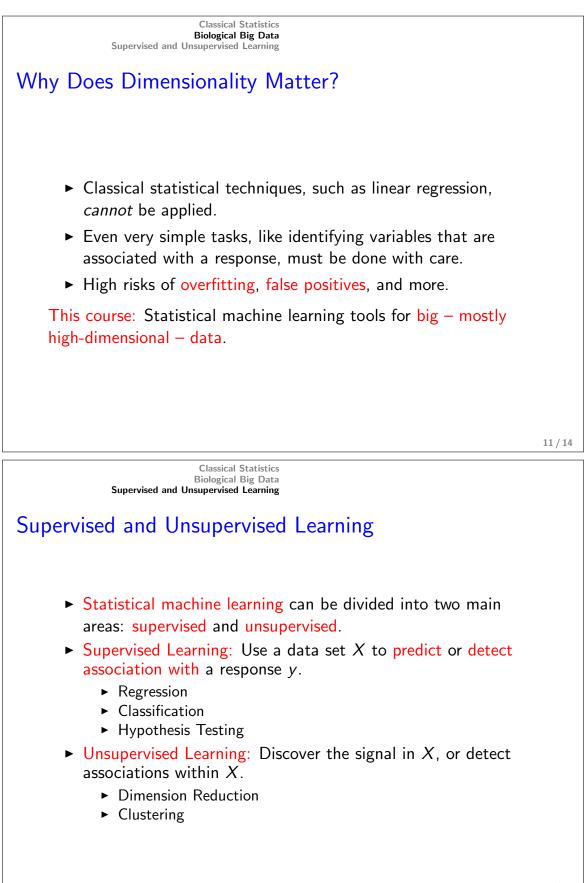


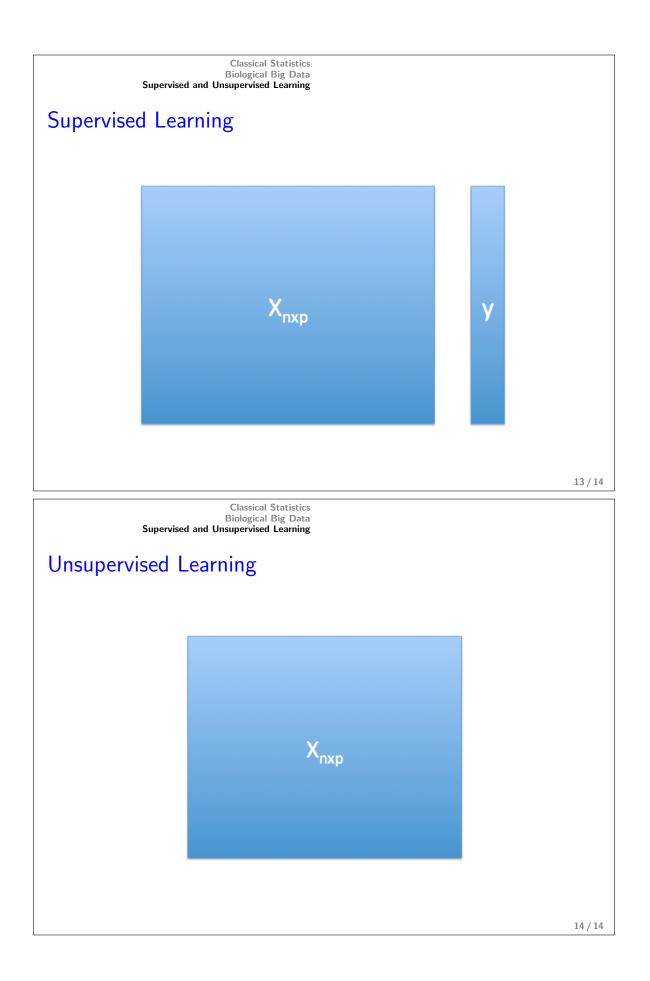


progression_measure \approx 152.9+77.1×LDL-487.6×HDL+477.6×GLU.







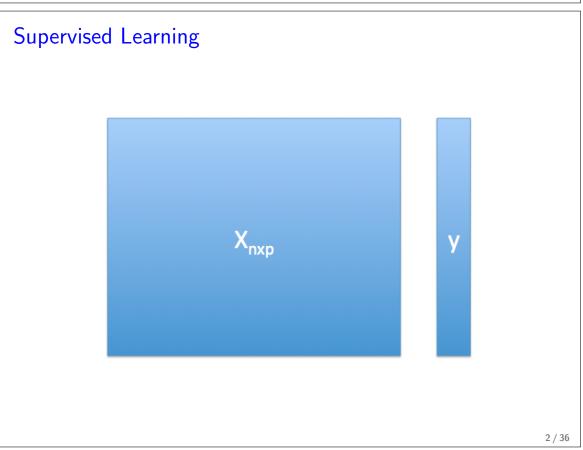


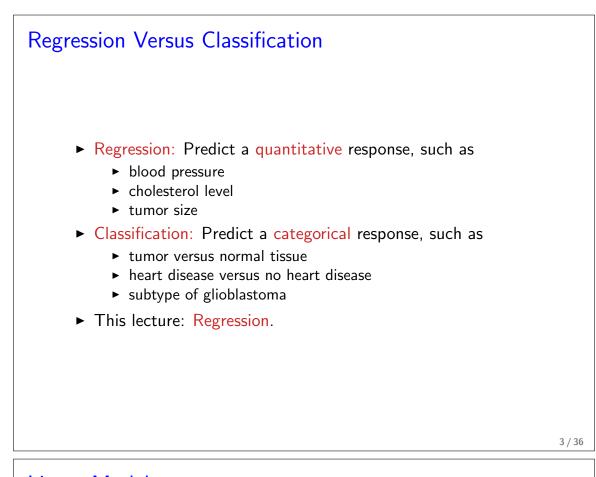
High-Dimensional Statistical Learning: Bias Variance Tradeoff and the Test Error

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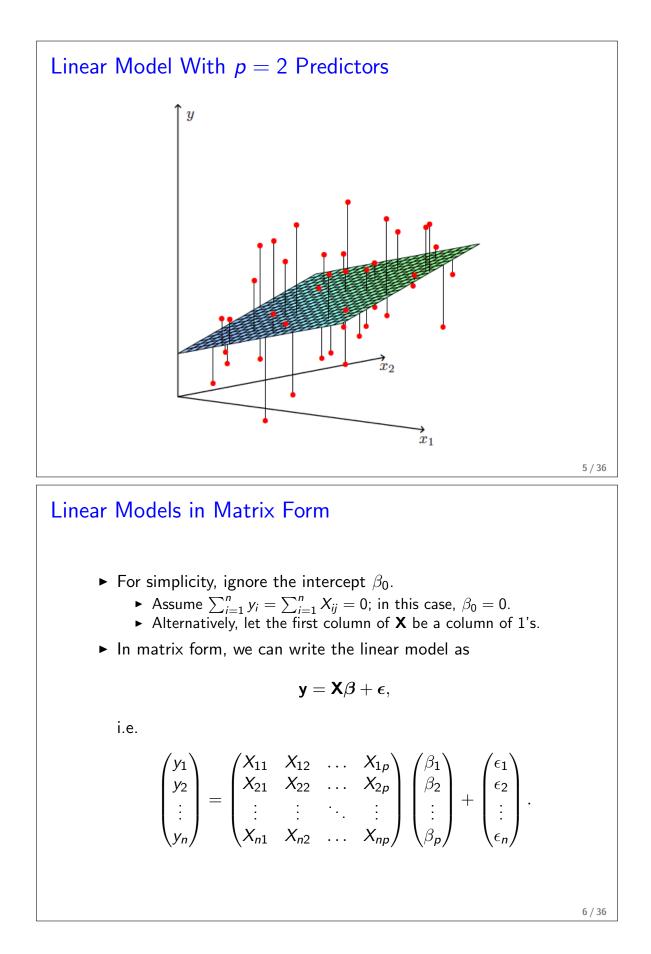


Linear Models

- ► We have *n* observations, for each of which we have *p* predictor measurements and a response measurement.
- ► Want to develop a model of the form

$$y_i = \beta_0 + \beta_1 X_{i1} + \ldots + \beta_p X_{ip} + \epsilon_i.$$

- Here ϵ_i is a noise term associated with the *i*th observation.
- Must estimate $\beta_0, \beta_1, \ldots, \beta_p$ i.e. we must fit the model.



Least Squares Regression

► There are many ways we could fit the model

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon}.$$

Most common approach in classical statistics is least squares:

minimize
$$\{ \| \mathbf{y} - \mathbf{X} \boldsymbol{\beta} \|^2 \}$$

Here $\|\mathbf{a}\|^2 \equiv \sum_{i=1}^n a_i^2$. • We are looking for β_1, \dots, β_p such that

$$\sum_{i=1}^n (y_i - (\beta_1 X_{i1} + \ldots + \beta_p X_{ip}))^2$$

is as small as possible, or in other words, such that

$$\sum_{i=1}^n (y_i - \hat{y}_i)^2$$

is as small as possible, where \hat{y}_i is the *i*th predicted value.

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Least Squares Regression

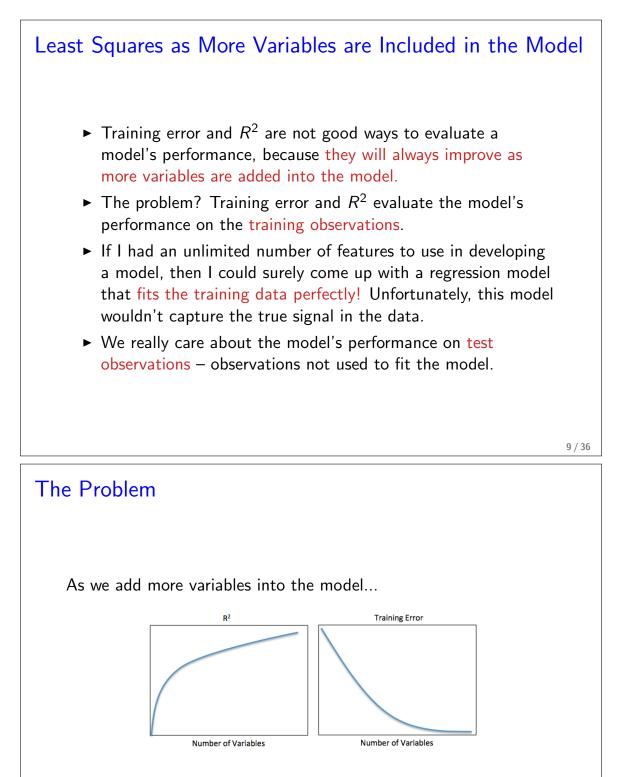
- ► When we fit a model, we use a training set of observations.
- We get coefficient estimates $\hat{\beta}_1, \ldots, \hat{\beta}_p$.
- ► We also get predictions using our model, of the form

$$\hat{y}_i = \hat{\beta}_1 X_{i1} + \ldots + \hat{\beta}_p X_{ip}.$$

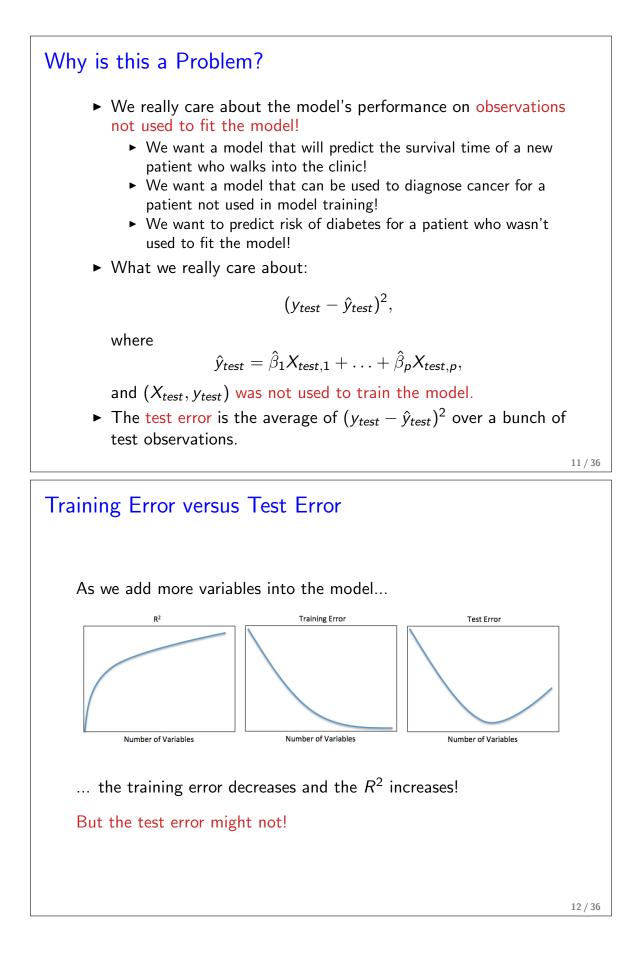
- ▶ We can evaluate the training error, i.e. the extent to which the model fits the observations used to train it.
- One way to quantify the training error is using the mean squared error (MSE):

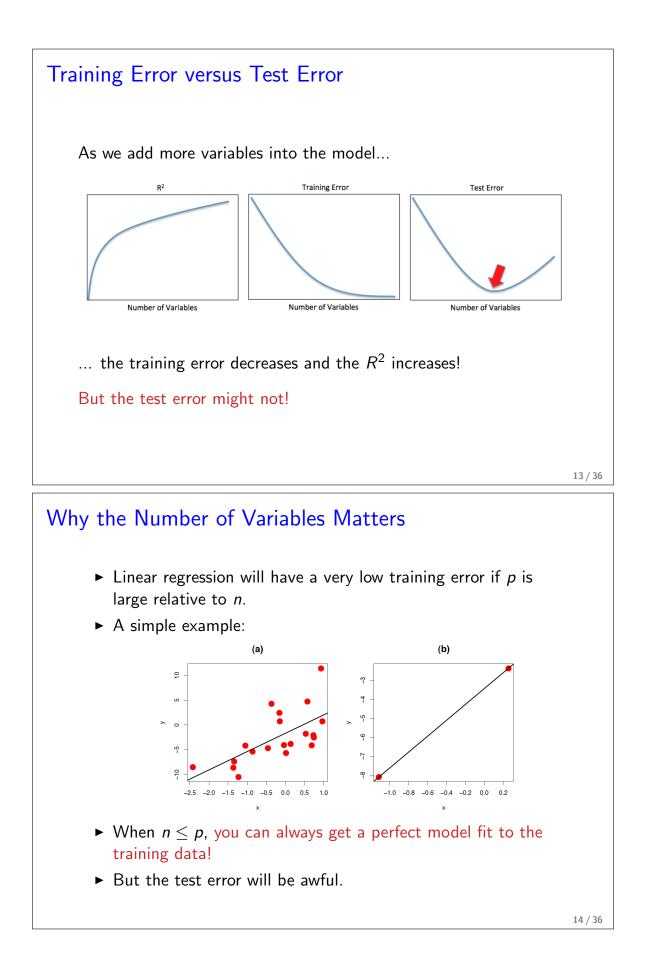
$$MSE = \frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{y}_i)^2 = \frac{1}{n} \sum_{i=1}^{n} (y_i - (\hat{\beta}_1 X_{i1} + \ldots + \hat{\beta}_p X_{ip}))^2.$$

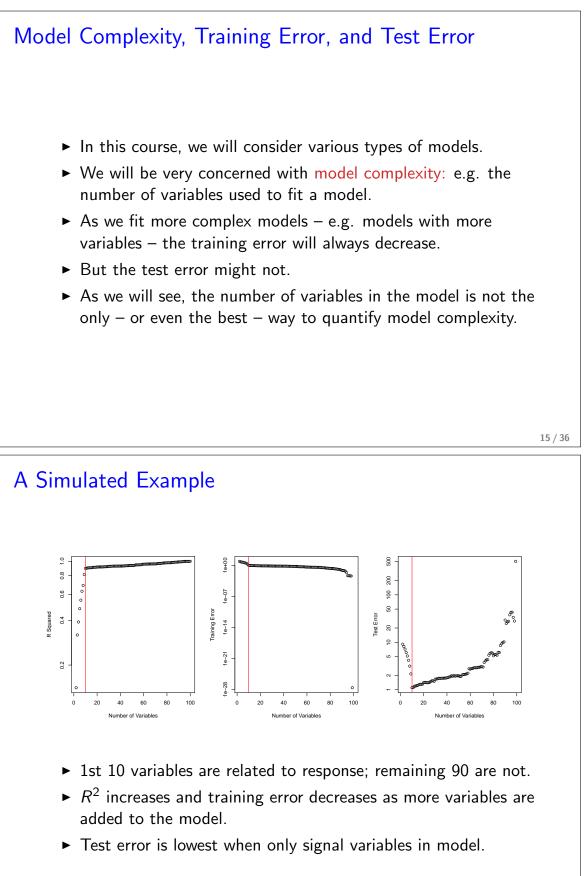
- The training error is closely related to the R^2 for a linear model - that is, the proportion of variance explained.
- Big $R^2 \Leftrightarrow$ Small Training Error.



... the training error decreases and the R^2 increases!







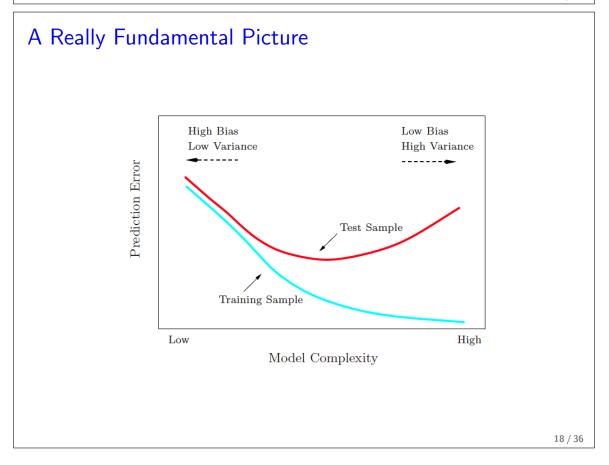
Bias and Variance

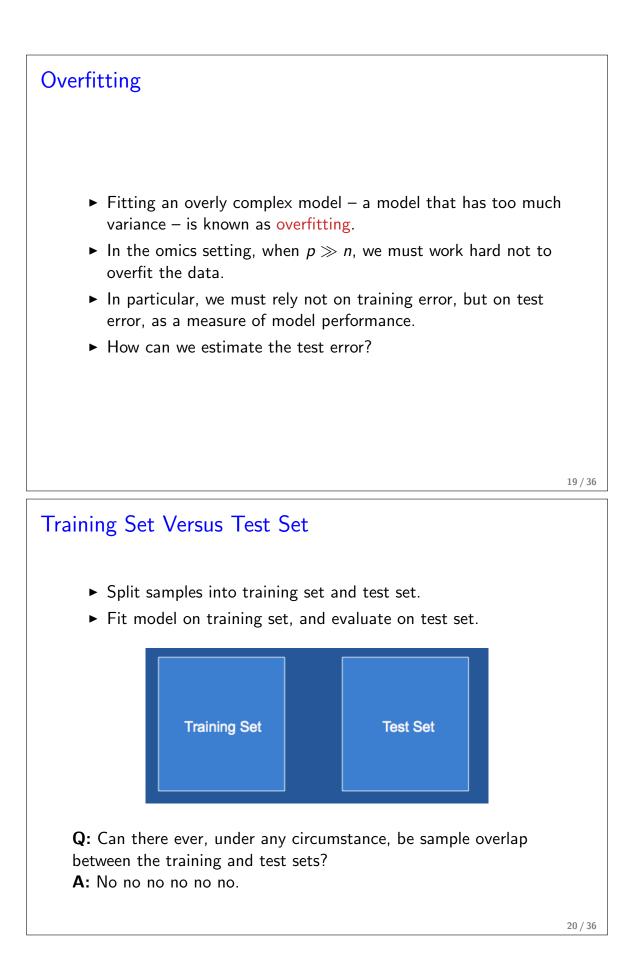
- As model complexity increases, the bias of β̂ the average difference between β and β̂, if we were to repeat the experiment a huge number of times will decrease.
- But as complexity increases, the variance of $\hat{\beta}$ the amount by which the $\hat{\beta}$'s will differ across experiments will increase.
- The test error depends on both the bias and variance:

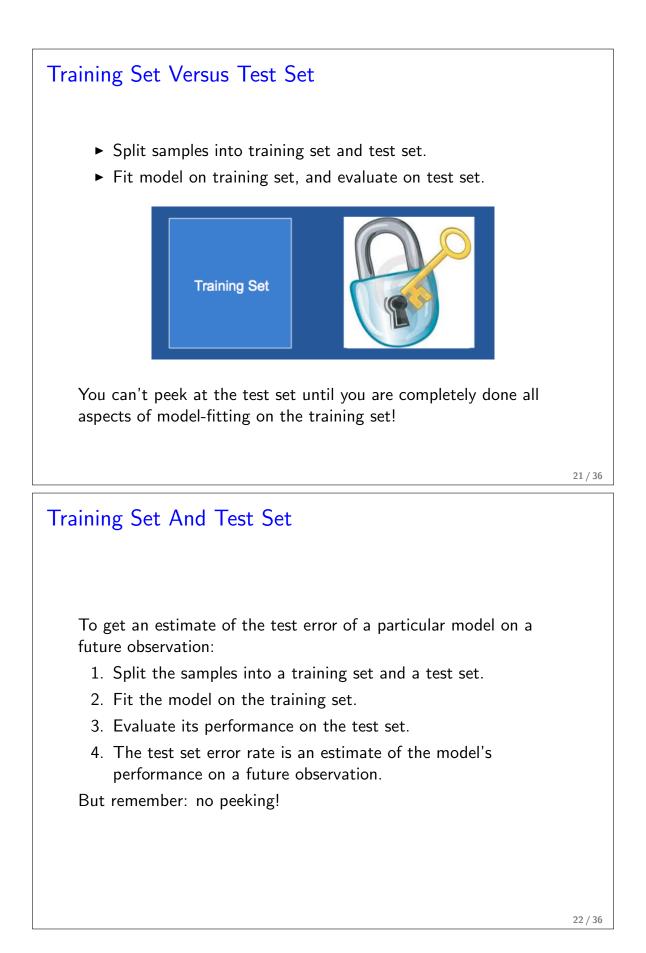
Test $Error = Bias^2 + Variance$.

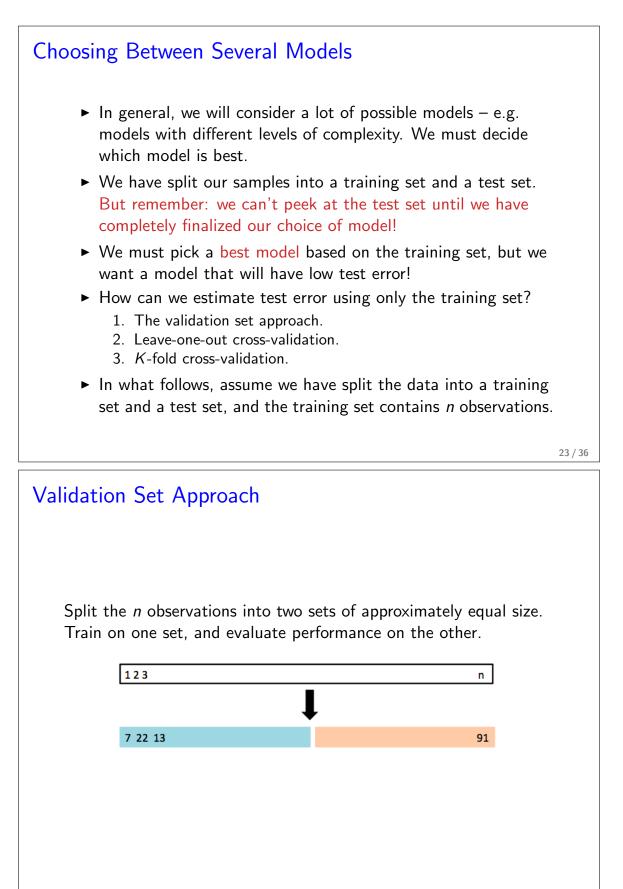
There is a bias-variance trade-off. We want a model that is sufficiently complex as to have not too much bias, but not so complex that it has too much variance.

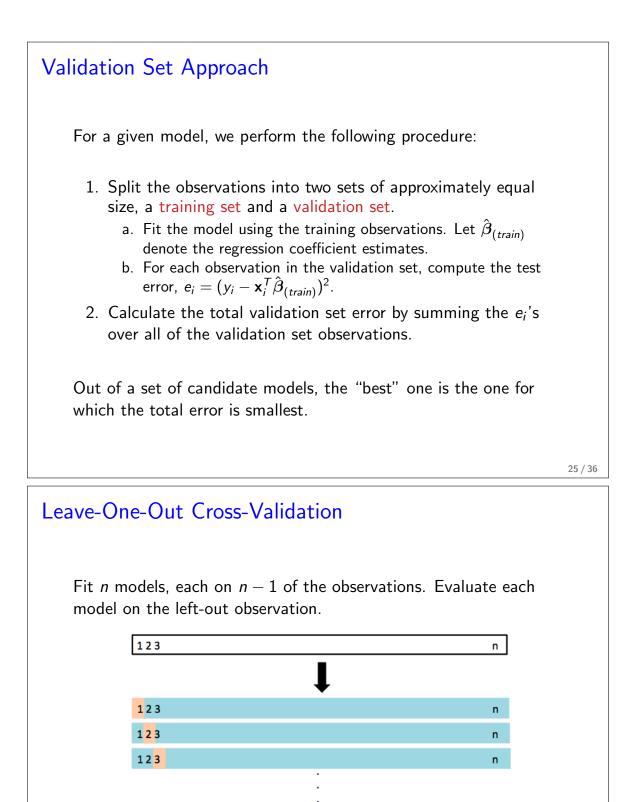












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Leave-One-Out Cross-Validation

For a given model, we perform the following procedure:

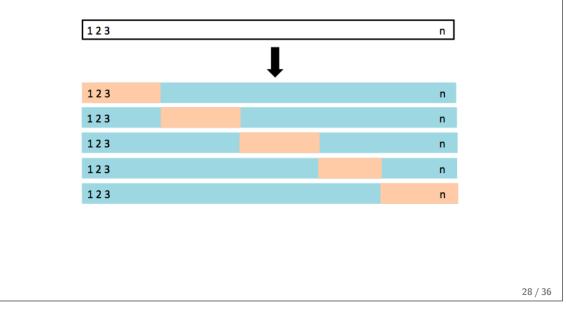
- 1. For i = 1, ..., n:
 - a. Fit the model using observations $1, \ldots, i 1, i + 1, \ldots, n$. Let $\hat{\beta}_{(i)}$ denote the regression coefficient estimates.
 - b. Compute the test error, $e_i = (y_i \mathbf{x}_i^T \hat{\boldsymbol{\beta}}_{(i)})^2$.
- 2. Calculate $\sum_{i=1}^{n} e_i$, the total CV error.

Out of a set of candidate models, the "best" one is the one for which the total error is smallest.

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5-Fold Cross-Validation

Split the observations into 5 sets. Repeatedly train the model on 4 sets and evaluate its performance on the 5th.



K-fold cross-validation

A generalization of leave-one-out cross-validation. For a given model, we perform the following procedure:

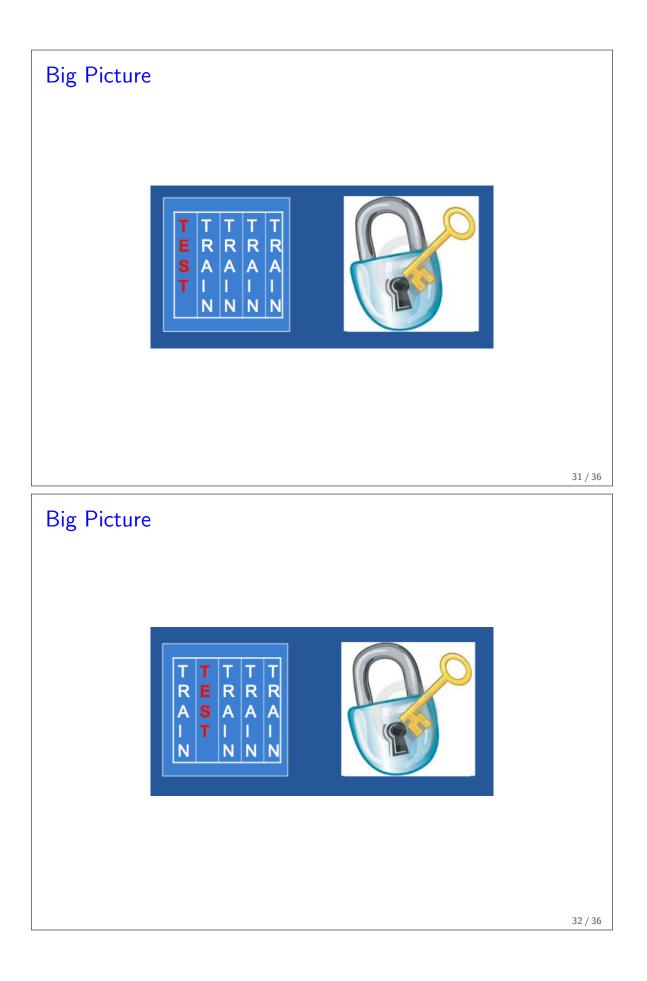
- 1. Split the n observations into K equally-sized folds.
- 2. For k = 1, ..., K:
 - a. Fit the model using the observations not in the kth fold.
 - b. Let e_k denote the test error for the observations in the *k*th fold.
- 3. Calculate $\sum_{k=1}^{K} e_k$, the total CV error.

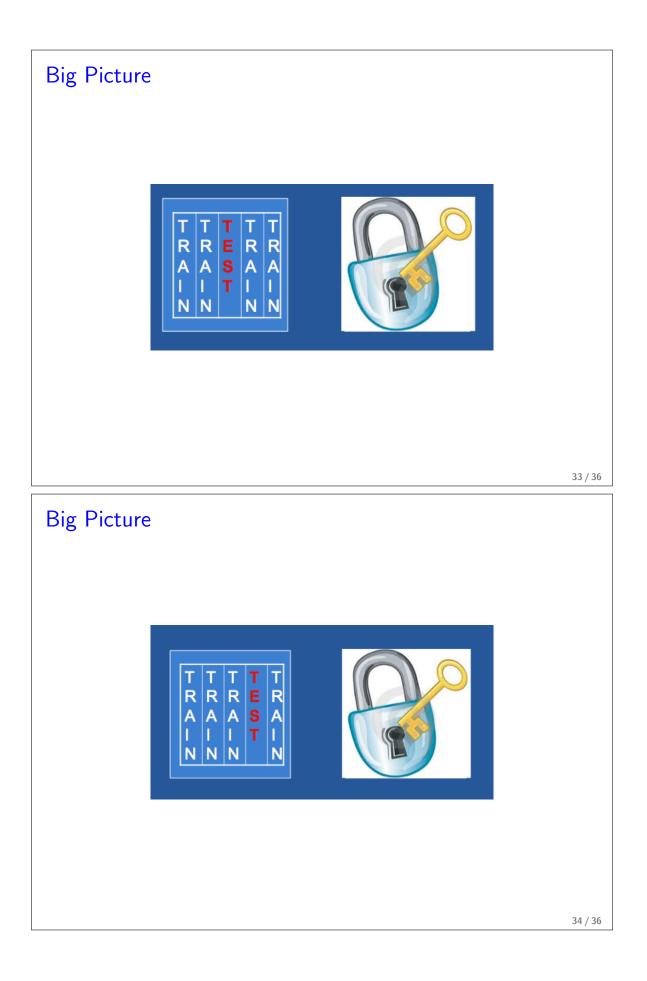
Out of a set of candidate models, the "best" one is the one for which the total error is smallest.

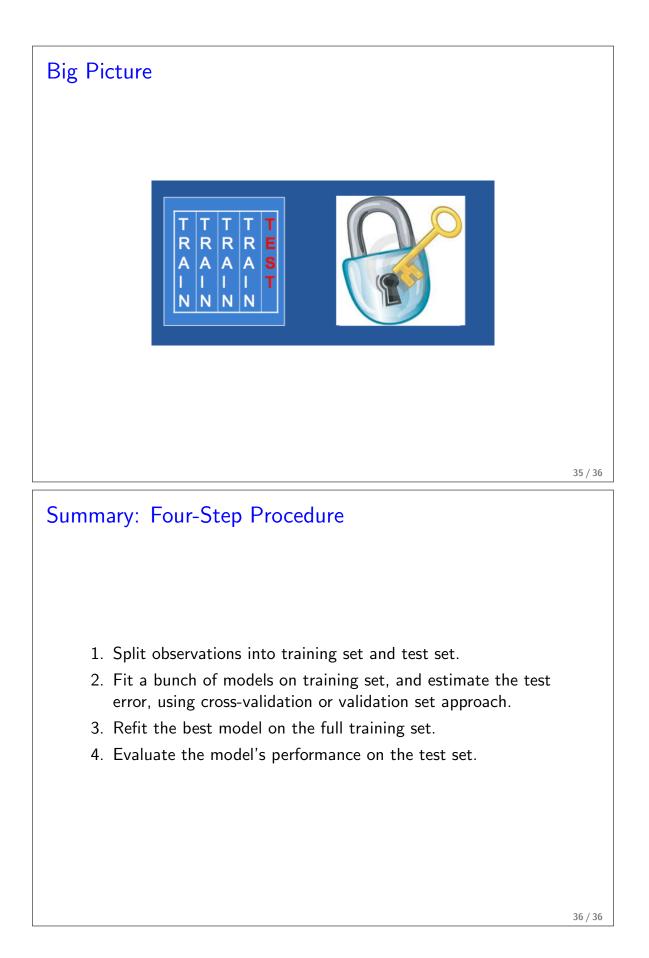
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After Estimating the Test Error on the Training Set...

After we estimate the test error using the training set, we refit the "best" model on all of the available training observations. We then evaluate this model on the test set.





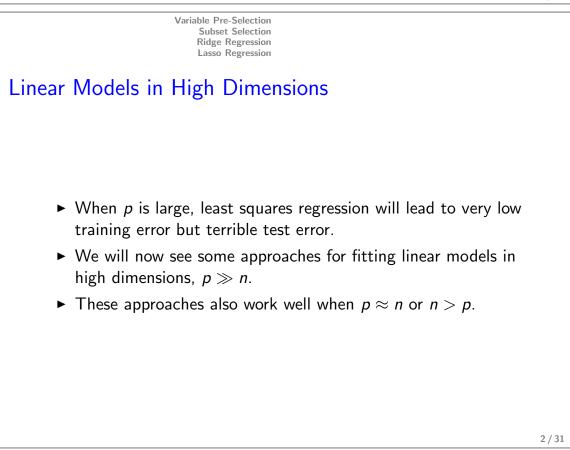


Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression

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Variable Pre-Selection	
Subset Selection	
Ridge Regression Lasso Regression	
Motivating example	
U	
 We would like to build a model to predict survival time for breast cancer patients using a number of clinical measurements (tumor stage, tumor grade, tumor size, patient age, etc.) as well as some biomarkers. For instance, these biomarkers could be: the expression levels of genes measured using a microarray. protein levels. mutations in genes potentially implicated in breast cancer. How can we develop a model with low test error in this setting? 	
setting?	
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Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression	
Remember	
We have <i>n</i> training observations.	
Our goal is to get a model that will perform well on future	
 Our goal is to get a model that will perform well on future test observations. 	
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Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression	
Variable Pre-Selection	
 The simplest approach for fitting a model in high dimensions: 1. Choose a small set of variables, say the q variables that are most correlated with the response, where q < n and q < p. 2. Use least squares to fit a model predicting y using only these q variables. This approach is simple and straightforward. 	
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Variable Pre-Selection	
Subset Selection Ridge Regression Lasso Regression	
How Many Variable to Use?	
We need a way to choose q, the number of variables used in the regression model.	
We want q that minimizes the test error.	
For a range of values of q, we can perform the validation set approach, leave-one-out cross-validation, or K-fold cross-validation in order to estimate the test error.	
 Then choose the value of q for which the estimated test error is smallest. 	

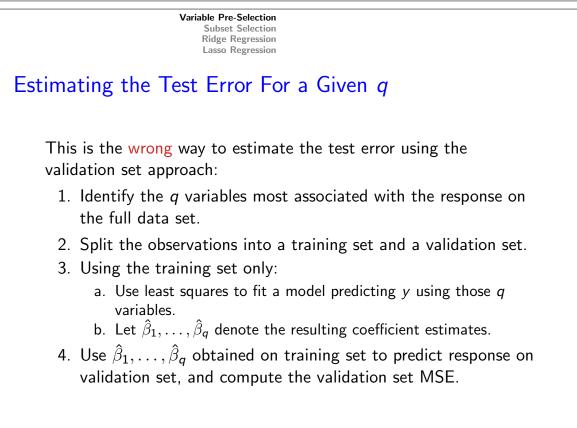
Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression

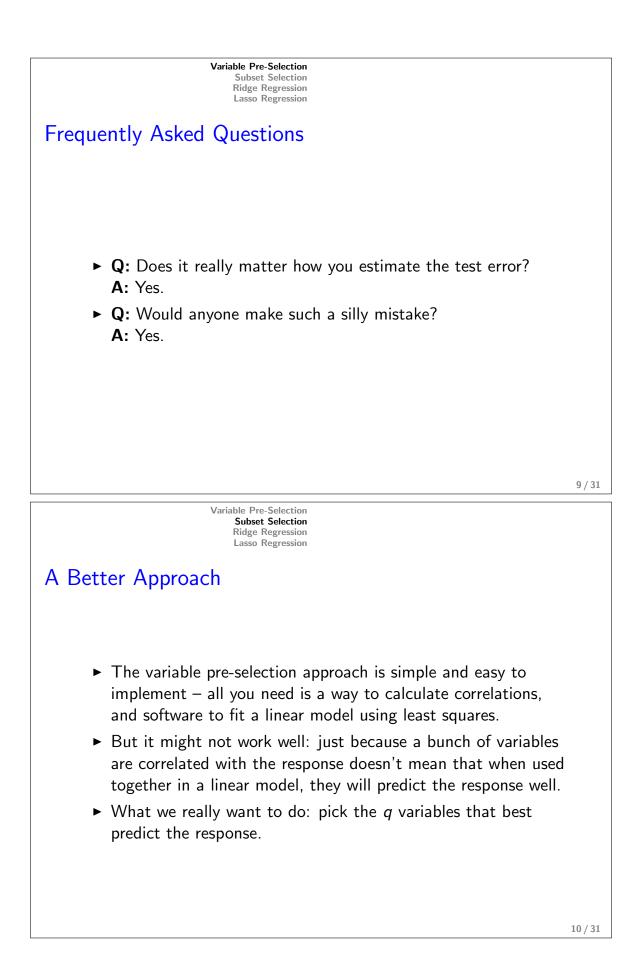
Estimating the Test Error For a Given \boldsymbol{q}

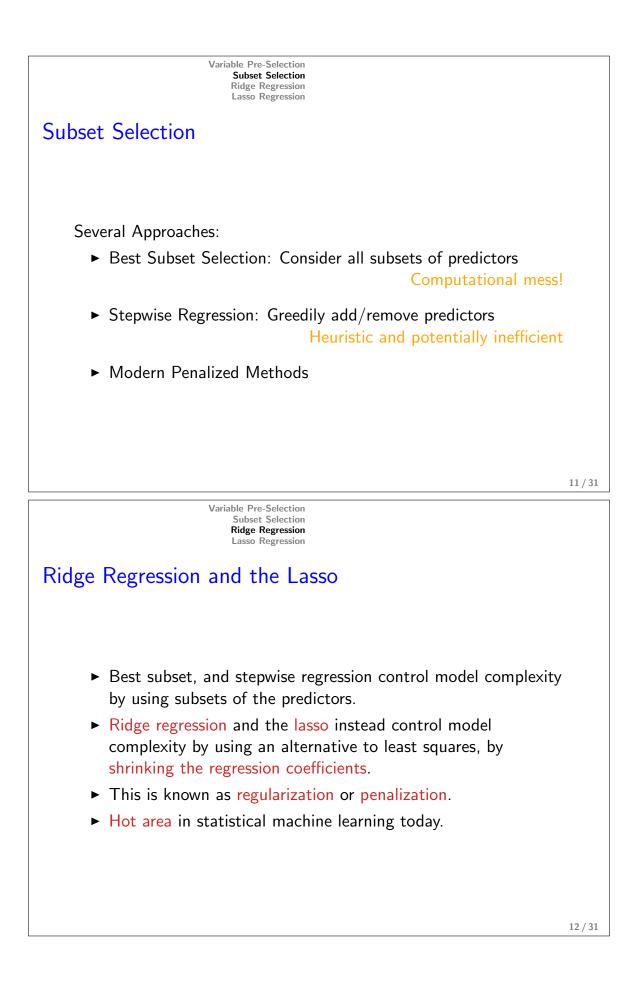
This is the right way to estimate the test error using the validation set approach:

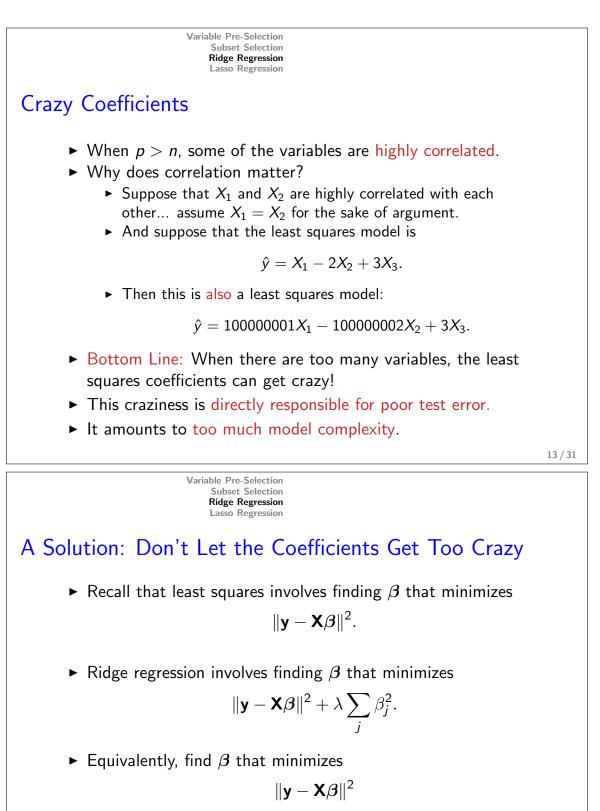
- 1. Split the observations into a training set and a validation set.
- 2. Using the training set only:
 - a. Identify the q variables most associated with the response.
 - b. Use least squares to fit a model predicting y using those q variables.
 - c. Let $\hat{\beta}_1, \ldots, \hat{\beta}_q$ denote the resulting coefficient estimates.
- 3. Use $\hat{\beta}_1, \ldots, \hat{\beta}_q$ obtained on training set to predict response on validation set, and compute the validation set MSE.





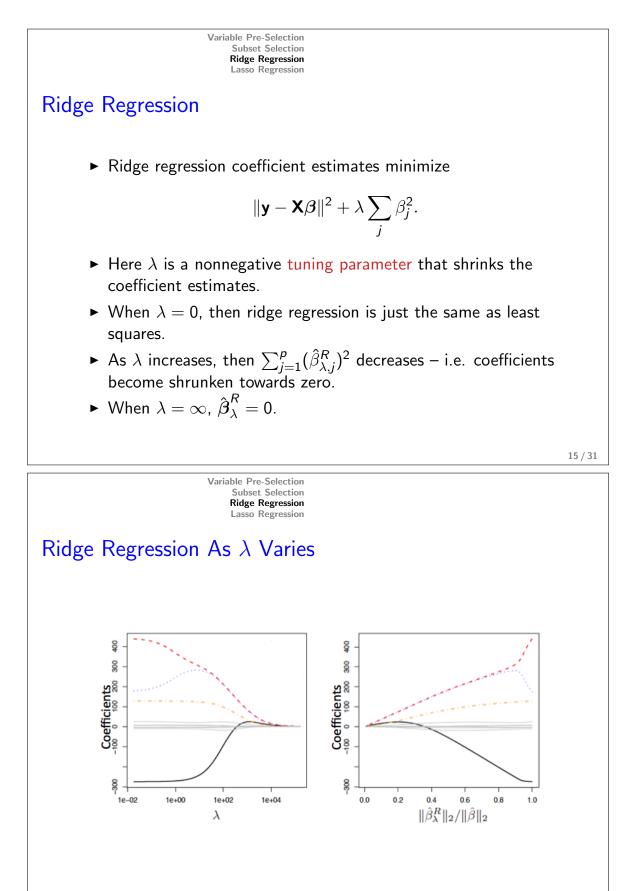


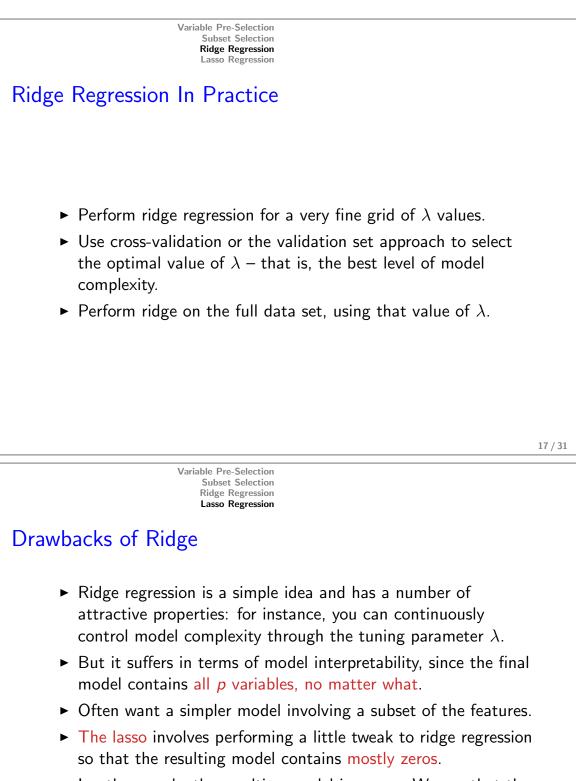




subject to the constraint that

$$\sum_{j=1}^p \beta_j^2 \le s.$$





- In other words, the resulting model is sparse. We say that the lasso performs feature selection.
- The lasso is a very active area of research interest in the statistical community!

Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression

The Lasso

• The lasso involves finding β that minimizes

$$\|\mathbf{y} - \mathbf{X}\boldsymbol{\beta}\|^2 + \lambda \sum_j |\beta_j|.$$

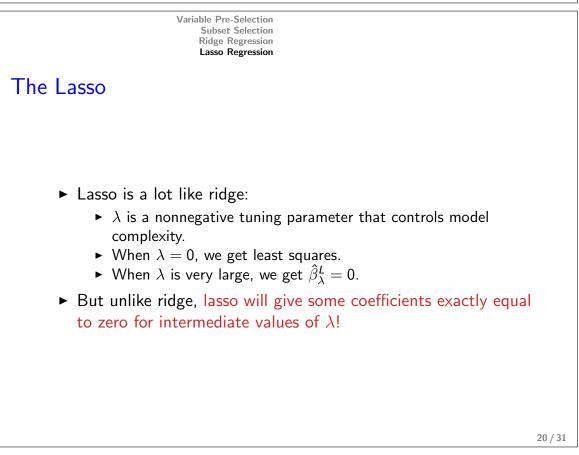
• Equivalently, find β that minimizes

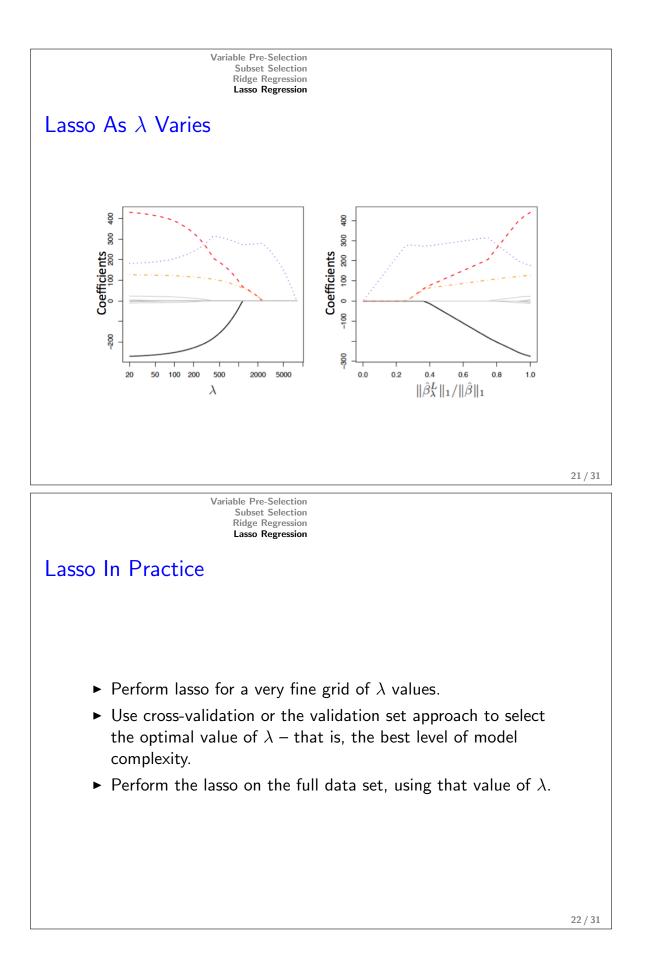
$$\|\mathbf{y} - \mathbf{X}\boldsymbol{\beta}\|^2$$

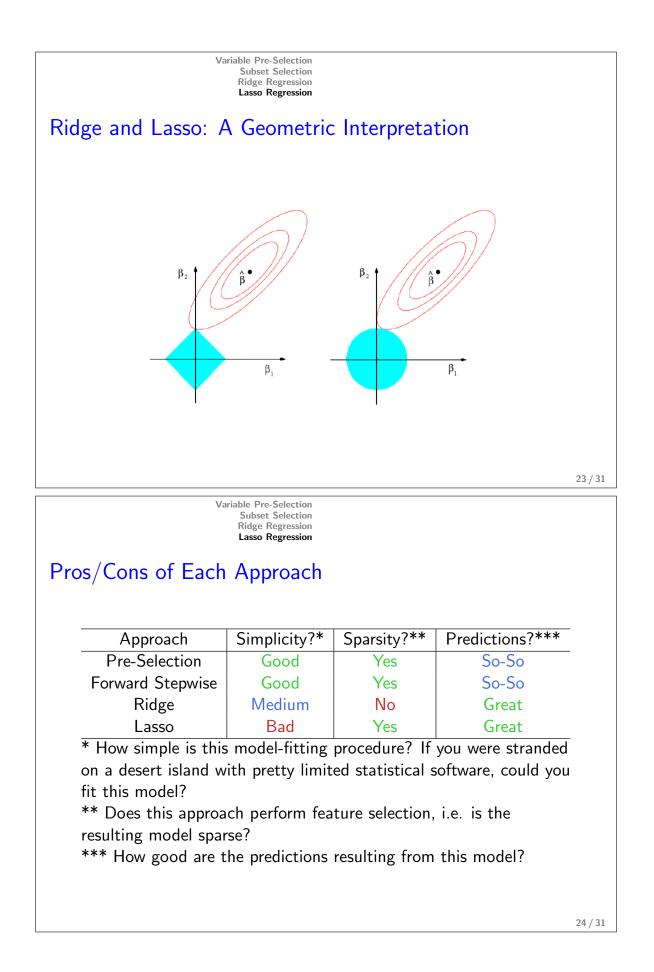
subject to the constraint that

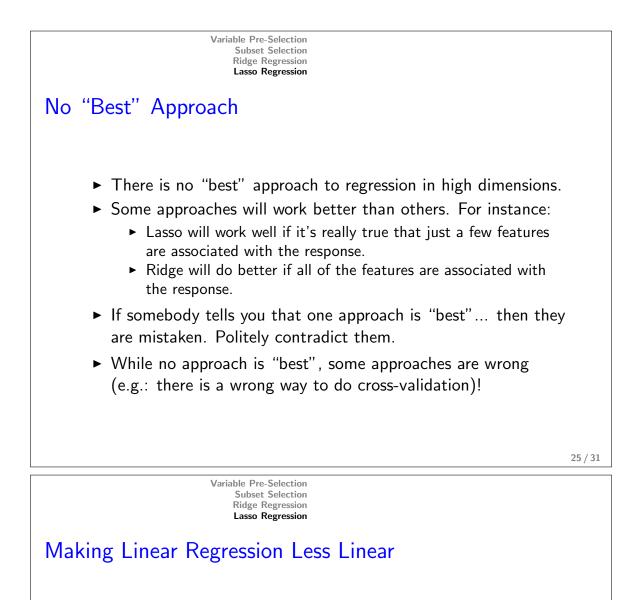
$$\sum_{j=1}^{p} |\beta_j| \le s.$$

So lasso is just like ridge, except that β²_j has been replaced with |β_j|.









What if the relationship isn't linear?

$$y = 3\sin(x) + \epsilon$$

$$y = 2e^{x} + \epsilon$$

$$y = 3x^{2} + 2x + 1 + \epsilon$$

If we know the functional form we can still use "linear regression"

Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression

Making Linear Regression Less Linear

$$y = 3\sin(x) + \epsilon:$$

$$\begin{pmatrix} x \\ x \end{pmatrix} \rightarrow \begin{pmatrix} \sin(x) \\ \sin(x) \end{pmatrix}$$

$$y = 3x^2 + 2x + 1 + \epsilon:$$

$$\begin{pmatrix} x \\ x \end{pmatrix} \rightarrow \begin{pmatrix} x \\ x^2 \end{pmatrix}$$

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Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression

Making Linear Regression Less Linear

What if we don't know the right functional form?

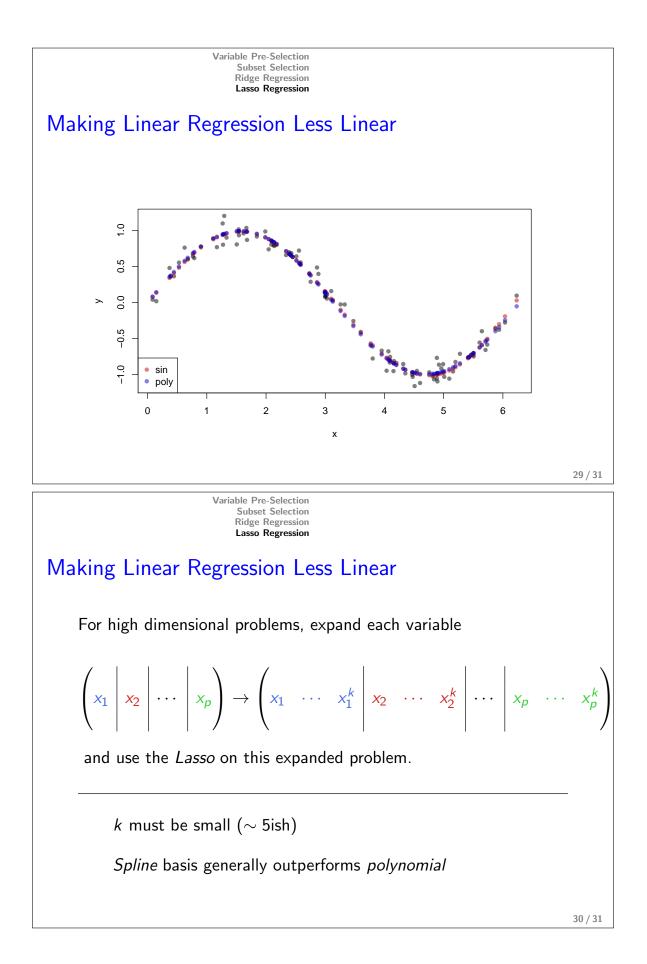
Use a flexible basis expansion:

polynomial basis

$$\left(x\right) \to \left(x \left| x^2 \right| \cdots \left| x^k \right)\right)$$

hockey-stick (/spline) basis

$$\begin{pmatrix} x \end{pmatrix} \rightarrow \begin{pmatrix} x \ x - t_1 \end{pmatrix}_+ \cdots \begin{vmatrix} (x - t_k)_+ \end{pmatrix}$$



Variable Pre-Selection Subset Selection Ridge Regression Lasso Regression	
Bottom Line	
Much more important than what model you fit is how you fit it.	
 Was cross-validation performed properly? Did you select a model (or level of model complexity) based 	
on an estimate of test error?	
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Classification

High-Dimensional Statistical Learning: Classification

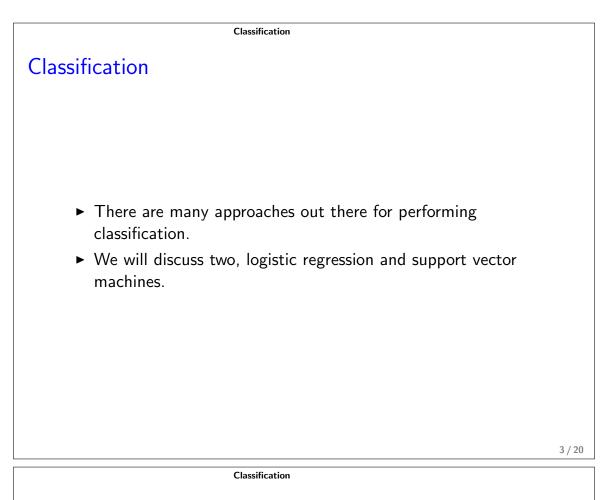
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Classification

Classification Regression involves predicting a continuous-valued response, like tumor size. Classification involves predicting a categorical response: Cancer versus Normal Tumor Type 1 versus Tumor Type 2 versus Tumor Type 3 Classification problems tend to occur even more frequently than regression problems in the analysis of omics data. Just like regression, Classification cannot be blindly performed in high-dimensions because you will get zero training error but awful test error; Properly estimating the test error is crucial; and There are a few tricks to extend classical classification approaches to high-dimensions, which we have already seen in the regression context!



Logistic Regression

- Logistic regression is the straightforward extension of linear regression to the classification setting.
- ► For simplicity, suppose y ∈ {0,1}: a two-class classification problem.
- ► The simple linear model

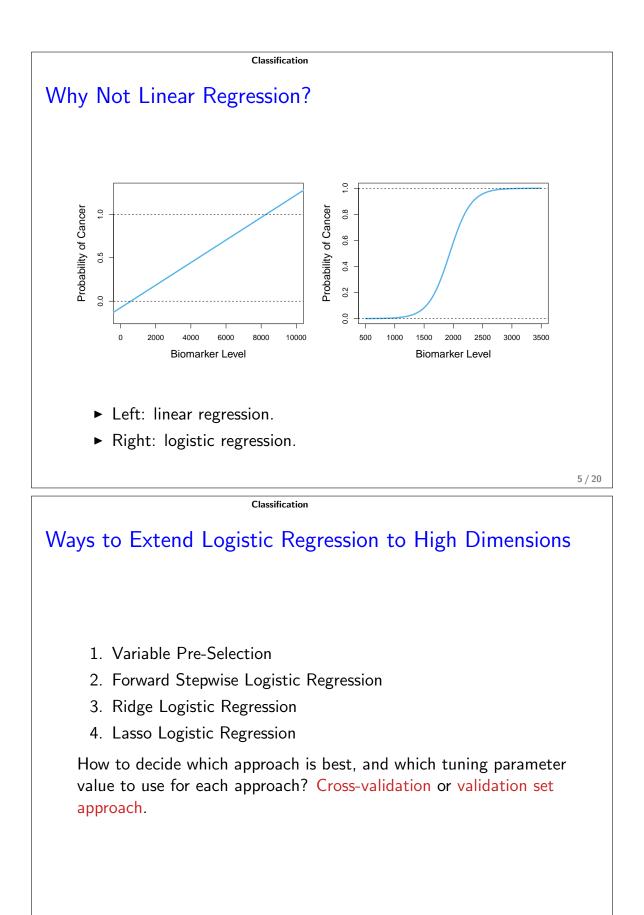
$$\mathbf{y} = \mathbf{X}\boldsymbol{eta} + \boldsymbol{\epsilon}$$

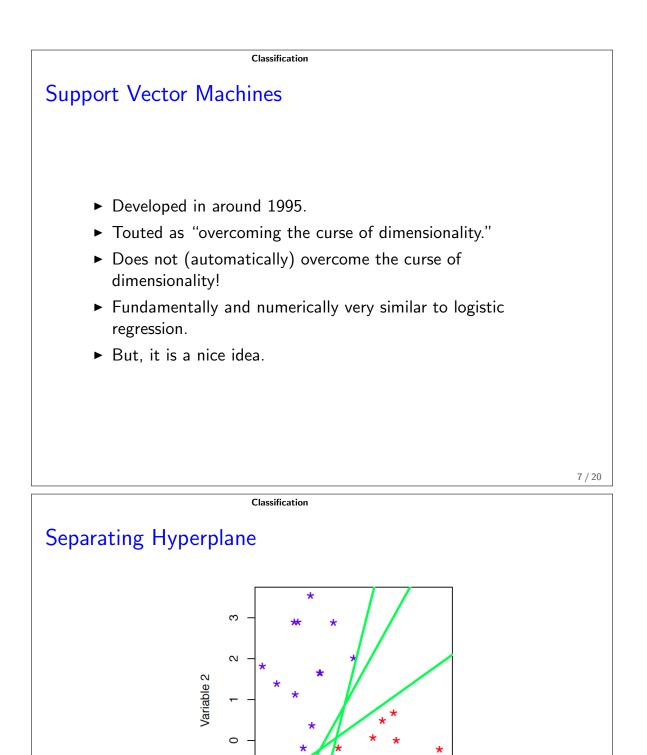
doesn't make sense for classification.

► Instead, the logistic regression model is

$$P(y=1|X) = \frac{\exp(X^{T}\beta)}{1 + \exp(X^{T}\beta)}.$$

 We usually fit this model using maximum likelihood – like least squares, but for logistic regression.





1

Variable 1

0

-1

2

3

T

