Classical Statistics Biological Big Data Supervised and Unsupervised Learning

Supervised Learning: Introduction

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Classical Statistics Biological Big Data

Supervised and Unsupervised Learning

Slides

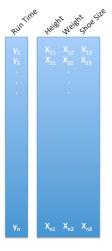
Slides and (some) codes are available at

https://github.com/SISBID/Module4

A Simple Example

- ▶ Suppose we have n = 500 kids for whom we have p = 3 measurements: height, weight, and shoe size.
- ► We wish to predict these kids' 1600-meter run times using these measurements.

A Simple Example



Notation:

- ▶ *n* is the number of observations.
- ▶ *p* the number of variables/features/predictors.
- y is a n-vector containing response/outcome for each of n observations.
- ightharpoonup X is a $n \times p$ data matrix.

Linear Regression on a Simple Example

➤ You can perform linear regression to develop a model to predict run time using height, weight, and shoe size:

$$y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \epsilon$$

where y is run time, X_1, X_2, X_3 are height, weight, and shoe size, and ϵ is a noise term.

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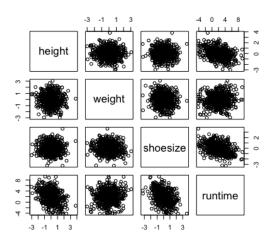
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- ➤ You can look at the coefficients, p-values, and t-statistics for your linear regression model in order to interpret your results.
- ► You learned everything (or most of what) you need to analyze this data set in AP Statistics!

A Relationship Between the Variables?



Linear Model Output

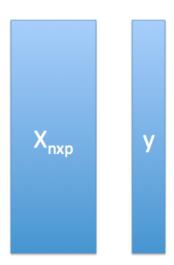
	Estimate	Std. Error	T-Stat	P-Value
Intercept	1.94179	0.09590	20.247	<2e-16 ***
height	-0.87704	0.09489	-9.243	<2e-16 ***
weight	0.07961	0.09105	0.874	0.382
shoesize	-1.00405	0.09530	-10.535	<2e-16 ***

RunTime $\approx 1.94 - 0.88 \times \text{Height} + 0.08 \times \text{Weight} - 1.00 \times \text{ShoeSize}$.

Low-Dimensional Versus High-Dimensional

- ▶ The data set that we just saw is low-dimensional: $n \gg p$.
- ▶ Lots of the data sets coming out of modern biological techniques are high-dimensional: $n \approx p$ or $n \ll p$.
- ➤ This poses statistical challenges! AP Statistics no longer applies.

Low Dimensional



High Dimensional



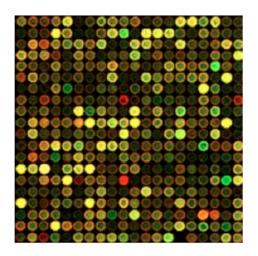
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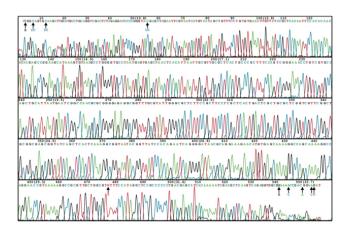
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- ▶ When $p \approx n$ or p > n, overfitting is guaranteed unless we are very careful.

Gene Expression Data



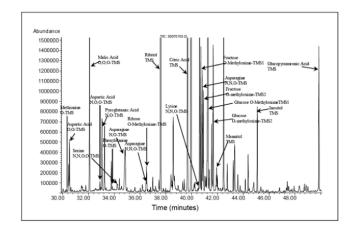
DNA Sequence Data



DNAse Hypersensitivity Data



Metabolomic Data



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- ▶ Predict risk of diabetes on the basis of DNA sequence data.... using n = 1000 patients and p = 3000000 variables.
- ► Cluster tissue samples on the basis of DNase hypersensitivity... using n = 200 cell types and p = 1000000000 variables.
- ▶ Identify genes whose expression is associated with survival time... using n = 250 cancer patients and p = 20000 variables.

Why Does Dimensionality Matter?

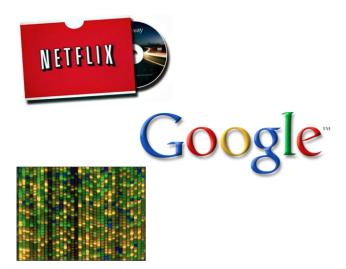
- Classical statistical techniques, such as linear regression, cannot be applied.
- ► Even very simple tasks, like identifying variables that are associated with a response, must be done with care.
- ► High risks of overfitting, false positives, and more.

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This course: Statistical machine learning tools for big – mostly high-dimensional – data.

Statistical Machine Learning



Supervised and Unsupervised Learning

► Statistical machine learning can be divided into two main areas: supervised and unsupervised.

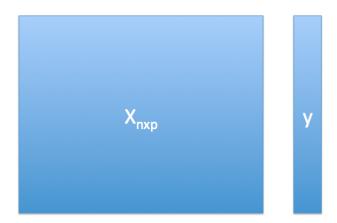
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- ► Supervised Learning: Use a data set *X* to predict or detect association with a response *y*.
 - ► Regression
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- ► Supervised Learning: Use a data set *X* to predict or detect association with a response *y*.
 - ► Regression
 - ► Classification
- ► Unsupervised Learning: Discover the signal in *X*, or detect associations within *X*.
 - ► Dimension Reduction
 - Clustering
 - ► Hypothesis Testing

Supervised Learning



Unsupervised Learning

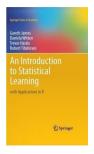


This Course

- ► We will cover the big ideas in supervised learning for big data.
- ► The best way to use these methods: learn R.



"Course Textbook" . . . with applications in R



- ► Available for (free!) download from www.statlearning.com.
- ► An accessible introduction to statistical machine learning, with an R lab at the end of each chapter!!
- We will go through some of these R labs in class.
- ► To learn more, go through them on your own!

Let's Try Out Some R!

Chapter 2 R lab www.statlearning.com

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- ► Classification: Predict a categorical response, such as
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 - subtype of glioblastoma
- ► We will start with Regression.

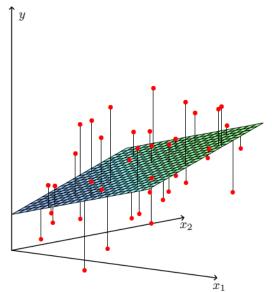
Linear Models

- ▶ 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

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

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

Linear Model With p = 2 Predictors



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- ► This is a linear model:

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► This is not a linear model:

$$y_i = \beta_1^{X_{i1}} + \sin(\beta_2 X_{i2}) + \epsilon_i.$$

Linear Models in Matrix Form

- ▶ For simplicity, ignore the intercept β_0 .
 - Assume $\sum_{i=1}^{n} y_i = \sum_{i=1}^{n} X_{ij} = 0$; in this case, $\beta_0 = 0$.
 - ► Alternatively, let the first column of X be a column of 1's.

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 - ► Alternatively, let the first column of X be a column of 1's.
- In matrix form, we can write the linear model as

$$y = X\beta + \epsilon$$
,

i.e.

$$\begin{pmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{pmatrix} = \begin{pmatrix} X_{11} & X_{12} & \dots & X_{1p} \\ X_{21} & X_{22} & \dots & X_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ X_{n1} & X_{n2} & \dots & X_{np} \end{pmatrix} \begin{pmatrix} \beta_1 \\ \beta_2 \\ \vdots \\ \beta_p \end{pmatrix} + \begin{pmatrix} \epsilon_1 \\ \epsilon_2 \\ \vdots \\ \epsilon_n \end{pmatrix}.$$

Least Squares Regression

- ▶ There are a lot of ways we could fit the model $y = X\beta + \epsilon$.
- ► Most common approach in classical statistics is least squares:

$$\mathop{\mathsf{minimize}}_{\boldsymbol{\beta}} \left\{ \| \mathbf{y} - \mathbf{X} \boldsymbol{\beta} \|^2 \right\}.$$

Here
$$\|\mathbf{a}\|^2 \equiv \sum_{i=1}^n a_i^2$$
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Here $\|a\|^2 \equiv \sum_{i=1}^n a_i^2$.

▶ We are looking for β_1, \ldots, β_p such that

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

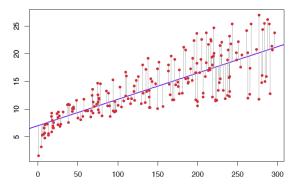
is as small as possible.

► Equivalently, we're looking for coefficient estimates 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.

Least Squares



► Horizontal axis: predictor

► Vertical axis: response

► Red dots: observations

► Purple line: least squares line

Purple line minimizes sum of squared lengths of the gray lines.

Let's Try Out Least Squares in R!

Chapter 3 R lab www.statlearning.com