

# **CSCI446/946 Big Data Analytics**

## **Week 3      Data Analytic Methods Using R**

School of Computing and Information Technology  
University of Wollongong Australia

# Data Analytic Methods Using R

- Introduction to R
  - R, RStudio, Data I/O, Attribute and Data Types
  - Descriptive statistics
- Exploratory Data Analysis
  - Visualization before analysis
  - Visualizing single or multiple variables
- Statistical Methods for Evaluation
  - Hypothesis Testing, ANOVA

All the figures, tables and codes are from the book “[Data Science and Big Data Analytics: Discovering, Analyzing, Visualizing and Presenting Data](#)” unless indicated otherwise.

# Data Analytic Methods Using R

- The success of a data analysis project requires a deep understanding of the data
- It requires a toolbox for mining and presenting the data
  - Basic statistical measures
  - Creation of graphs and plots
  - Identify relationships and patterns
- R: popularity and versatility

# Introduction to R

- A programming language and software framework for statistical analysis and graphics
- [Comprehensive R Archive Network](#)
- An overview the basic functionality of R
- We begin with understanding the flow of a basic R script to address an analytic problem
  - Command-line interface (CLI)
  - Graphical user interface (GUI)

# Introduction to R

- The first example

```
# import a csv file of the total annual sales for each  
customer  
sales <- read.csv("./yearly_sales.csv")
```

```
# examine the imported dataset  
head(sales)  
summary(sales)
```

```
# plot num_of_orders vs. sales  
plot(sales$num_of_orders,sales$sales_total,  
      main="Number of Orders vs. Sales")
```

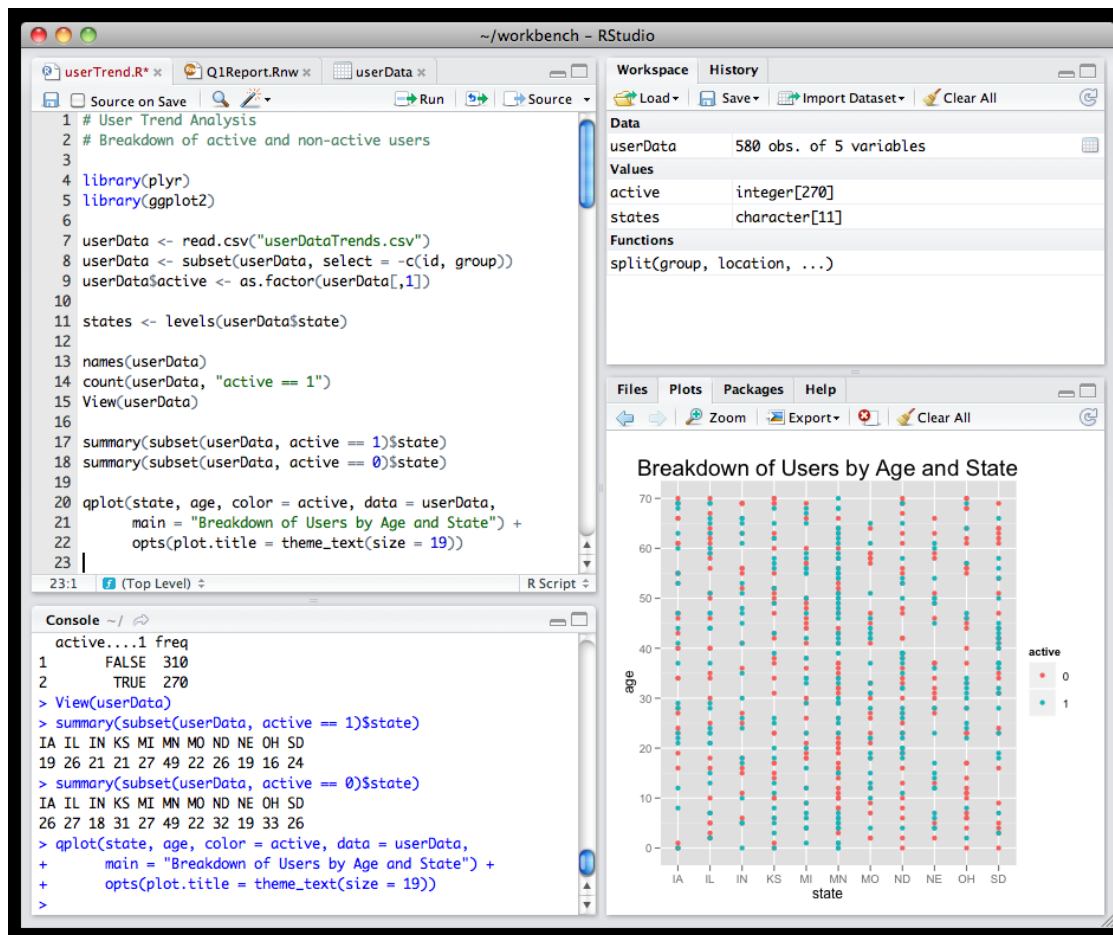
# Introduction to R

- The first example

```
# perform a statistical analysis (fit a linear  
regression model)  
results <- lm(sales$sales_total ~ sales$num_of_orders)  
results  
summary(results)  
  
# perform some diagnostics on the fitted model  
# plot histogram of the residuals  
hist(results$residuals, breaks = 800)
```

# Introduction to R

- R Graphical User Interface (RStudio)



- Scripts
- Workspace
- Plots
- Console

# Introduction to R

- **Help** functionality
  - `Help(lm)` or `?lm`
- `Edit()` and `fix()`
  - Allow to update the contents of an R variable
- `Save.image()` function to **create** .Rdata file
- `Load.image()` function to **load** .Rdata file
- Please install R and RStudio to try out the R examples



# Introduction to R

- Data Import and Export

```
sales <- read.csv("c:/data/yearly_sales.csv")
```

```
setwd("c:/data/")
```

```
sales <- read.csv("yearly_sales.csv")
```

```
# add a column for the average sales per order
```

```
sales$per_order <- sales$sales_total/sales$num_of_orders
```

```
# export data as tab delimited without the row names
```

```
write.table(sales,"sales_modified.txt", sep="\t",  
row.names=FALSE)
```

# Introduction to R

- Automatically save plots

```
# export a histogram to a jpeg
jpeg(file="c:/data/sales_hist.jpeg") # create a new jpeg
file
hist(sales$num_of_orders) # export histogram to jpeg
dev.off() # shut off the graphic device
```

- More information
  - <https://cran.r-project.org/doc/manuals/r-release/R-data.html>

# Introduction to R

- Attribute and Data Types
- Attributes: Nominal, Ordinal, Interval, and Ratio (NOIR)

	Categorical (Qualitative)		Numeric (Quantitative)	
	Nominal	Ordinal	Interval	Ratio
Definition	The values represent labels that distinguish one from another.	Attributes imply a sequence.	The difference between two values is meaningful.	Both the difference and the ratio of two values are meaningful.
Examples	ZIP codes, nationality, street names, gender, employee ID numbers, TRUE or FALSE	Quality of diamonds, academic grades, magnitude of earthquakes	Temperature in Celsius or Fahrenheit, calendar dates, latitudes	Age, temperature in Kelvin, counts, length, weight
Operations	=, ≠	=, ≠, <, ≤, >, ≥	=, ≠, <, ≤, >, ≥, +, -	=, ≠, <, ≤, >, ≥, +, -, ×, ÷

# Introduction to R

- Data Types
  - Numeric, character, logical (and list)

```
i <- 1                # create a numeric variable
sport <- "football"    # create a character variable
flag <- TRUE           # create a logical variable

class(i)              # returns "numeric"
typeof(i)             # returns "double"
class(sport)          # returns "character"
typeof(sport)         # returns "character"
class(flag)           # returns "logical"
typeof(flag)          # returns "logical"
```

# Introduction to R

- Vectors
  - A basic building block for data in R
  - Simple R variables are actually vectors
  - Can only consist of values in the same class

```
# Vectors
```

```
is.vector(i)
```

```
# returns TRUE
```

```
is.vector(flag)
```

```
# returns TRUE
```

```
is.vector(sport)
```

```
# returns TRUE
```



# Introduction to R

- `vector()` function, by default, create a logical vector

```
a <- vector(length=3)      # create a logical vector of length 3
a                          # returns FALSE FALSE FALSE
b <- vector(mode="numeric", 3) # create a numeric vector of length 3
typeof(b)                  # returns "double"
b[2] <- 3.1                 # assign 3.1 to the 2nd element
b                          # returns 0.0 3.1 0.0
c <- vector(mode="integer", 0) # create an integer vector of length 0
c                          # returns integer(0)
length(c)                  # returns 0
```

# Introduction to R

- Arrays and Matrices

```
# the dimensions are 3 regions, 4 quarters, and 2 years
quarterly_sales <- array(0, dim=c(3,4,2))
quarterly_sales[2,1,1] <- 158000
quarterly_sales
```

```
sales_matrix <- matrix(0, nrow = 3, ncol = 4)
sales_matrix
```

```
install.packages("matrixcalc") # install, if necessary
library(matrixcalc)
```

```
# build a 3x3 matrix
M <- matrix(c(1,3,3,5,0,4,3,3,3),nrow = 3,ncol = 3)
M %%% matrix.inverse(M) # multiply M by inverse(M)
```



# Introduction to R

- Data Frames

- A structure for storing and accessing several variables of possibly **different** data types
- Preferred input format for many R functions

```
sales <- read.csv("c:/data/yearly_sales.csv")  
is.data.frame(sales)                # returns TRUE
```

```
is.vector(sales$cust_id)             # returns TRUE  
is.vector(sales$sales_total)        # returns TRUE  
is.vector(sales$num_of_orders)      # returns TRUE  
is.vector(sales$gender)             # returns FALSE  
is.factor(sales$gender)             # returns TRUE
```

# Introduction to R

- **List**: a collection of objects that can be of various types, including other lists

```
sales <- read.csv("c:/data/yearly_sales.csv")  
class(sales)  
typeof(sales)
```

```
# build an assorted list of a string, a numeric,  
# a list, a vector, and a matrix  
housing <- list("own", "rent")  
assortment <- list("football", 7.5, housing, v, M)  
assortment
```

```
str(assortment)
```

# Introduction to R

- **Factors**: a **categorical** variable, typically with a few **finite levels** such as “F” and “M”
- **Factors** can be ordered or not ordered

```
# Factors
```

```
class(sales$gender)      # returns "factor"  
is.ordered(sales$gender) # returns FALSE
```

- Use of **factors** is important in R statistical modelling functions

# Introduction to R

- Contingency Tables

- A class of objects used to store the observed counts across the factors for a given dataset
- The basis for performing a statistical test on the independence of the factors

```
# build a contingency table based on the gender and  
spender factors  
sales_table <- table(sales$gender,sales$spender)  
sales_table
```

# Introduction to R

- Contingency Tables

- A class of objects used to store the observed counts across the factors for a given dataset
- The basis for performing a statistical test on the independence of the factors

```
class(sales_table)           # returns "table"  
typeof(sales_table)          # returns "integer"  
dim(sales_table)              # returns 2 3  
  
# performs a chi-squared test  
summary(sales_table)
```

# Introduction to R

- Descriptive Statistics

- `Summary()` function: mean, median, min, max
- R functions include descriptive statistics

```
# to simplify the function calls, assign
```

```
x <- sales$sales_total
```

```
y <- sales$num_of_orders
```

```
cor(x,y)           # returns 0.7508015 (correlation)
```

```
cov(x,y)           # returns 345.2111 (covariance)
```

```
IQR(x)             # returns 215.21 (interquartile range)
```

```
mean(x)            # returns 249.4557 (mean)
```

```
median(x)          # returns 151.65 (median)
```

```
range(x)           # returns 30.02 7606.09 (min max)
```

```
sd(x)              # returns 319.0508 (std. dev.)
```

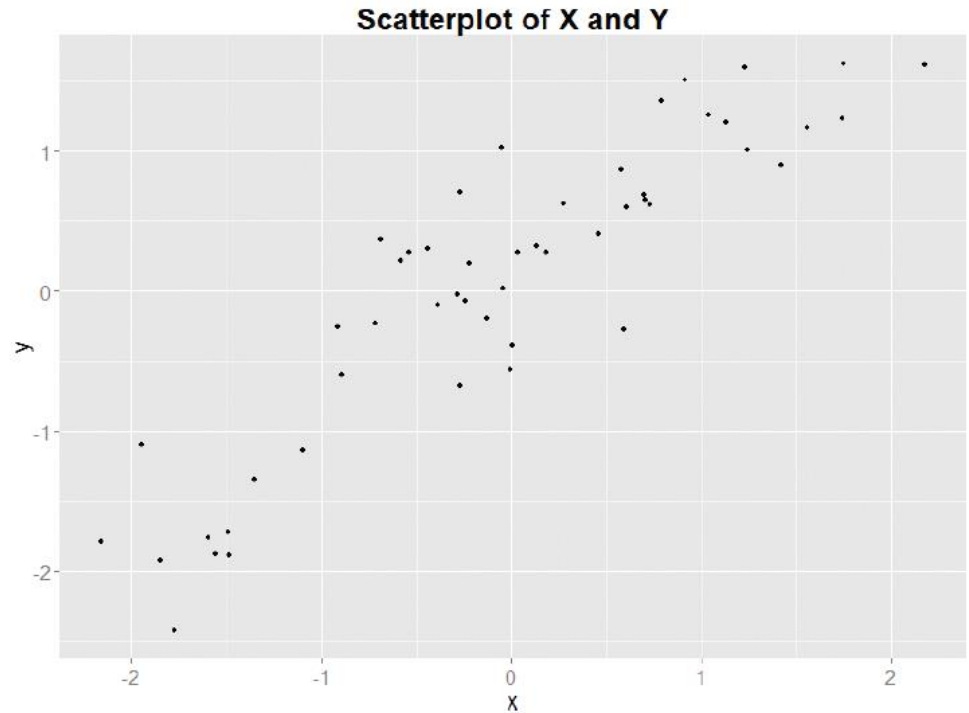
```
var(x)             # returns 101793.4 (variance)
```

# Exploratory Data Analysis

- Linear relationship and distributions are more difficult to see from descriptive statistics

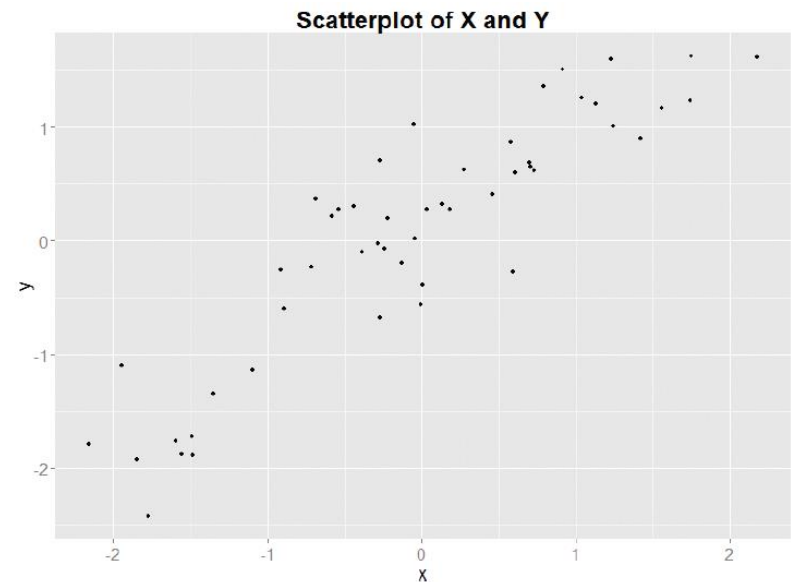
```
summary(data)
```

x	y
Min. : -1.90483	Min. : -2.16545
1st Qu.: -0.66321	1st Qu.: -0.71451
Median : 0.09367	Median : -0.03797
Mean : 0.02522	Mean : -0.02153
3rd Qu.: 0.65414	3rd Qu.: 0.55738
Max. : 2.18471	Max. : 1.70199



# Exploratory Data Analysis

- Detect patterns and anomalies in the data
  - Through exploratory data analysis by **visualization**
  - **Visualization** gives a succinct, holistic view
  - **Visualization** is an important facet at the initial data exploration



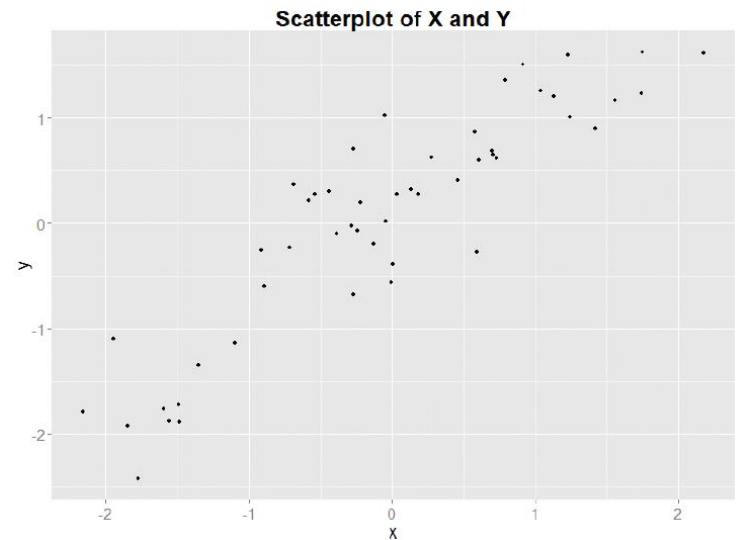


# Exploratory Data Analysis

```
# Figure 3-5  
x <- rnorm(50)  
y <- x + rnorm(50, mean=0, sd=0.5)
```

```
data <- as.data.frame(cbind(x, y))  
summary(data)
```

```
library(ggplot2)  
ggplot(data, aes(x=x, y=y)) +  
  geom_point(size=2) +  
  ggtitle("Scatterplot of X and Y") +  
  theme(axis.text=element_text(size=12),  
        axis.title = element_text(size=14),  
        plot.title = element_text(size=20, face="bold"))
```



# Exploratory Data Analysis

- Visualization Before Analysis

# 1		# 2		# 3		# 4	
x	y	x	y	x	y	x	y
4	4.26	4	3.10	4	5.39	8	5.25
5	5.68	5	4.74	5	5.73	8	5.56
6	7.24	6	6.13	6	6.08	8	5.76
7	4.82	7	7.26	7	6.42	8	6.58
8	6.95	8	8.14	8	6.77	8	6.89
9	8.81	9	8.77	9	7.11	8	7.04
10	8.04	10	9.14	10	7.46	8	7.71
11	8.33	11	9.26	11	7.81	8	7.91
12	10.84	12	9.13	12	8.15	8	8.47
13	7.58	13	8.74	13	12.74	8	8.84
14	9.96	14	8.10	14	8.84	19	12.50

**FIGURE 3-6** *Anscombe's quartet*

# Exploratory Data Analysis

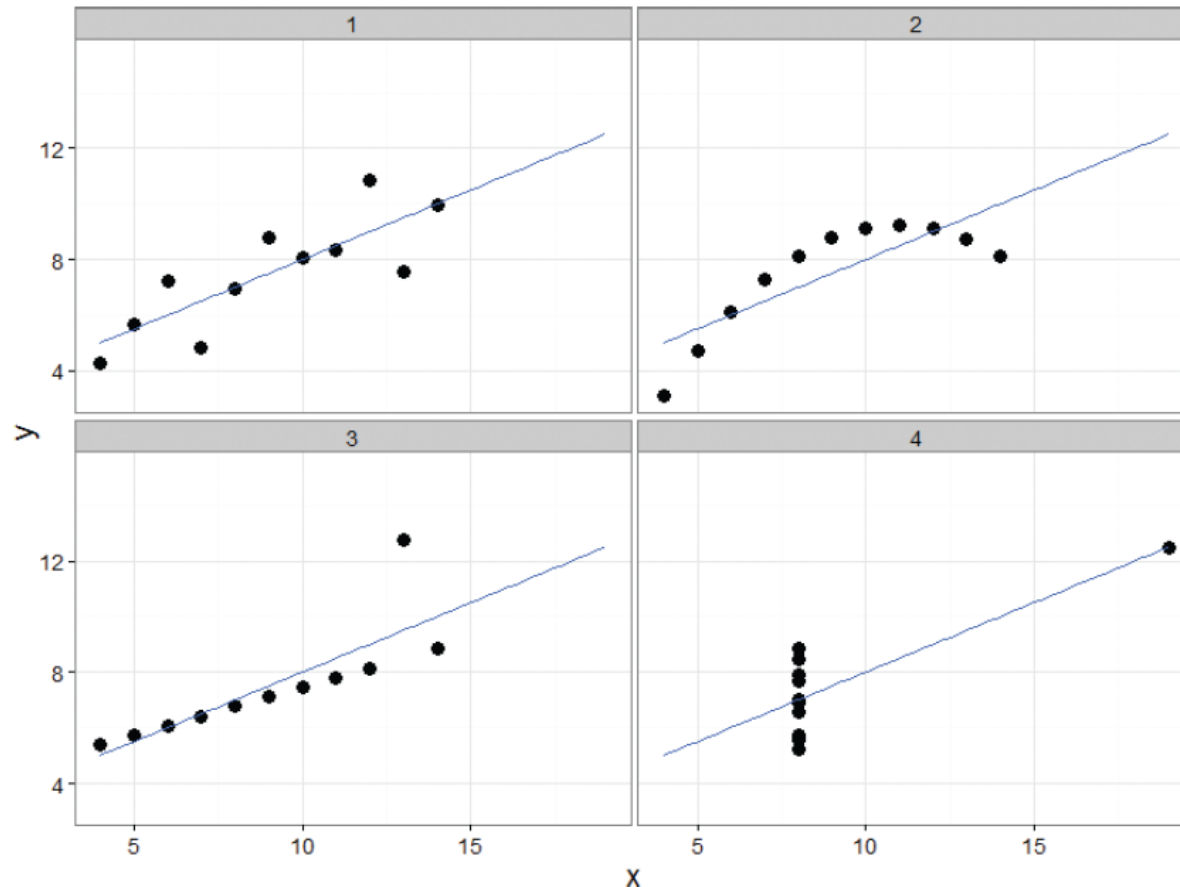
- The four data sets have **nearly identical** statistical properties

**TABLE 3-3** *Statistical Properties of Anscombe's Quartet*

Statistical Property	Value
Mean of $x$	9
Variance of $y$	11
Mean of $y$	7.50 (to 2 decimal points)
Variance of $y$	4.12 or 4.13 (to 2 decimal points)
Correlations between $x$ and $y$	0.816
Linear regression line	$y = 3.00 + 0.50x$ (to 2 decimal points)

# Exploratory Data Analysis

- However, the **reality** is a **different** story...

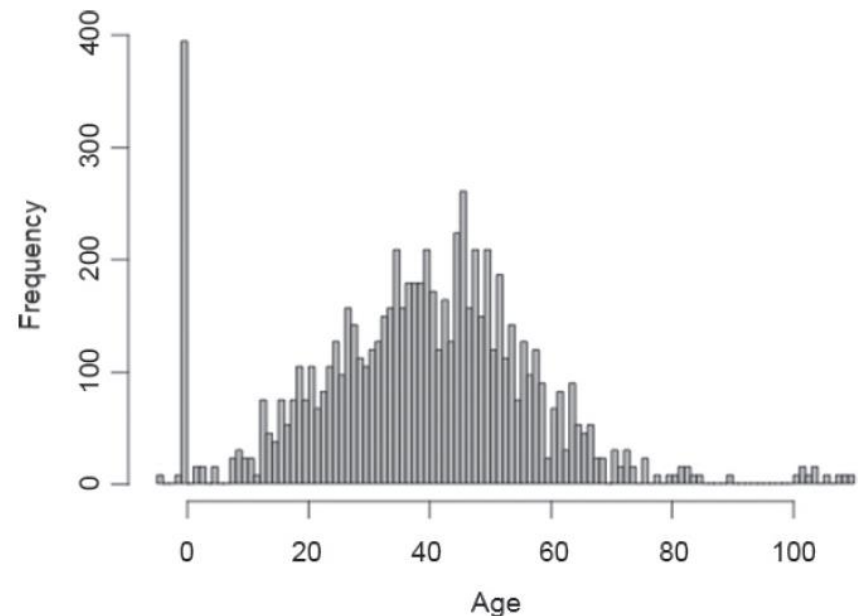


# Exploratory Data Analysis

- Dirty Data

- Detect dirty data with visualization
- Look for **anomalies**, verify with **domain** knowledge
- **Clean** the data appropriately

```
hist(age, breaks=100,  
main="Age Distribution of  
Account Holders",  
xlab="Age",  
ylab="Frequency", col="gray")
```



# Exploratory Data Analysis

- Any **dirty data**?



```
hist(mortgage, breaks=10, xlab="Mortgage Age", col="gray",  
     main="Portfolio Distribution, Years Since Origination")
```

# Exploratory Data Analysis

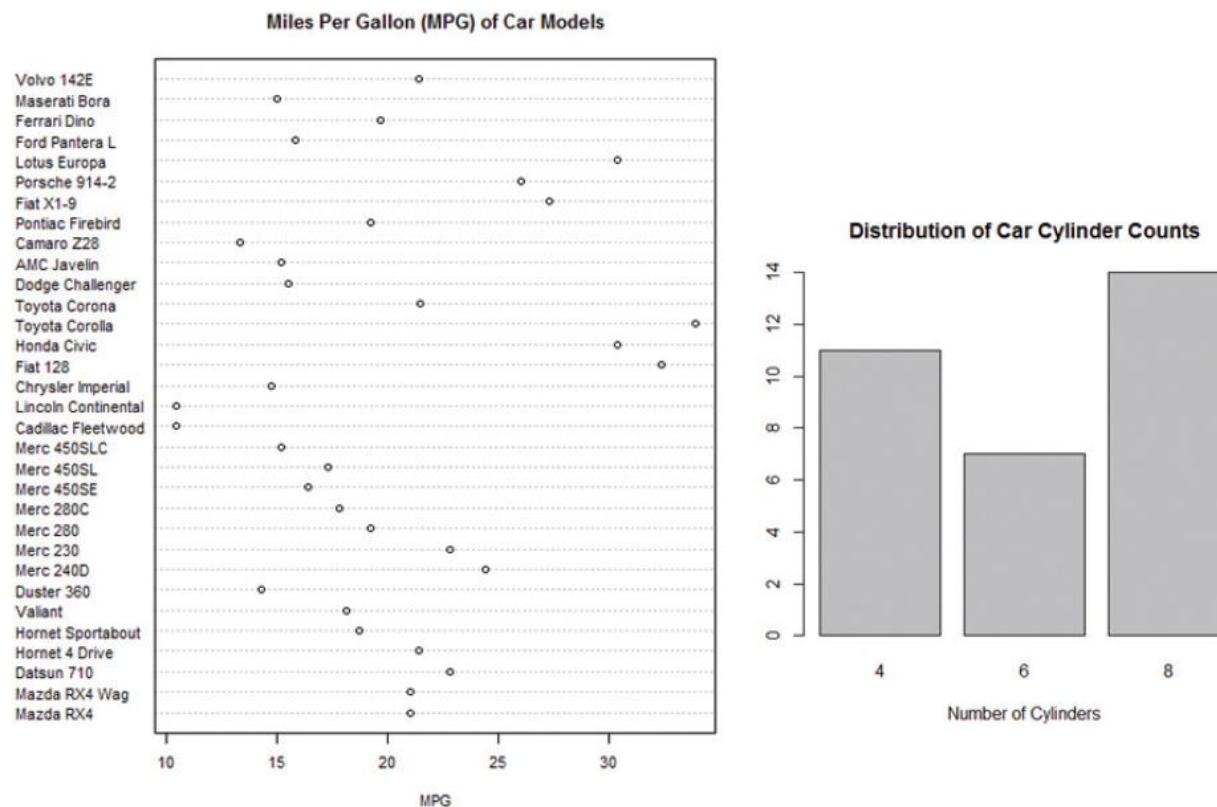
- Visualizing a Single Variable

**TABLE 3-4** *Example Functions for Visualizing a Single Variable*

Function	Purpose
<code>plot(data)</code>	Scatterplot where x is the index and y is the value; suitable for low-volume data
<code>barplot(data)</code>	Barplot with vertical or horizontal bars
<code>dotchart(data)</code>	Cleveland dot plot [12]
<code>hist(data)</code>	Histogram
<code>plot(density(data))</code>	Density plot (a continuous histogram)
<code>stem(data)</code>	Stem-and-leaf plot
<code>rug(data)</code>	Add a rug representation (1-d plot) of the data to an existing plot

# Exploratory Data Analysis

- Visualizing a Single Variable



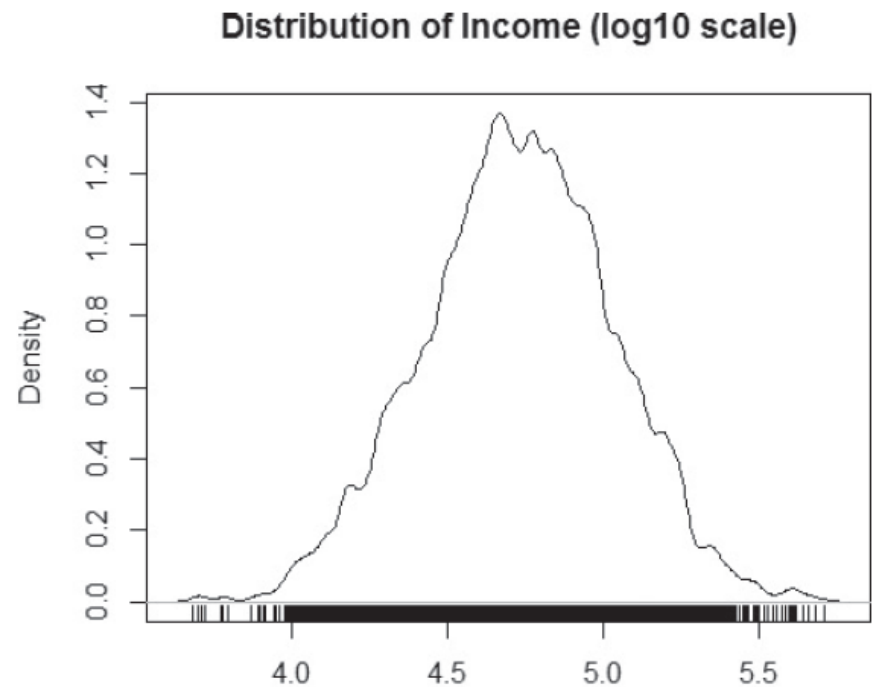
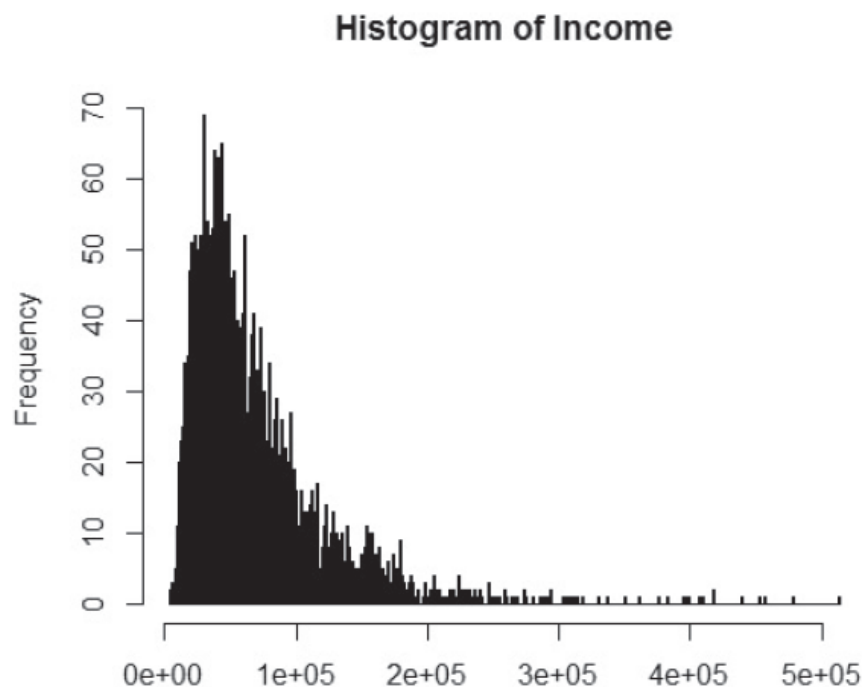
## Dotchart and Barplot ##

```
dotchart(mtcars$mpg, labels=row.names(mtcars), cex=.7, main="Miles Per Gallon (MPG) of Car Models", xlab="MPG")  
barplot(table(mtcars$cyl), main="Distribution of Car Cylinder Counts", xlab="Number of Cylinders")
```



# Exploratory Data Analysis

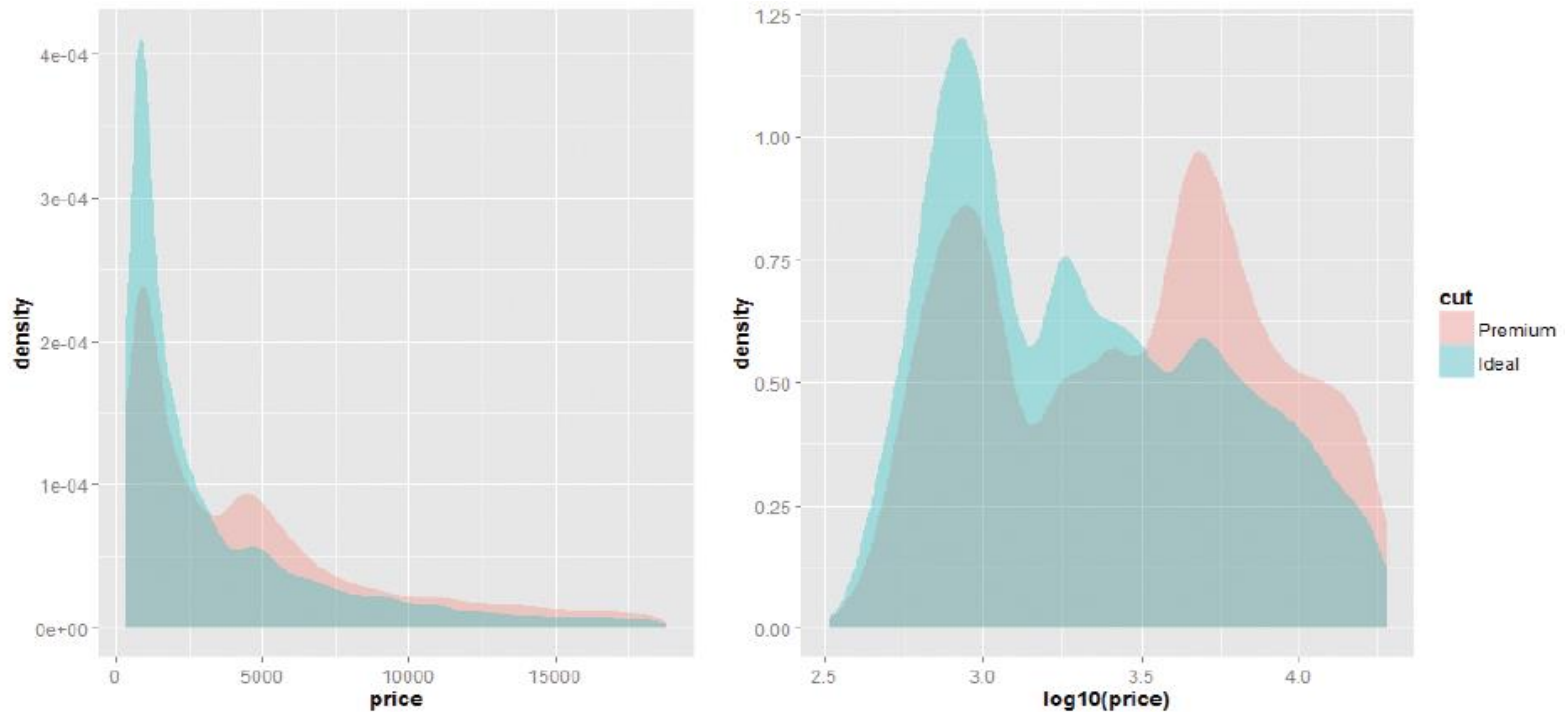
- Visualizing a Single Variable (log transformation)



```
# plot the histogram
hist(income, breaks=500, xlab="Income", main="Histogram of Income")
# density plot
plot(density(log10(income), adjust=0.5), main="Distribution of Income (log10 scale)")
# add rug to the density plot
rug(log10(income))
```

# Exploratory Data Analysis

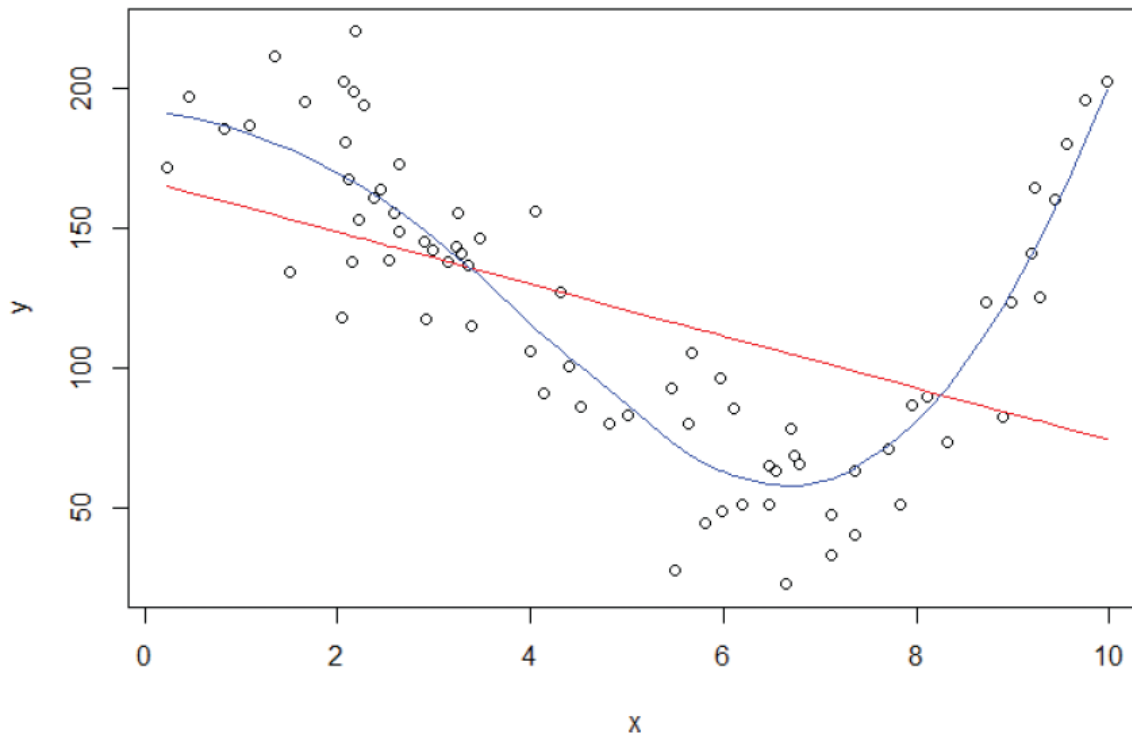
- Visualizing a Single Variable (unimodal or multimodal?)



```
# plot density plot of diamond prices
ggplot(niceDiamonds, aes(x=price, fill=cut)) + geom_density(alpha = .3, color=NA)
# plot density plot of the log10 of diamond prices
ggplot(niceDiamonds, aes(x=log10(price), fill=cut)) + geom_density(alpha = .3, color=NA)
```

# Exploratory Data Analysis

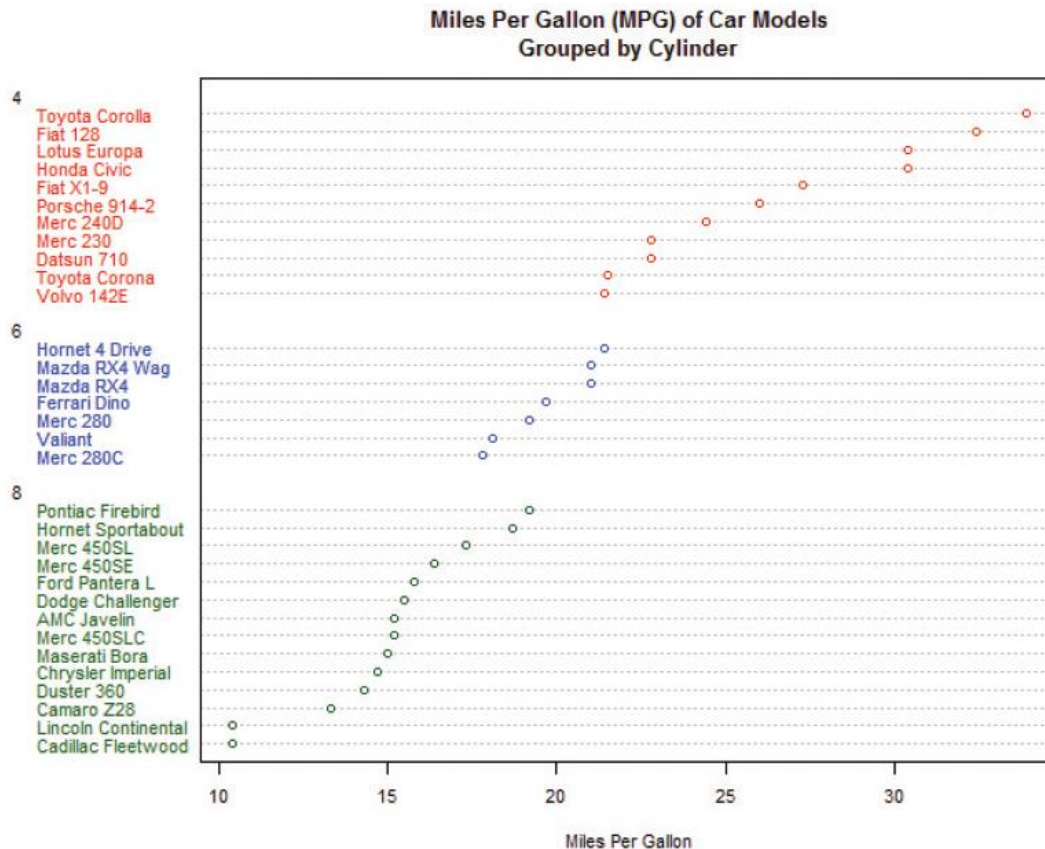
- Examining Multiple Variable



```
# 75 numbers between 0 and 10 of
uniform distribution
x <- runif(75, 0, 10)
x <- sort(x)
y <- 200 + x^3 - 10 * x^2 + x +
rnorm(75, 0, 20)
lr <- lm(y ~ x) # linear
regression
poly <- loess(y ~ x) # LOESS
fit <- predict(poly) # fit a
nonlinear line
plot(x,y)
# draw the fitted line for the
linear regression
points(x, lr$coefficients[1] +
lr$coefficients[2] * x,
      type = "l", col = 2)
# draw the fitted line with LOESS
points(x, fit, type = "l", col =
4)
```

# Exploratory Data Analysis

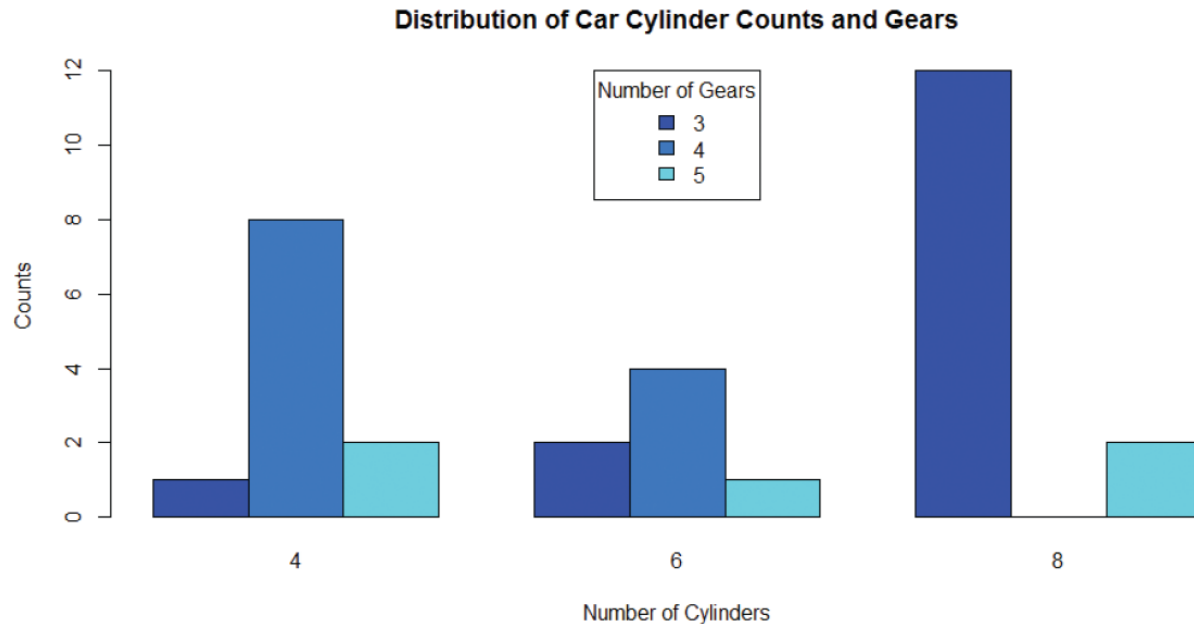
- Examining Multiple Variable



```
# sort by mpg
cars <-
mtcars[order(mtcars$mpg),]
# grouping variable must be a
factor
cars$cyl <- factor(cars$cyl)
cars$color[cars$cyl==4] <-
"red"
cars$color[cars$cyl==6] <-
"blue"
cars$color[cars$cyl==8] <-
"darkgreen"
dotchart(cars$mpg,
labels=row.names(cars),
cex=.7, groups= cars$cyl,
main="Miles Per
Gallon (MPG) of Car
Models\nGrouped by Cylinder",
xlab="Miles Per
Gallon", color=cars$color,
gcolor="black")
```

# Exploratory Data Analysis

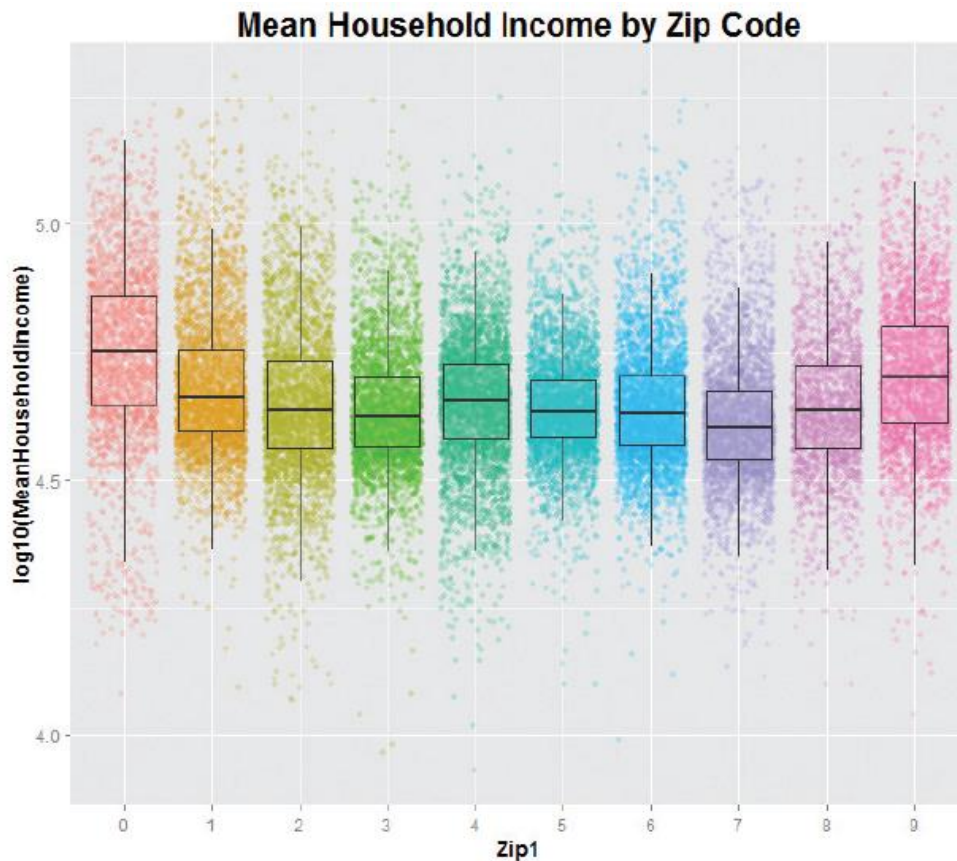
- Examining Multiple Variable



```
counts <- table(mtcars$gear, mtcars$cyl)
barplot(counts, main="Distribution of Car Cylinder Counts and Gears",
        xlab="Number of Cylinders", ylab="Counts",
        col=c("#0000FFFF", "#0080FFFF", "#00FFFFFF"),
        legend = rownames(counts), beside=TRUE,
        args.legend = list(x="top", title = "Number of Gears"))
```

# Exploratory Data Analysis

- Examining Multiple Variable (box-and-whisker plot)



## Box-and-Whisker Plot ##

```
DF <- read.csv("c:/data/zipIncome.csv", header=TRUE, sep=",")
```

```
# Remove outliers
```

```
DF <- subset(DF, DF$MeanHouseholdIncome > 7000 &  
DF$MeanHouseholdIncome < 200000)  
summary(DF)
```

```
library(ggplot2)
```

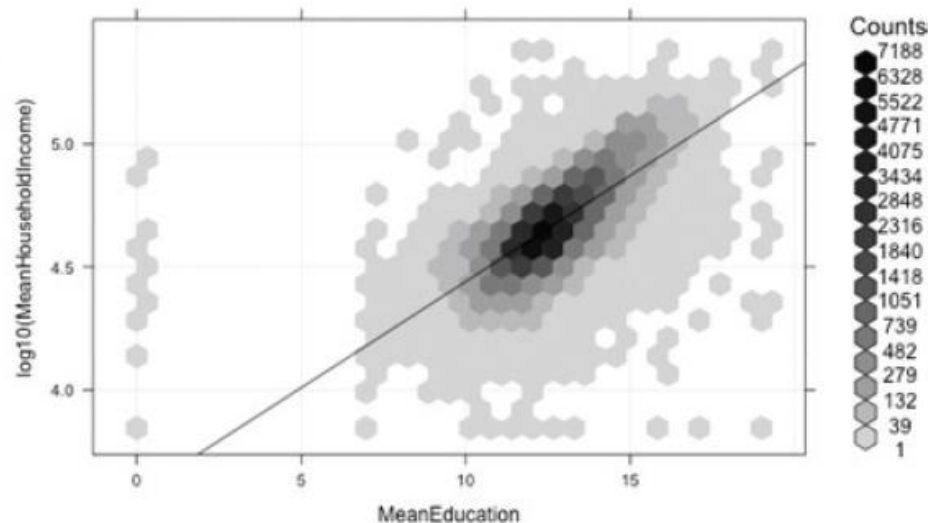
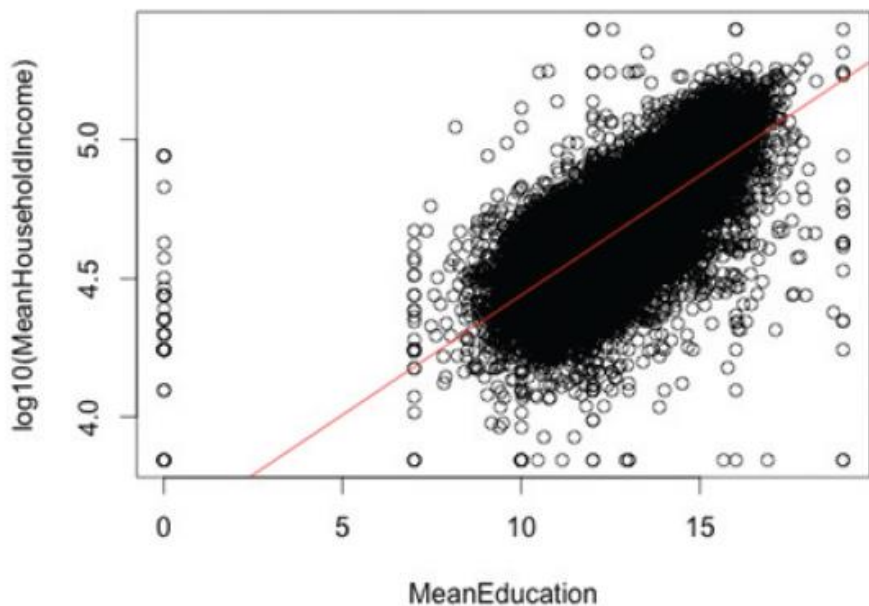
```
# plot the jittered scatterplot w/ boxplot  
# color-code points with zip codes  
# the outlier.size=0 prevents the boxplot from  
plotting the outlier  
ggplot(data=DF, aes(x=as.factor(Zip1),  
y=log10(MeanHouseholdIncome))) +  
  geom_point(aes(color=factor(Zip1)), alpha=0.2,  
position="jitter") +  
  geom_boxplot(outlier.size=0, alpha=0.1) +  
  guides(colour=FALSE) +  
  ggtitle ("Mean Household Income by Zip Code")
```

```
# simple boxplot
```

```
boxplot(log10(MeanHouseholdIncome) ~ Zip1, data=DF)  
title ("Mean Household Income by Zip Code")
```

# Exploratory Data Analysis

- Examining Multiple Variable (hexbinplot for large data)

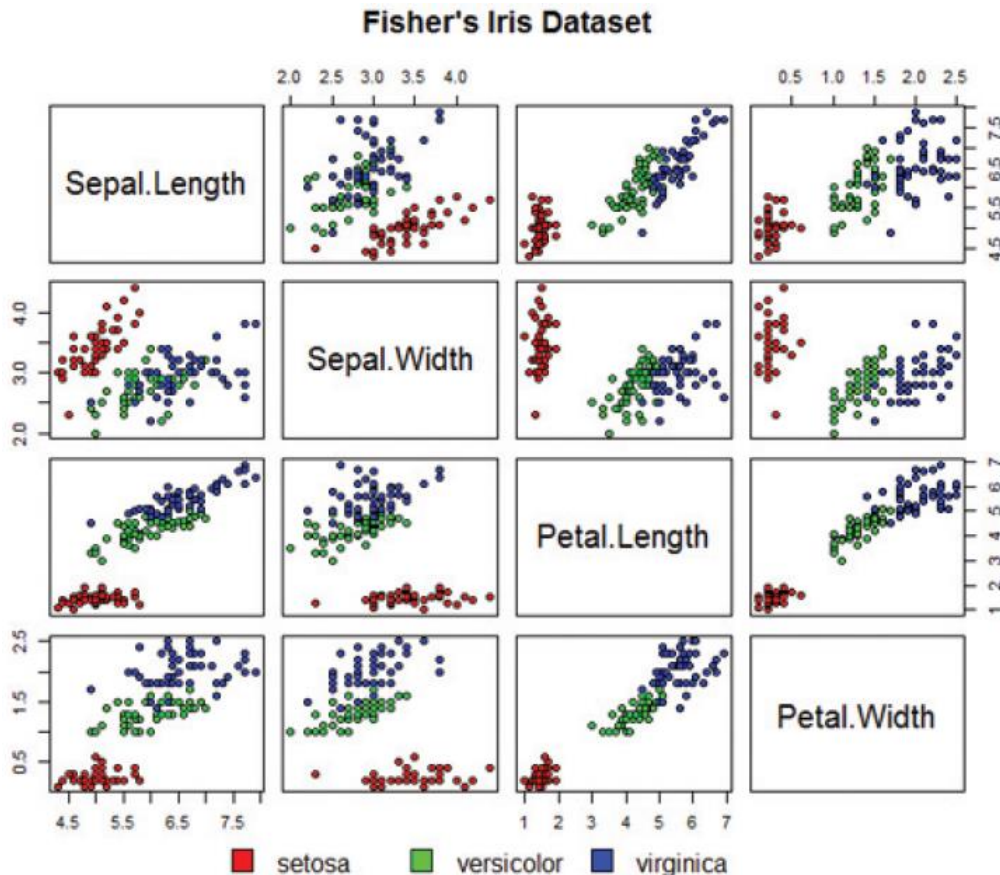


```
# plot the data points
plot(log10(MeanHouseholdIncome) ~ MeanEducation, data=DF)
# add a straight fitted line of the linear regression
abline(lm(log10(MeanHouseholdIncome) ~ MeanEducation, data=DF), col='red')
install.packages("hexbin")
library(hexbin)
# "g" adds the grid, "r" adds the regression line; sqrt transform on the count gives more dynamic range to the shading;
# inv provides the inverse transformation function of trans
hexbinplot(log10(MeanHouseholdIncome) ~ MeanEducation, data=DF, trans = sqrt, inv = function(x) x^2, type=c("g", "r"))
```



# Exploratory Data Analysis

- Examining Multiple Variable (scatterplot matrix)



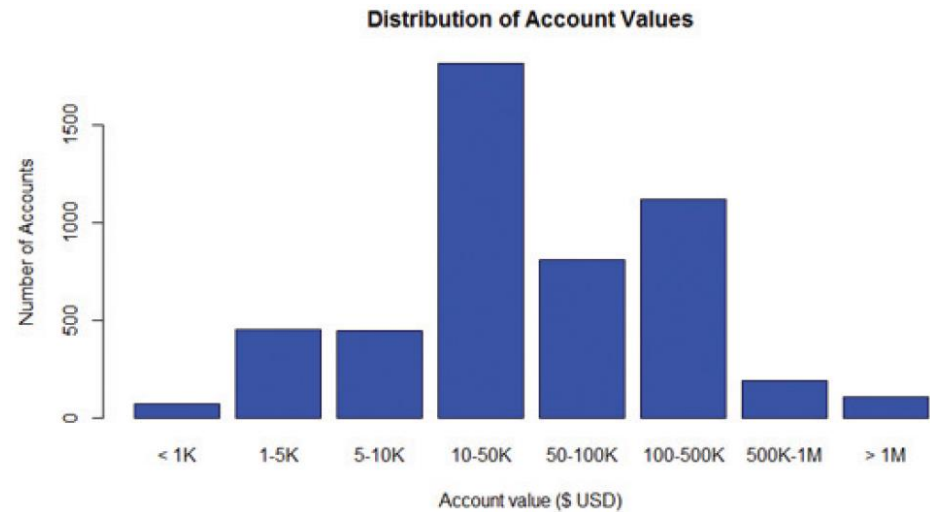
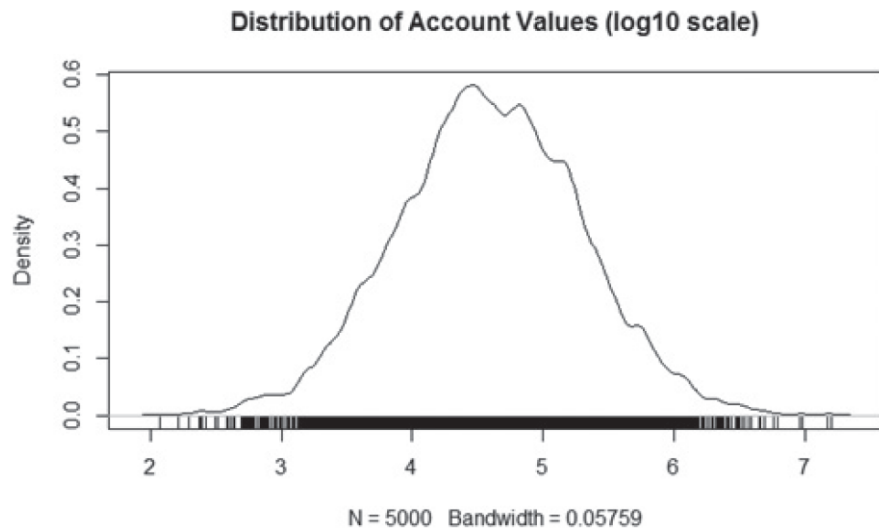
```
# define the colors
colors <- c("red", "green",
"blue")
```

```
# draw the plot matrix
pairs(iris[1:4], main = "Fisher's
Iris Dataset",
      pch = 21, bg =
colors[unclass(iris$Species)] )
# set graphical parameter to clip
plotting to the figure region
par(xpd = TRUE)
# add legend
legend(0.2, 0.02, horiz = TRUE,
as.vector(unique(iris$Species)),
      fill = colors, bty = "n")
```



# Exploratory Data Analysis

- Data Exploration Versus Presentation



Presenting the **same** data to **different** audience

# Statistical Methods for Evaluation

- **Statistics** is crucial because it may exist **throughout** the entire Data Analytics Lifecycle
  - Initial data exploration and data preparation
  - Model building and planning
    - Best input variables, predictability
  - Evaluation of the final models
    - Accuracy, better than guess or another one?
  - Assessment of the new models when deployed
    - Sound prediction? Have desired effect?

# Statistical Methods for Evaluation

- Hypothesis Testing
  - Form an **assertion** and test it with data
  - Common assumption (there is **no difference**)
  - Null hypothesis ( $H_0$ )
  - Alternative hypothesis ( $H_A$ )
- **Example**: identify the effect of **drug A** compared to **drug B** on patients
  - What are the  $H_0$  and  $H_A$ ?

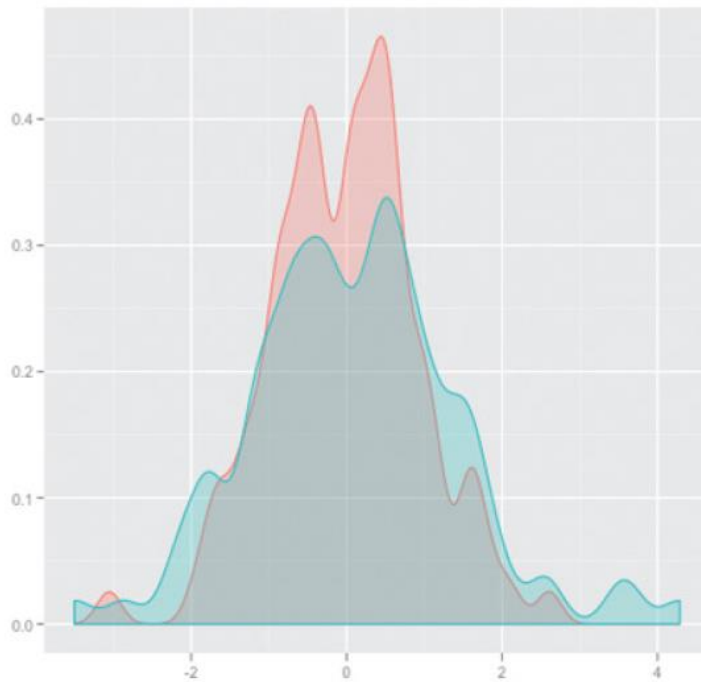
# Statistical Methods for Evaluation

- Hypothesis Testing
  - Clearly state Null and Alternative hypotheses
  - **Either** reject the null hypothesis in favour of the alternative **or** not reject the null hypothesis

Application	Null Hypothesis	Alternative Hypothesis
Accuracy Forecast	Model X <i>does not predict</i> better than the existing model.	Model X <i>predicts</i> better than the existing model.
Recommendation Engine	Algorithm Y <i>does not produce</i> better recommendations than the current algorithm being used.	Algorithm Y <i>produces</i> better recommendations than the current algorithm being used.
Regression Modeling	This variable <i>does not affect</i> the outcome because its coefficient is zero.	This variable <i>affects</i> outcome because its coefficient is not zero.

# Statistical Methods for Evaluation

- **Difference of Means** (A common hypothesis test)
  - Whether two populations are different?
  - Compare their means based on sampled data



– What are  $H_0$  and  $H_A$  ?

# Statistical Methods for Evaluation

- Student's *t*-test
  - Assume that distributions of the two populations have **equal but unknown variance**
  - If each population is **normally** distributed with the **same** mean and with the **same** variance, then

$$T = \frac{\bar{X}_1 - \bar{X}_2}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}$$

$$S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2}$$

T (the *t*-statistic) follows a *t*-distribution with  $(n_1 + n_2 - 2)$  degree of freedom

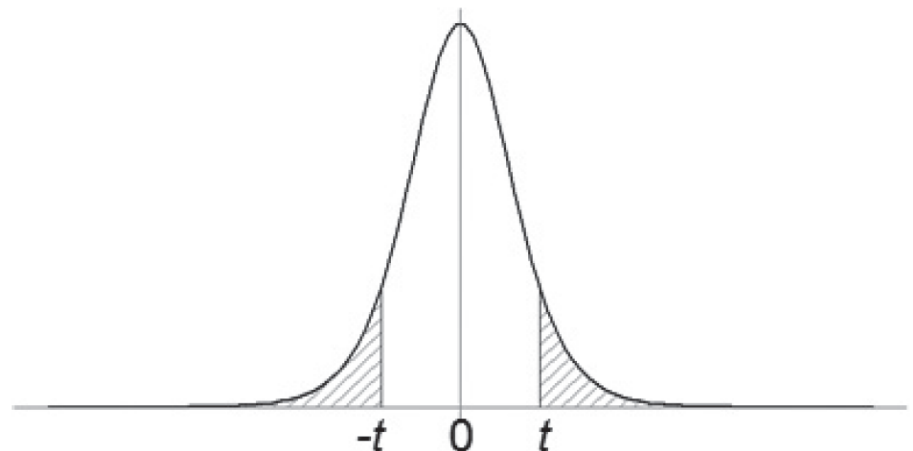
# Statistical Methods for Evaluation

- Student's *t*-test

- If the observed *T* is far from zero such that the probability of observing such a value of *T* is unlikely, one would reject the null hypothesis

$$T = \frac{\bar{X}_1 - \bar{X}_2}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}$$

$$S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2}$$

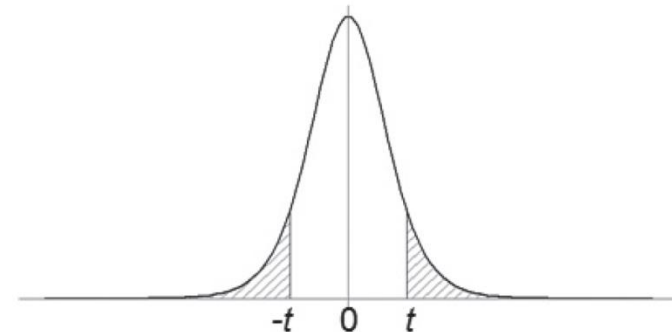


# Statistical Methods for Evaluation

- Student's *t*-test

- Significance level of the test ( $\alpha$ ): the probability of **rejecting** the null hypothesis, when the null hypothesis is **actually TRUE**
- So, what does it mean by setting  $\alpha = 0.05$ ?
- Find  $T^*$  such that  $P(|T| \geq T^*) = \alpha$
- Reject  $H_0$  if  $|T| \geq T^*$

$$T = \frac{\bar{X}_1 - \bar{X}_2}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}$$
$$S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2}$$





# Statistical Methods for Evaluation

- Student's *t*-test (an example)

```
# generate random observations from the two populations
x <- rnorm(10, mean=100, sd=5)      # normal distribution centered at 100
y <- rnorm(20, mean=105, sd=5)     # normal distribution centered at 105

t.test(x, y, var.equal=TRUE)       # run the Student's t-test

Two Sample t-test

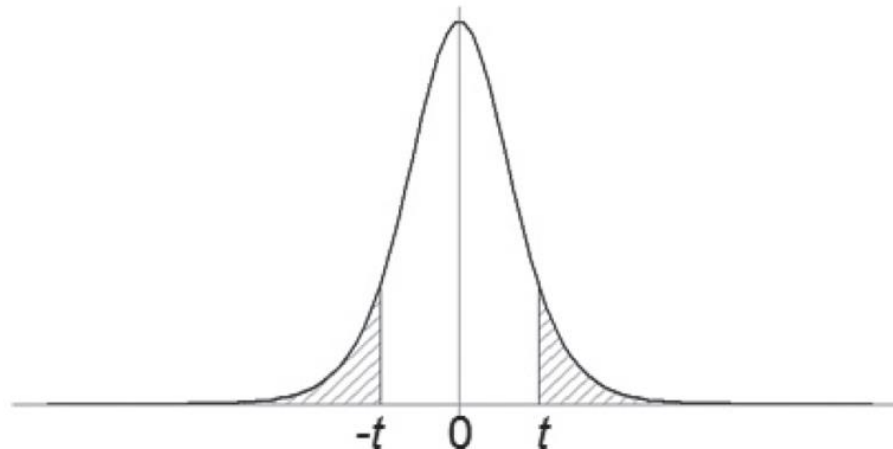
data:  x and y
t = -1.7828, df = 28, p-value = 0.08547
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -6.1611557  0.4271893
sample estimates:
 mean of x mean of y
102.2136  105.0806
```

# Statistical Methods for Evaluation

- Student's *t*-test (an example)

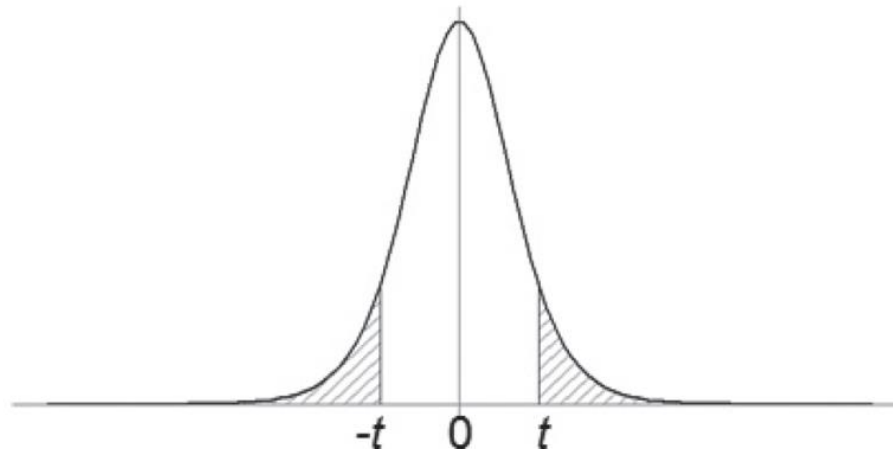
```
# obtain t value for a two-sided test at a 0.05 significance level  
qt(p=0.05/2, df=28, lower.tail= FALSE)  
2.048407
```

- Shall we **reject or accept** the null hypothesis?
- What does the “**two-sided** test” mean?



# Statistical Methods for Evaluation

- Student's *t*-test (an example)
  - What does the “p-value” mean?  
 $t = -1.7828, df = 28, p\text{-value} = 0.08547$
  - The sum of  $P(T \leq -t)$  and  $P(T \geq t)$
  - p-value offers the probability of observing  $|T| \geq t$  given the null hypothesis is TRUE



# Statistical Methods for Evaluation

- Student's *t*-test (an example)

- What is the “95 percent confidence interval”?

95 percent confidence interval:

-6.1611557    0.4271893

- A confidence level is an interval estimate of a population parameter based on sample data

- The above “95 percent confidence interval” straddles the TRUE value of the difference of the population means 95% of the time

# Statistical Methods for Evaluation

- Welch's *t*-test
  - Shall be used when the equal population variance assumption is NOT justified
  - It uses the sample variance for each population instead of the pooled sample variance
  - Still assume two populations are normal with the same mean

$$T_{welch} = \frac{\bar{X}_1 - \bar{X}_2}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}}$$

# Statistical Methods for Evaluation

- Welch's  $t$ -test

```
t.test(x, y, var.equal=FALSE)           # run the Welch's t-test
```

```
Welch Two Sample t-test
```

```
data:  x and y
```

```
t = -1.6596, df = 15.118, p-value = 0.1176
```

```
alternative hypothesis: true difference in means is not equal to 0
```

```
95 percent confidence interval:
```

```
 -6.546629  0.812663
```

```
sample estimates:
```

```
 mean of x mean of y
```

```
102.2136  105.0806
```

# Statistical Methods for Evaluation

- Wilcoxon Rank-Sum Test
  - What if the two populations are **not normal**?
- Parametric test
  - **Makes assumptions** about the population distributions from which the samples are drawn
- Nonparametric test
  - Shall be used if the populations **cannot** be assumed (or transformed) to be **normal**

# Statistical Methods for Evaluation

- Wilcoxon Rank-Sum Test
  - A nonparametric test to check whether two populations are identically distributed
  - It uses “ranks” instead of numerical outcomes to avoid specific assumption about the distribution
- How to conduct the test
  - Rank two samples as if they are from one group
  - Sum assigned ranks for one population's sample
  - Determine the significance of the rank-sums



# Statistical Methods for Evaluation

- Wilcoxon Rank-Sum Test

```
wilcox.test(x, y, conf.int = TRUE)
```

```
Wilcoxon rank sum test
```

```
data: x and y
```

```
W = 55, p-value = 0.04903
```

```
alternative hypothesis: true location shift is not equal to 0
```

```
95 percent confidence interval:
```

```
-6.2596774 -0.1240618
```

```
sample estimates:
```

```
difference in location
```

```
-3.417658
```

**p-value:** the probability of the rank-sums of this magnitude being observed assuming that the population distributions are identical

# Statistical Methods for Evaluation

- Type I and Type II Errors

- Type I error: the **rejection** of the null hypothesis when the null hypothesis is **TRUE**
- The probability of type I error is denoted by  $\alpha$
- Type II error: the **acceptance** of the null hypothesis when the null hypothesis is **FALSE**
- The probability of type II error is denoted by  $\beta$

- Power (statistical power)

- The probability of **correctly rejecting** the null hypothesis ( $1 - \beta$ )

# Statistical Methods for Evaluation

- ANOVA (Analysis of Variance)
  - What if there are more than two populations?
  - Multiple  $t$ -test may not perform well now
- A generalization of the hypothesis testing
  - ANOVA tests if any of the population means differ from the other population means
  - Each population is assumed to be normal and have the same variance

# Statistical Methods for Evaluation

- ANOVA (Analysis of Variance)

$$H_0: \mu_1 = \mu_2 = \dots = \mu_n$$

$$H_A: \mu_i \neq \mu_j \text{ for at least one pair of } i, j$$

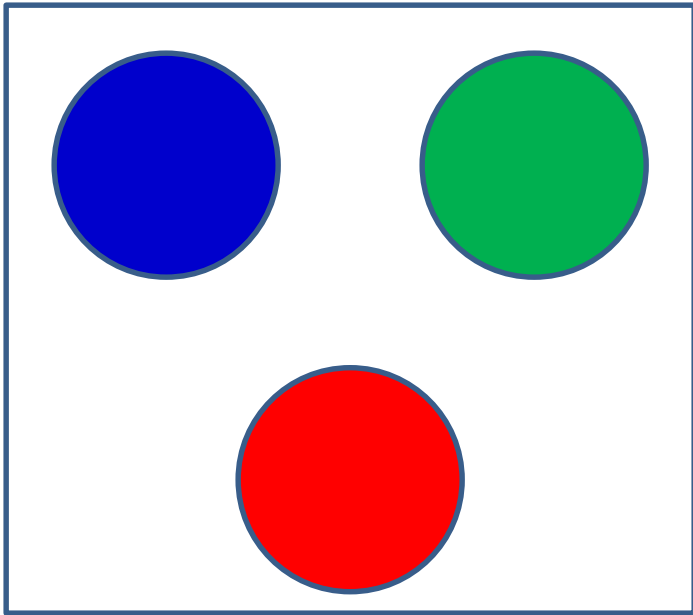
- Compute **F-test statistic**

- Between-groups mean sum of squares
- Within-groups mean sum of squares

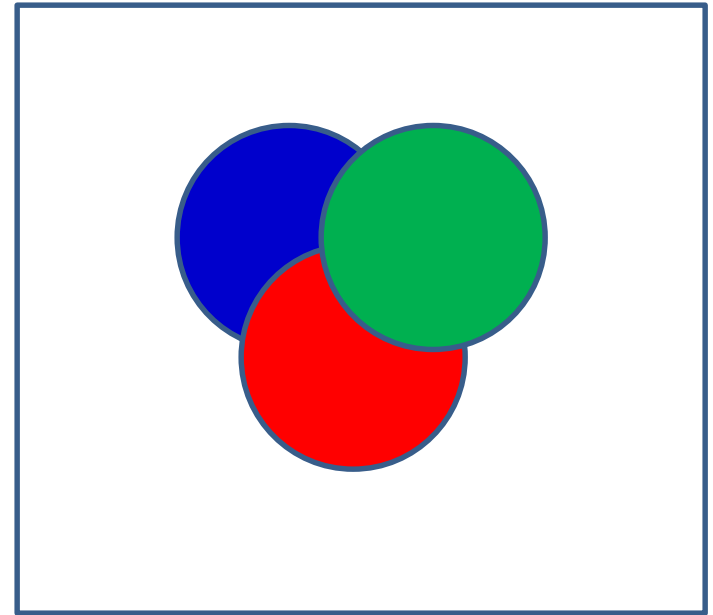
$$S_B^2 = \frac{1}{k-1} \sum_{i=1}^k n_i \cdot (\bar{x}_i - \bar{x}_0)^2 \quad S_W^2 = \frac{1}{n-k} \sum_{i=1}^k \sum_{j=1}^{n_i} (x_{ij} - \bar{x}_i)^2$$

# Statistical Methods for Evaluation

- ANOVA (Analysis of Variance)



$$F = \frac{S_B^2}{S_W^2}$$



$$S_B^2 = \frac{1}{k-1} \sum_{i=1}^k n_i \cdot (\bar{x}_i - \bar{x}_0)^2$$

$$S_W^2 = \frac{1}{n-k} \sum_{i=1}^k \sum_{j=1}^{n_i} (x_{ij} - \bar{x}_i)^2$$

# Statistical Methods for Evaluation

- ANOVA (Analysis of Variance)
  - Measures how **different** the means are **relative to** the **variability** within each group
  - The **larger** the  $F$ -test statistic, the **greater** the likelihood that the difference of means are due to something **other than chance** alone
  - The  **$F$ -test** statistic follows an  **$F$ -distribution**

$$F = \frac{S_B^2}{S_W^2}$$

# Statistical Methods for Evaluation

- ANOVA (Analysis of Variance)

```
# fit ANOVA test
model <- aov(purchase_amt ~ offers, data=offertest)

summary(model)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
offers	2	225222	112611	130.6	<2e-16 ***
Residuals	497	428470	862		

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

- Shall we **accept or reject** the null hypothesis?

# Statistical Methods for Evaluation

- ANOVA (Analysis of Variance)
  - Additional tests for each pair of groups
  - Tukey's Honest Significant Difference (HSD)

```
TukeyHSD(model)
```

```
Tukey multiple comparisons of means  
95% family-wise confidence level
```

```
Fit: aov(formula = purchase_amt ~ offers, data = offertest)
```

```
$offers
```

		diff	lwr	upr	p adj
offer1-nopromo	40.961437	33.4638483	48.45903	0.0000000	
offer2-nopromo	48.120286	40.5189446	55.72163	0.0000000	
offer2-offer1	7.158849	-0.4315769	14.74928	0.0692895	



# Recap: Data Analytic Methods Using R

- Introduction to R
  - R, RStudio, Data I/O, Attribute and Data Types
  - Descriptive statistics
- Exploratory Data Analysis
  - Visualization before analysis
  - Visualizing single or multiple variables
- Statistical Methods for Evaluation
  - Hypothesis Testing, ANOVA

