Introduction to R

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Topics to be Covered

- R Language Basics
- Vectors and Selections
- Matrices and Data Frames
- Writing and Reading Data
- Plotting
- Using Packages
 - Installing
 - Loading
 - Viewing Help

Basics

 From: https://github.com/Data-Camp/introduction_to_R/blob/master/chapter1.Rmd

Simple calculations

```
2 + 2
[1] 4
```

Commenting Code

```
# This is a comment
2 + 2
[1] 4
# Addition
5 + 5
[1] 10
# Subtraction
5 - 5
[1] 0
# Multiplication
[1] 15
# Division
(5 + 5) / 2
[1] 5
```

Variable Assignments

```
my_variable <- 4
my_variable
```

```
[1] 4
```

Basic Data Types

```
# What is the answer to the universe?
my_numeric <- 42

# The quotation marks indicate that the variable is of type character
my_character <- "some text"

# Change the value of my_logical
my_logical <- TRUE</pre>
```

Help

?mean

Vectors

• From: https://github.com/Data-Camp/introduction_to_R/blob/master/chapter2.Rmd

Creating a vector

```
numeric_vector <- c(1, 2, 3)
character_vector <- c("a", "b", "c")
boolean_vector <- c(TRUE, FALSE, FALSE)</pre>
```

Selection by index

```
numeric_vector[c(1, 3)]
[1] 1 3
```

Selection by logical

```
my_variable <- 2
result <- numeric_vector[numeric_vector > my_variable]
result
```

```
[1] 3
```

Matrices

 From: https://github.com/Data-Camp/introduction_to_R/blob/master/chapter3.Rmd

Matrices from vectors

```
first_row <- c(6,8,7,9,9,10)
second_row <- c(6,8,7,5,9,6)
third_row <- c(5,4,6,6,7,8)
fourth_row <- c(4,5,3,4,6,8)

# Combine multiple vectors to form a matrix
theater <- rbind(first_row, second_row,
third_row, fourth_row)
row_scores <- rowSums(theater)
scores <- cbind(theater, row_scores)</pre>
```

Naming a Matrix

```
rownames(scores) <- c("row1", "row2", "row3", "row4")
colnames(scores) <- c("col1", "col2", "col3", "col4",
"col5", "col6", "total")
scores</pre>
```

```
      col1 col2 col3 col4 col5 col6 total

      row1
      6
      8
      7
      9
      9
      10
      49

      row2
      6
      8
      7
      5
      9
      6
      41

      row3
      5
      4
      6
      6
      7
      8
      36

      row4
      4
      5
      3
      4
      6
      8
      30
```

Size of Matrix

```
ncol(scores)
[1] 7
nrow(scores)
[1] 4
dim(scores)
[1] 4 7
```

Selecting Elements

Select rows and columns

```
i <- 1
j <- 1
scores[i,]
 col1 col2 col3 col4 col5 col6 total
scores[,j]
row1 row2 row3 row4
scores[i,j]
[1] 6
```

Data Frames

data(iris)
head(iris)

```
Sepal.Length Sepal.Width Petal.Length Petal.Width
Species
           5.1
                       3.5
                                                 0.2
                                     1.4
setosa
                       3.0
           4.9
                                                 0.2
                                     1.4
setosa
           4.7
                       3.2
                                     1.3
                                                 0.2
setosa
                                     1.5
           4.6
                       3.1
                                                 0.2
setosa
           5.0
                                                 0.2
                       3.6
                                     1.4
setosa
           5.4
                       3.9
                                     1.7
                                                 0.4
setosa
```

Rename data.frame Columns

```
numeric_vector <- c(1, 2, 3)
character_vector <- c("a", "b", "c")
boolean_vector <- c(TRUE, FALSE, FALSE)

df <- data.frame(numbers=numeric_vector,
characters=character_vector, boolean=boolean_vector)

df</pre>
```

Selecting Columns by Name

```
iris[,"Sepal.Length"]
[137] 6.3 6.4 6.0 6.9 6.7 6.9 5.8 6.8 6.7 6.7 6.3 6.5 6.2 5.9
iris$Sepal.Length
[137] 6.3 6.4 6.0 6.9 6.7 6.9 5.8 6.8 6.7 6.7 6.3 6.5 6.2 5.9
```

Exporting Data

Writing files

```
write.table(iris, file="iris.txt", sep="\t",
row.names=TRUE, col.names=TRUE, quote=FALSE)
```

Reading files

```
df <- read.table("iris.txt", sep="\t",
header=TRUE)</pre>
```

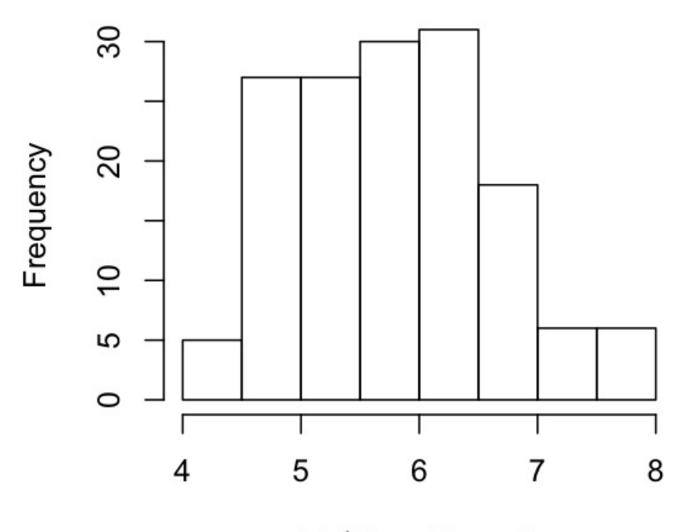
Plotting

 From: https://github.com/Data-Camp/introduction_to_R/blob/master/chapter7.Rmd

Histogram

hist(iris\$Sepal.Length)

Histogram of iris\$Sepal.Length

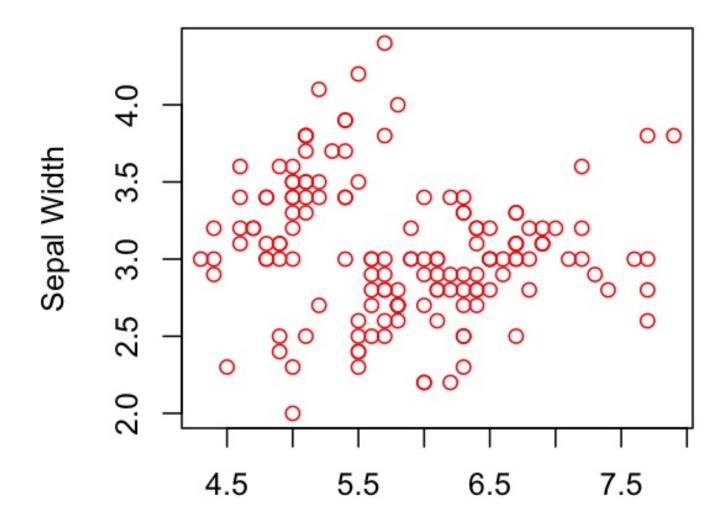


iris\$Sepal.Length

Scatterplot

```
plot(x=iris$Sepal.Length,
    y=iris$Sepal.Width,
    main = "Sepal Length versus Sepal Width",
    xlab = "Sepal Length",
    ylab = "Sepal Width",
    col = "red")
```

Sepal Length versus Sepal Width



R Packages

• From:

http://www.jkarreth.net/files/RPOS517_Day1_IntroR.pdf

Install packages from repositories

 NOTE: These commands are commented out since these packages are already installed

```
# From CRAN (for general packages)
install.packages("httr")

# From Bioconductor (for biology-related packages)
source("https://bioconductor.org/biocLite.R")
biocLite("rcellminer")
```

Load Package

```
library(rcellminer)

# Check if package was loaded
sessionInfo()
```

Package Help

```
help(package="rcellminer")
```

Getting Help

- Stack Overflow
 - http://stackoverflow.com/
- Cross-Validated Stats Exchange
 - Part of Stack Overflow
 - http://stats.stackexchange.com/
- Biostars
 - https://www.biostars.org