### Introduction to R

**ALY-6050** 

### **Skills**

- Distinguish the differences between R and Excel for data analysis and recognize the advantages of R
- Navigate the RStudio working environment, using syntax and opening files
- Understand the basic R data types and data objects, syntax of creation and expression and typical applications
- Differentiate between and apply basic data manipulation, such as indexing, subsetting and reshaping data in R

### Introduction to R (I)

- R is an open-source statistical programming language
- Supported by CRAN, the Comprehensive R Archive Network
  - Core code is distributed to users for free
  - CRAN maintains the source code and documentation
  - User community can contribute to R by writing packages to increase the functionality of R
- Free integrated development environment (IDE), RStudio



### Introduction R (II)

- R is easy to learn
- R is most useful for statistical analysis, data manipulation, data analytics and creating customized, high-end graphics and visualizations
- Many resources from the community:
  - http://www.r-bloggers.com/
  - https://www.r-project.org/

### Advantages of R over Excel (I)

- Excel is imited to relatively small data sets, 1,048,576 rows by 16.384 columns, and large .xlsx files can be very slow to navigate and edit
  - R addresses these limitations in two ways:
    - Scripting and setting up plotting and plot formatting can be performed without opening file storing the data, large dataset is not in RAM during scripting
    - Data can be allocated to up to 8TB of RAM, and there are R packages available that allow users to store data in hard drives and analyze it in chunks

### Advantages of R over Excel (II)

- Excel has limited functionality for statistical computing, whereas R continues to grow in this regard through package submissions from the user community
- Difficult to reproduce analysis in Excel:
  - Copying and pasting is prone to human error
  - Raw data is often next to cells requiring data entry or modification, equations are easily, accidentally overwritten
  - Using R scripts, the same analysis can be performed on multiple datasets by changing a single input
  - Similarly, the same graphic can easily be created for more than one data set using R
  - Code can be commented, tested and versioned

### Correct tool for the task

#### Excel

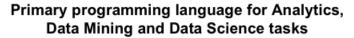
- Simple descriptive or inferential statistics on a relatively small dataset
- Collaborating on a project with people who are not data analysts (and don't know R), so long as the formulas and functions are simple

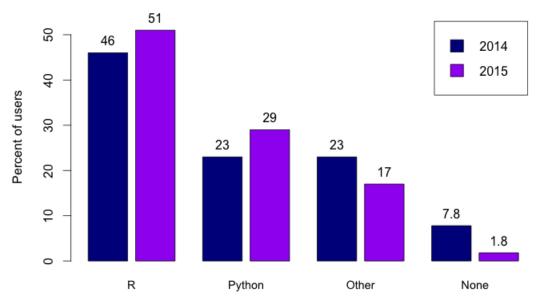
#### • R

- Large datasets
- Advanced statistical computations
- Same analysis on more than one data set
- Complex data visualization
- Analyses which must be reproducible, require advanced functionality or plotting capabilities

### R vs. other languages (II)

- R has similar advantages over Excel to other languages
- R is popular, growing and relatively easy to learn
- It is also free and has a strong support community



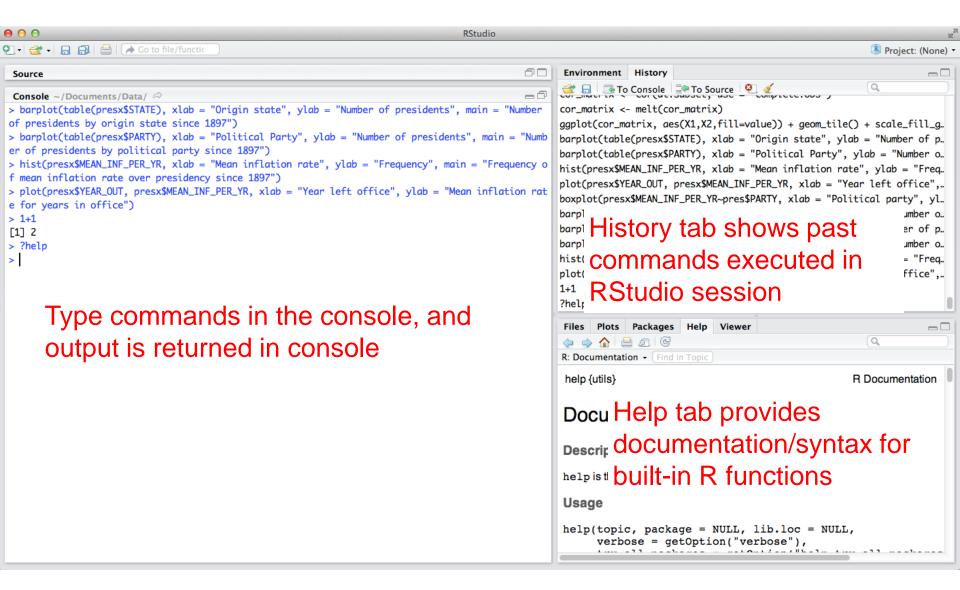


Note: "Other" includes MATLAB, Java, SAS, Scala, etc.

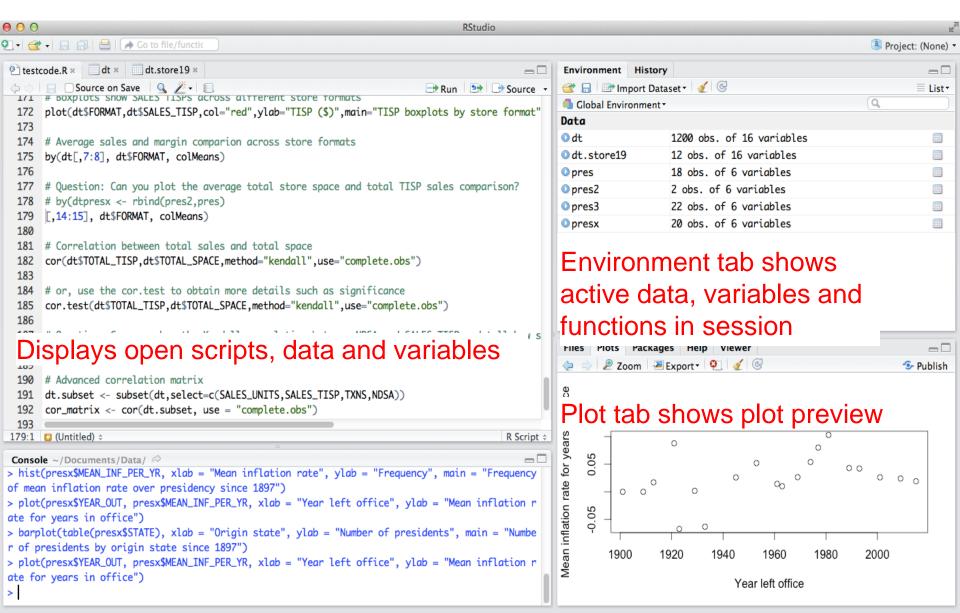
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# RStudio GUI (I)



### RStudio GUI (II)



### RStudio GUI (III)

 In this presentation, as well as in RStudio, we will use the following font and color scheme:

Input to command line
Printed in console
Error or warning
Commented code

- Check out and follow the google style guide for R:
  - Makes code easier to read/share with others
  - https://google.github.io/styleguide/Rguide.xml

### Basic syntax guidelines for R

- # begins a line of comment
- R is case sensitive: x is not the same as x
- arguments go between parenthesis (and are separated by commas), for example help(lapply)
- multiple arguments are separated with commas
- <- is the assignment operator, analogous to equals sign</li>

### Useful functions

- print() displays argument value(s) in console
- install.packages() installs (attaches) add-on packages to the basic installation of R, note: packages remain installed until uninstalled
- installed.packages() with empty parentheses, lists packages already installed
- library() opens library of functions within a package, note: empty parenthesis lists libraries/packages available for opening in an R session in upper-left window; note: libraries are closed when R session is closed
- search() with no arguments, lists open packages in console

# Getting help in R

- help() or ? opens help tab for argument, shows syntax options
- example() runs all the R code from the Examples section of the help documentation for argument
- browseVignettes() lists available longer form documentation for argument in a web browser
- RSiteSearch() searches for key words or phrases in help pages, vignettes or task views, using http://search.rproject.org and views them in a web browser
- apropos() returns the names of all objects matching argument

# Working directory (I)

- Working directory is the folder currently in use in R
- Location from which files are opened and location to which files are saved

 Important to know and be able to change so that files are read and written to known location

### Working directory (II)

- getwd() get working directory, returns current working directory in console
- setwd() set working directory, assigns working directory
  - setwd("...") up one folder
  - setwd("../..") up two folders
  - setwd("Folder1/Folder2/Folder3") –
     specifies path within current working directory
- Create a folder on your computer for R projects

### Check your knowledge (VIII)

 Create a folder on your computer for R projects, and set it to your working directory, example solution with extra navigation:

```
getwd()
[1] "/Users/YourName/Documents/Data"
setwd("..")
getwd()
[1] "/Users/YourName/Documents"
setwd("../..")
getwd()
[1] "/Users"
setwd("YourName/Documents/RData")
getwd()
[1] "/Users/YourName/Documents/RData"
```

# Reference - functions to read and write files in R (I)

• read.table() - default delimiter is white space

read.csv() – default delimiter is a comma

 read.csv2() – default delimiter is a semicolon, and commas are read as decimal points

read.delim() – default delimiter is a tab

 read.delim2() – default delimiter is a tab, and commas are read as decimal points

# Reference - functions to read and write files in R (I)

 read.table(file.choose()) – opens window to browse for file

 See help for: write.table(), write.csv() and write.csv2() for information about saving data in R to file

# Reading Excel files in R with XLConnect, xlsx and gdata (I)

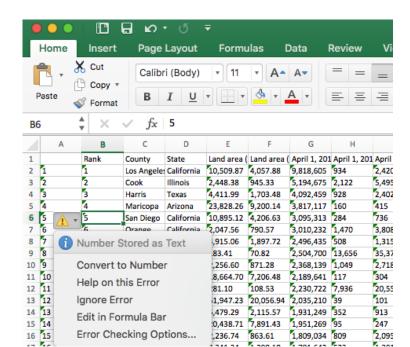
- Many ways to read worksheets and workbooks from Excelinto R:
  - readWorksheetFromFile() reads data from worksheets in Excel into data frame (XLConnect package)
  - loadworkbook() reads in entire Excel workbook (XLConnect package)
  - read.xlsx() reads data from worksheets in Excel into data frame (xlsx package)

# Reading Excel files in R with XLConnect, xlsx and gdata (II)

- Many ways to read worksheets and workbooks from Excelinto R:
  - read.xlsx2() similar to read.xlsx() but faster for large datasets (xlsx package)
  - loadworkbook() reads in entire Excel workbook (xlsx package)
  - can write to Excel files with write.xlsx() and write.xlsx2() (xlsx package)
  - read.xls() read Excel file into a data frame using Perl (gdata package)

# A note about data types in R

- We must be conscious of and intentional about data types in R
  - Soon we will learn how to specify, check and change data types in R



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# Data types in R

Atomic Type	Example	typeof()	mode()	class()
character	"hello"	character	character	character
real	1.34	double	numeric	numeric
integer	3L	integer	numeric	integer
complex	3.0 - 4.3i	complex	complex	complex
logical	TRUE or T	logical	logical	logical

### Check your knowledge (I)

Match the output of class(x) with the corresponding command

$$2. x < -50L$$

$$3. x < -50$$

- A. integer
- B. numeric
- c. logical
- D. character
- E. complex

# Data objects (I)

- Data objects store groups of elements in R, and this classification defines what type of operations can be performed on these elements
- Vectors store elements of the same data type in a single column or row
- Matrices store elements of the same data type in multiple columns or rows
- Lists store different types of data objects in the form of a vector (single column)
- Factors store integer elements to represent discrete categories, known in R as levels
- Data frames similar to a single sheet in Excel
  - Store data in a tabular format
  - Each column can be a different data type

### Check your knowledge (VIII)

Match the data types to the output, using each exactly once

- 1. Vector
- Matrix
- 3. List
- 4. Factor
- 5. Data frame

```
A. [1] hello hello welcome Levels: hello welcome
```

```
B [1] "hello" "welcome" "goodbye"
```

```
C. hello...welcome.. c.1..2.

1 hello 1
2 welcome 2
```

```
D. [[1]]
[1] "hello" "hello"

[[2]]
[1] hello welcome
Levels: hello welcome
```

```
[,1] [,2]
[1,] "hello" "goodbye"
[2,] "welcome" "hello"
```

### Basic arithmetic in R

- R can perform all of the basic arithmetic of a scientific calculator, order of operations (PEMDAS) applies
- In addition to using R for statistics, may need to add, subtract or multiple columns, much like in Excel

rowMeans() and colMeans() – computes the arithmetic mean for rows or columns of a matrix or data frame (2D data object)

rowSums() and colsums() – computes the numeric sum of rows or columns of a matrix or data frame (2D data object)

sum() and prod() - returns the sum of all values in arguments and the product of all values in arguments, respectively

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### Subsetting data in R (I)

- Elements within data objects in R are indexed in the format [row, column]
- A particular element or set of elements can be returned from a data object using the name of the data object, followed by the square bracket with the desired indices
- Consider a 5x4 data frame, df:
  - df[1,] returns first row of df
  - df[,2] returns second column of df
  - df[1,2] returns single element in the first row and second column of df

### Subsetting data in R (II)

- Because the indices of data objects are integers, a colon, :, is used to specify more than one row
  - Note: the colon returns all integers between the integers on either side of the colon, inclusive
- Consider a 5x4 data frame, df:
  - df[3:5,] returns third through fifth rows of df
  - df[,1:3] returns first through third columns of df
  - df[3:5,1:3] returns nine elements of df spanning the third through fifth rows and the first through third columns

### Subsetting data in R (III)

- Elements within data objects in R are indexed in the format [row, column]
- Elements with only one dimension, such as vectors can be subsetted without the colon
- Consider vector of length 6, a:
  - a[1] returns first element of a
  - a[3:5] returns third through fifth elements of a

### Subsetting data in R (IV)

- Elements within data objects in R are indexed in the format [row, column]
- Exception for *lists*: list elements are subsetted with two sets of square brackets, followed by the subsetting rules for other *vectors*, *matrices*, or *data frames*
- Consider *list* of 6 3x3 *matrices*, b:
  - b[[3]][2,] returns second row from the third 3x3 matrix in b

### Subsetting data in R (V)

- \$ is important for accessing data in data frames
  - data frames have row and column names, they are automatically generated and can be reassigned
  - Reassign row and column names to something useful
  - Access rows and columns using the name of the data frame, followed by the dollar sign, \$, followed by the name of the row or column if interest

### Check your knowledge (VIII)

```
Matching, x is a data frame of A. x[2:4,] dimensions 5x5
```

- Element in Row 2, Column
   of x
- 2. Row 2 of **x**
- 3. Column 2 of x B
- 4. Rows 2, 3 and 4 of x A
- 5. Columns 2, 3 and 4 of x E
- 6. Columns 2 and 4 of x
- 7. Rows 2 and 4 of x

```
B. x[,2]

C. x[seq(2,4,2),]

D. x[2,2]

E. x[,c(2,3,4)]

F. x[2,]

G. x[,c(2,4)]
```

### Useful querying functions in R (I)

attributes() - returns information about shape, size
and type of data object in argument

dim() - returns dimensions of argument

length() - returns length of argument, number of columns for a matrix or data frame

str() - returns structure of argument

head() - returns first rows of argument

### Useful querying functions in R (II)

tail() - returns final rows of argument

colnames()/names() - returns or assigns column
names of argument

rownames () – returns or assigns row names of argument

unique() - returns unique elements within argument, can be applied to factors as well as character vectors

## Useful querying functions in R (III)

duplicated() - returns logical vector indicating which
elements are duplicated in a given vector or data frame

**summary()** – returns summary descriptive statistics of argument for numerical data, and tabulates number of elements in each factor level of argument for categorical data

# Appending, modifying and removing data in R (I)

subset() - returns a subset of data from argument based
on a selection argument

merge() – returns a merged data frame from two input data frame arguments, merged by common columns or row names

cbind()/rbind() - used to combine a sequence of
vectors, matrices or data frames, into a single large data
object

- cbind() combines by columns
- rbind() combines by rows

# Which functions for subsetting data in R (I)

which functions return indices of elements that match some argument

which() - returns indices of the elements elements that
match the argument

which.min() - returns indices of element that has the
minimum value in the argument

which.max() - returns indices of element that has the
maximum value in the argument

### Sorting data in R (I)

- Useful to sort a data set by some variable
  - Combine with head() or tail() to see the smallest or largest values

sort() - returns sorted argument, use for datasets with a single column

order() - returns sorted argument, use for datasets with multiple columns

# Some of the many statistics functions in R (I)

- There are many descriptive and inferential statistics functions in R
  - Below are some simple functions which can be used in combination with more advanced functions to calculate statistics for numeric data
  - Can operate every column, row or groups of the data split by a categorical variable

mean() – returns mean of argument

median() – returns median of argument

# Some of the many statistics functions in R (II)

 There are many descriptive and inferential statistics functions in R

var() – returns sample variance of argument

cov() - returns covariance of argument(s)

cor() - returns correlation coefficient of argument(s)

### Apply functions in R (I)

- apply() functions, which are built-in to R, apply a formula or built-in R function to multiple elements within a data object
  - Select between apply functions based on type of data objects we have for our dataset and type of data object we want as the output

apply() - applies formula or function to a vector or matrix
argument, across some dimension (rows or columns) and
returns output in a matrix or vector

lapply() - applies formula or function to a list, vector or matrix argument and returns output in a list

### Apply functions in R (II)

sapply() - applies formula or function to a list, vector or matrix argument and returns output in a vector or matrix

tapply() – applies formula or function to categorical subsets of data in a vector or matrix argument and returns output in a list

# Built-in functions for processing data in R (I)

with() - applies function to selected data in data frame,
does not affect data frame

within() - applies function to selected data in data
frame, appends results to data frame

by() – applies function to subsets of data specified by factor or list of factors

aggregate() - splits data into subsets by factor, returns
descriptive statistics

#### NA and NaN (I)

Nonsensical coercion brings us special and missing values in R

- NA is a missing value
  - NA has a data type, such as integer or logical, but no specific value
- NaN is defined as zero divided by zero, stands for "not a number"
- NaN is NA, but NA is not NaN

### NA and NaN (II)

is.na() - returns a logical element; TRUE if argument is NA, FALSE if argument is not NA

is.nan() - returns a logical element; TRUE if argument is NaN, FALSE if argument is not NaN

#### NA and NaN (III)

complete.cases() - returns logical element for each
row in matrix/data frame or each element in vector, TRUE if
no NA or NaN values are present, FALSE if at least one NA
or NaN is present

- Useful for calculating statistics for large samples with missing values
- Also check out anyNA()
- Note: many built-in R functions will return NA if any of the elements are NA, such as mean()
  - Some of these functions accept additional arguments, such as na.rm = T, to remove NA values in calculations