Before the workshop

Download Data Files: https://github.com/julipetal/Intro_to_R

Introduction to R

Part I

Part I

Please interrupt with questions

Basic Introduction

Good Coding Practice

Data Handling

Basic Data Visualization

Basic Introduction

What is R?

Free software environment for statistical computing and graphics

Runs on most UNIX platforms, Windows and MacOS

Website: http://www.r-project.org/

CRAN - Comprehensive R Archive Network

Popular in academia, and growing software in industry

Frequently releases updated versions with funny names (like 'Bug in Your Hair'), check every 6 month or so to ensure you are using an up-to-date version

Tons of different libraries (of which some do the same thing)

R code

Any R-script (fileName.r) can be opened in an editor of your choice

 For windows: Notepad++ is good because it highlights different attributes of the code, unlike Notepad

Copy and past R-code from scripts into the R window, also called R console

Directly write your code into the R window

Using UP and DOWN arrows will reveal lines of code that have been typed before within the current R history

To execute a line of code press enter

Help!!!

- > help.start()
- > help(<functionName>)

Replace < functionName > with which

> ?<functionName>

Again try which and then for

- > help.search("<functionName>")
- > example(<functionName>)

- > demo()
- > demo(graphics)

Easy examples and explanations for basic R functions

http://www.statmethods.net/

RStudio as an editor

https://www.rstudio.com/products/rstudio/download/#download

Build in Functions

Standard math operations and functions

```
o +,-,*,/
o exp(), log(), sin(), ...
```

Standard statistics

```
o mean(), sd(), density()
```

Assigning numbers to variables

```
0 > X = 65

0 > X = 43

0 > X + X
```

NOTE: R is case sensitive

You will also see <- instead of an equal sign, it does the same thing

$$\circ$$
 > y <- 2

R Reference Card: https://cran.r-project.org/doc/contrib/Short-refcard.pdf

Closing an R Session/Console

To close a session

> q()

R will promote you if you want to save your session

If you kept a script and saved important data to file, there is no need to save the session.

Otherwise you might want to say yes. If you do, an .Rhistory file will be created. This will overwrite any previously created file if not renamed. The .Rhistory file includes all lines of code without output and also will include nonfunctional code.

Good Coding Tip: Instead of saving the R session, keep a clean script file with comments.

Good Coding Practice

Good Practice

Save work and code by using scripts

Keep these scripts clean

- Avoid saving code that did not work or produced unwanted results
- Avoid saving R output within these script files (.r)
- Comment your code using #
 - Write a bullet point of what the line of code does for future reference

Find a way to structure and organize your scripts and data

Save often ... save again

Open your R window in a work directory away from backups and original files - you do not want to accidently overwrite them. R does not ask if "you are sure".

Script files your notebook of code

Here is an example of a script file viewed in Notepad++

Comments start with #

R recognizes # and will not try to execute these lines. Also notepad++ highlights comment lines in green

Different colors make the code visually easier to read

```
# Checking for zero data entries
     length (as.vector (which (as.vector (expM) == 0)))
     ### MM file created, checked if all measured were included
     ### split MM into 6 files for computational reasons
16
     library (petal)
     ### first sub-matrix done step-wise to double-check output
     m1 = readinTable("smallMM 1.txt")
     vec = as.numeric(m1[,3])
     indexRM = as.vector(which(abs(as.numeric(m1[,3]))<0.5))
     newCurrentMM = m1[-indexRM,]
     rm (indexRM, m1)
24
     ### integrating remaining 5 files automatically
    □for(i in 2:6) {
          currentMM = readinTable(paste("smallMM ",i,".txt",sep=""))
          tmpVec = as.numeric(currentMM[,3])
          vec = c(vec, tmpVec)
          indexRM = as.vector(which(abs(as.numeric(currentMM[,3]))<0.5))
          tmpnewCurrentMM = currentMM[-indexRM,]
          newCurrentMM = rbind(newCurrentMM, newCurrentMM)
          rm (currentMM, tmpVec, indexRM, tmpnewCurrentMM)
34
     ### checking distribution of association measure (Spearman)
     measures = vec
     min (measures, na.rm = TRUE) - 0.5, digits = 0)
     maxExp = round (max (measures, na.rm = TRUE) + 0.5, digits = 0)
     png (filename = "Histogram MM.png"), width = 7, height = 4.25, units = "in", pointsize = 12, bg = "white", res = 600)
     h = hist(measures, breaks = seg(-1, 1, by = 0.05))
     dev.off()
43
     min (measures)
     # [11 -0.872
     max (measures)
     # [1] 1
     dim (newCurrentMM)
     # [1] 33132192
     write.table(newCurrentMM, file = "MM reduced.txt", sep = "\t", quote=FALSE, row.names = FALSE, col.names = TRUE)
     save (newCurrentMM, file="newCurrentMM.RData")
```

Directory

Directory is just another word for folder

Make sure you know in which directory your R window is

Work in an directory away from backups and original data files

- > getwd()
- > setwd("C:/ <TAB>)
- > getwd()

You are learning a new language

Syntax

 \circ What is the difference between f(x) versus f[x]?

Vocabulary

- What functionality is part of R and what do I have to code myself?
- What are these functions called?

Thinking in that language rather than translating

• "Did you go to school?" Versus "Did you to school go?" (German)

Do not try to memorize everything - write scripts and know where to find them, plus google and other people's code are your best friends.

Variables

- Any letter, sequence of letters, or letters followed by a number can be used as variable names
 - X, xx, myData, myData1, myData2, my data
- A variable name cannot start with a number.
- Short variable names are great for typing, long variable names often make it easier to understand the code at a later time
- Do not overwrite R's predefined functions, in some cases an error message will appear
- <Tab> autocompletes variable names, function names, or filenames very handy
- R is case sensitive
- Avoid overriding variables

Data Handling

Indexing, Simple Manipulation, Uploading, Investigation, Exporting data

Vectors & Operations & Functions

A vector is an ordered sequence of numbers

$$> x = c(5, 64, 7, 3)$$

Multiple ways to create a vector

$$> y = c(34, 65, 0, 7)$$

$$> x = c(5, 64, 7, 3)$$

$$>$$
 new1 = cbind(x,y)

$$> y = 4:7$$

$$>$$
 new2 = rbind(x,y)

$$> z = seq(4,7)$$

$$> t = seq(1, 11, by=2)$$

$$> u = seq(1, 11, by=2)$$

$$> tmp = rep(10, 6)$$

$$> x[10] = 4$$

Indexing and Simple Manipulations

```
> women$height
R has some integrated dataset available
                                    > meanh = mean(women[,1])
> data()
> women
                                    > meanw = mean(women[,2])
> dim(women)
                                    > hmin = min(women[,1])
> rownames (women)
                                    > hmax = max(women[,1])
> colnames (women)
> women[1,1]
                                    > table(women)
> women[1,]
                                    > hist(women[,1])
> women[,2]
                                    > hist(women[,1], breaks = seq(57, 73, by=1))
> women[1:4,1:2]
```

Investigating Data & Correlation & Standard Stats

Refer to worksheet

LET'S TAKE A BREAK



Data Upload

Function		
read.csv()	Convenient function for .csv files Returns a list-object	read.csv("data.csv",
read.table()	Any delimited text files can be read-in, can be clunky at times, Returns a list-object	read.table("data.txt"
scan()	Reads file line by line Returns a vector-object	scan("data.txt", whars scan("data.txt", wha

, header = TRUE) t", header=TRUE, sep="\t") ", header=FALSE, sep="\t") at="", sep="\t", nline=1, skip=1) at="", sep="\t", n=15, skip=1)

Data Upload Script

```
fileName = ""
# make sure the file is in table format
# line 1 should have the same length as line 2
11 = scan(fileName, sep = "\t", what = "", nlines = 1)
tmpM= matrix(scan(fileName, what = "", sep = "\t", skip = 1), ncol = length(l1), byrow = TRUE)
myData = matrix(as.numeric(tmpM[, -1]), ncol = length(l1) -1, byrow = FALSE)
colnames(myData) = 11[-1]
rownames (myData) = tmpM[, 1]
# remove variables (objects) you no longer need
rm(fileName, tmpM, 11)
# check if the file was read in correctly
dim(myData)
head (myData)
# myData is a numeric matrix object
```

Exporting Data

Objects can also be saved as .RData files

```
> save(new_trees, file="trees.RData")
> q()
> load("trees.RData")
> objects()
```

Can save more than one object Good coding tip: give the objects meaningful names, not x and y - two days from now you won't know what x and y are :)

```
> save(x, y, file="dataObjects.RData")
```

Data Visualization

Histograms and Scatter Plots

Histogram - hist()
&

Scatterplot - plot()
&

Graphical Parameters

Refer to worksheet

Plotting two graphics intro one window

```
vec = as.vector(myData)
MIN = round(min(vec) - 0.5, digits = 0)
MAX = round(max(vec) + 0.5, digits = 0)
par(mfrow = c(2, 1))
hist(myData[,1], breaks=seq(MIN, MAX, by = (MAX-MIN)/numBins), col="green",
xlab="Control", main="Values for Control Group")
hist(myData[,2], breaks=seq(MIN, MAX, by = (MAX-MIN)/numBins), col="red",
xlab="Treatment", main="Values for Treatment Group")
```

Histogram - hist()

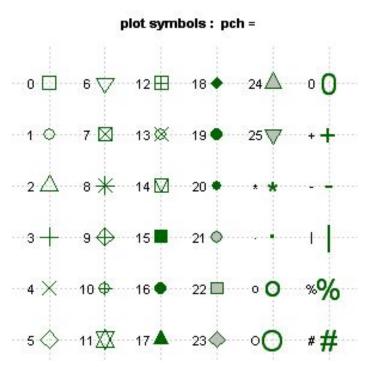
```
# number of histogram's subintervals
numBins = 100
# MIN and MAX will always be integers
MIN = round(min(myData[,1])-0.5, digits=0)
MAX = round(max(myData[,1])+0.5, digits=0)
h = hist(myData[,1], breaks=seq(MIN, MAX, by = (MAX-MIN)/numBins),
col="green", xlab="Control", main="Values for Control Group")
# see statistics of h
# barplot() versus hist()
barplot(myData[,1])
```

Histogram - hist()script

```
TTTTF =
xAXTS =
COLOR =
vec =
numBins =
                               # number of histogram's subintervals
fileName =
# MIN and MAX will always be integers
MIN = round(min(vec) - 0.5, digits = 0)
MAX = round(max(vec) + 0.5, digits = 0)
png(filename = paste(fileName, ".png", sep=""), width=7, height=8.5,
units="in", pointsize=12, bg="white", res=600)
h = hist(vec, breaks=seq(MIN, MAX, by=(MAX-MIN)/numBins), col=COLOR,
xlab=xAXIS, main=TITLE)
dev.off()
```

Plotting Symbols

Line Types



lty line type
lwd line thickness, default =1

Line Types: Ity=

