Introduction to R for Psychometrics

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Introduction to R

The R Project for Statistical Computing

What is R? Who is R? How can R help me?

Organization

This presentation will introduce the R computing environment. We will briefly address some history and fundamentals, but largely focus on applications: how R can be utilized for the tasks that we commonly deal with at MHS. This will include: data manipulation tasks, running basic and advanced statistical analyses, and key techniques for programming user defined functions.

What is R?

- R is an open source language and environment for statistical computing and graphics.
- Based upon S, which was developed by John Chambers and colleagues at Bell Laboratories in the late 1970s.
- R is a re-implementation of S that began development circa 1993 by Ross Ihaka and Robert Gentleman at the University of Auckland, New Zealand.



The current version of R is 3.1.3, code-named "Smooth Sidewalk".

Who is R?

- Since mid-1997, the R Development Core Team has been maintained with write access to the R source code
- The R Foundation, a not for profit organization with a board of 5 board members and 2 auditors, controls the design and evolution of base R.

However, the ${\tt R}$ Project is actually a large scale effort and features the work thousands of contributors world-wide.

- UseRs are able to develop their own collections of functions which can be "packaged" and hosted for public distribution on the Comprehensive R Archive Network (http://cran.r-project.org/)
- As of March 2015, the CRAN package repository hosts 6,417 available packages and counting!

What can R do for me?

- Interactivity and expressiveness.
- No black boxes! All code is available and open.
- Primarily designed as a programming language for statistical operations, and almost every statistical method, especially those on the cutting edge, have R implementations.
- One of R's strengths is the ease with which well-designed publication-quality plots can be produced.
- Additional packages provide a framework for everything from running spatial analyses, to text mining, to the construction of interactive visualizations, and even to the creation of...

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- this presentation!

Free?

Yes, free! R is an official part of the Free Software Foundation's GNU project with the standard General Public License. This means:

- you can obtain, copy, and re-use R without paying anything (freeware);
- you can access the source code and modify it (open source);
- you can redistribute your modifications back to the public for the improvement of the community at large (awesome).

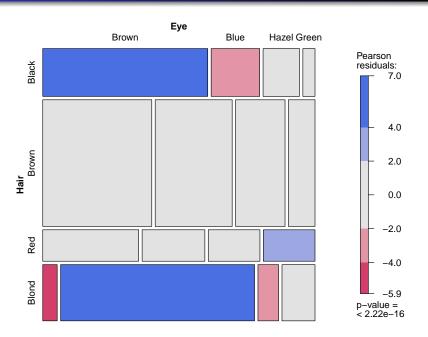
R's Future

As part of its mandate, R will always be free. However, a company called **Revolution Analytics** has proprietary rights for Revolution R.

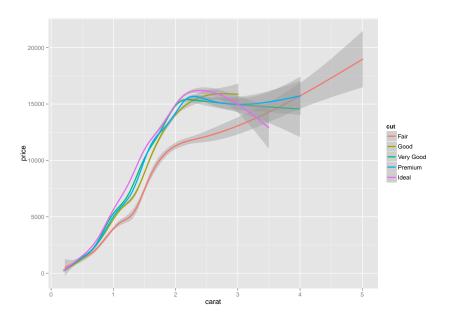
Revolution R is a parallel version that has been optimized for big data analytics. As testament to R's influence, Microsoft has recently purchased Revolution Analytics, and are rumoured to incorporate features that communicate with R into some of their applications.

...But why all of this interest?

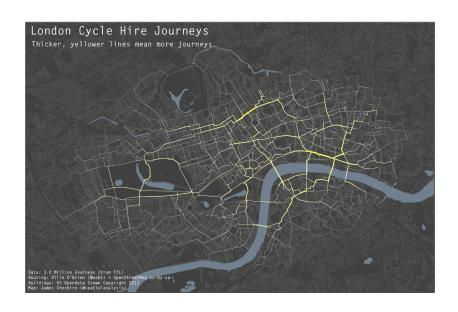
Categorical Data Analysis



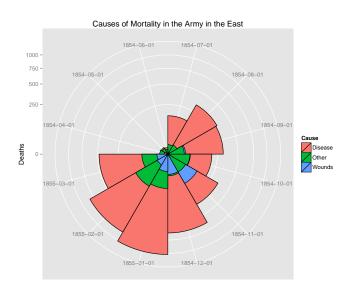
Non-linear Regression



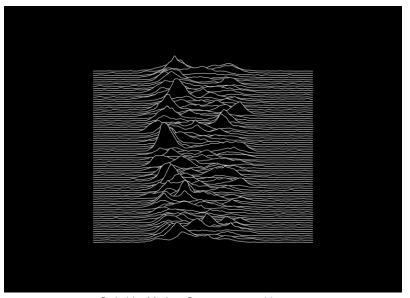
Spatial Mapping



Polar Coordinates



Unknown Pleasures



Coded by Markus Gesmann, magesblog.com

Oh, R...

- A challenge and strength of R is that it is entirely syntax based.
- Even those not interested in programming new functions must become familiar with handling objects and calling functions by writing scripts.
- While R operates on spreadsheets, it is not "spreadsheet software".
- Learning R will cause some frustration and require persistence, however it is well worth the effort!

"R is basically a big global family of fluffy altruistic people contributing to the goal of producing a versatile data analysis tool that is free for everyone to use."

— Andy Field, Discovering Statistics Using R, p. 63

"R is basically a big global family of fluffy curmudgeonly altruistic people contributing to the goal of producing a versatile data analysis tool that is free for everyone to use."

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- Large and active community of users who post tutorials online and answer questions on message boards.
- It rarely takes more than a quick Google search to find the solution to most R problems.
- Most packages are well-documented and package authors are generally accessible via online forum or e-mail.

Useful Resources

- Stack Exchange (http://www.stackexchange.com)
 - Stack Overflow (questions tagged [R])
 - Cross Validated (general statistics)
- R-Bloggers (www.r-bloggers.com)
- DataCamp (www.datacamp.com)
- Quick-R (www.statmethods.net) examples and tips
- R-seek (rseek.org) restricted search of R specific sites
- One Page R (onepager.togaware.com) cheat sheets

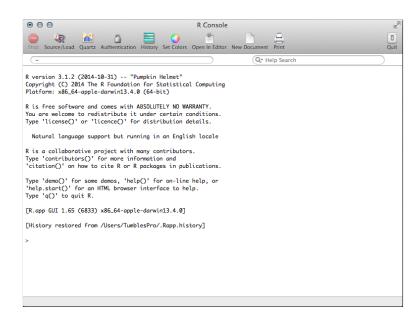
Venting

- Why 'R' is hard to learn
- The R Inferno

Get R

- R is available for Windows, Mac OS X, and Linux.
- R is downloaded from CRAN (cran.r-project.org/), and installs:
 - R.exe: a text-based command-line version, run via command prompt
 - RScript.exe: an interface designed to be used in scripts and CMD batch files
 - RGui.exe: both a command-line interface and a barebones GUI.
- We highly recommend (and assume) you will install the RStudio IDE promptly afterwards, which is also freely available for Windows, Mac OS X, and Linux (from rstudio.com).

The R Environment: RGui.EXE



The RStudio Environment

Unfortunately the user environment that comes with R leaves a lot to be desired. Fortunately, some other fluffy altruistic people developed something better: RStudio.

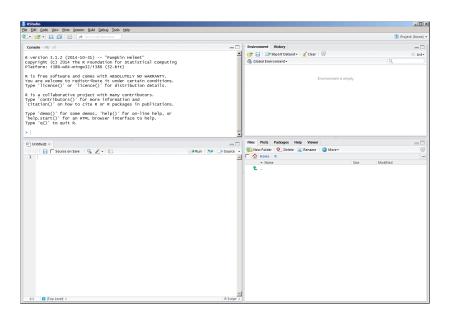
RStudio IDE is a powerful and productive user interface for R, currently on version 0.98.1103.

Tips for Installing R and RStudio

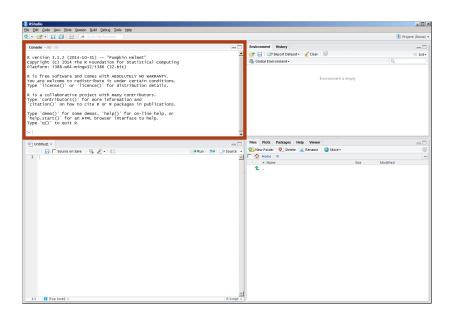
- When installing, the default options are acceptable.
- Upon launching RStudio, it can be useful to change some options (Tools -> Global Options)
- For example, set "Save Workspace to .RData on Exit" to Never
- Set Console to upper left corner, Source to lower left.

BONUS: RStudio keyboard shortcuts cheatsheet for even faster coding.

The RStudio Environment



The Console



The Console

The main window in R is called the Console. This is where R syntax is executed and you will see the results of the submitted code. We can interact with R directly through the Console window by typing commands at the > prompt and pressing <return>.

```
5 + 16
## [1] 21
```

Using the Console, we can also create variables. Variables will be discussed more extensively later, but for now we can type the following lines into the console and press <return>.

```
x <- c(3, 5, 5, 7, 9)
x
## [1] 3 5 5 7 9
```

The Console

The variable x is saved in the R Workspace, and it will still exist until it is deleted. You can check the contents of the Workspace by using the command 1s().

```
ls()
## [1] "x"
```

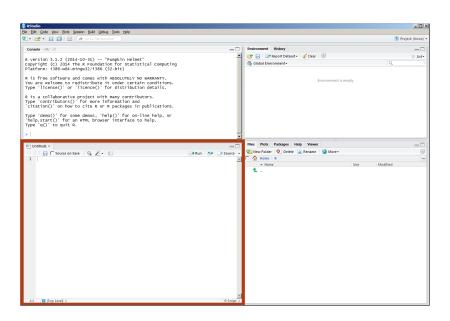
Assigning Values in R

Note: We typically use <- to assign variables rather than =. In most cases you can use =, but you will sometimes get some strange behaviour. As a rule it is safer to use <- when assigning values to variables.

R does not produce any output when you submit assignments. If you want to print the results you can add parentheses:

```
(x <- c(3, 5, 5, 7, 9))
## [1] 3 5 5 7 9
```

The Source Window (or Editor)



The Source Window (or Editor)

Upon launching RStudio, you may have to create a new script via File -> New File -> R Script or open a pre-existing script via File -> Open File.

.R files are plain text: can be modified with any text editor, cross-platform.

In the Editor, we can write a series of R commands that we would like to run. Once in the editor window, code can be run line by line by pressing <ctrl> + R or <ctrl> + Enter. Multiple lines can be run simultaneously by highlighting them and using the same shortcut.

We can also highlight the desired code and select 'Run' from the top right of the Editor.

The Source Window (or Editor)

```
FirstScript.R ×

☐ Source on Save 
☐ ✓ ✓ □
                                                                               Run 🕪 🕒 Source
  1 x < -c(3,5,5,7,9)
     y \leftarrow c(2,4,8,5,14)
    x2 <- c(9,10,11,12,15,12)
y2 <- c(1,2,3,4,5,6)
7 ls()
8
9 plot
     plot(x,y)
 10:1
       (Top Level) $
                                                                                                 R Script ¢
```

Coding Conventions

Basic structure of R syntax:

```
out <- t.test(y ~ x, data = dat, alternative = "less")</pre>
```

- the function we are calling is t.test()
- y ~ x is an example of R's formula interface, in which only variable names are provided
- data and alternative are arguments for t.test()
- dat and "less" are objects that are matched to their arguments
- output from t.test() is being returned and stored in another object, out, via <- (the assignment operator).

NOTES:

- No use of "." to designate end of line (can use ;, but is often unnecessary).
- No inclusion of anything like EXECUTE.

Coding Conventions

Comments in R

```
# The # symbol is used to demark comments in R
x <- c(3, 5, 5, 7, 9)
# y <- c(1, 2, 3, 4, 5)
x # print X
## [1] 3 5 5 7 9
y # print Y
## Error in eval(expr, envir, enclos): object 'y' not found</pre>
```

Commenting code inline is **highly encouraged**, and doubly so for collaborative projects.

Note: We will soon introduce a better way of managing your comments and code through the use of **Markdown**.

Coding Conventions

Spacing in R Syntax

R is almost always oblivious to spacing. There are a few notable exceptions, but they tend to come out in more advanced features (e.g., using if statements and embedded functions). The following will all produce the same results, so feel free to use whatever spacing structure you find most aesthetically appealing.

```
x <- c(3, 5, 5, 7, 9)

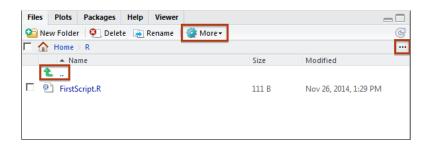
x <- c(3,5,5,7,9)

x <- c(3,5,5,7,9)

x <- c(3, 5, 5, 7, 9)
```

If you want to follow some leading sources on coding style, Google and Hadley Wickham (a leading R programmer) both have R-specific style guides you can follow.

File Manager



- Navigate by clicking on folders or the double dots/green arrow to go up a level.
- Use the ellipsis to open a file browser.
- Under More, lives the very useful Set As Working Directory command.

File Manager

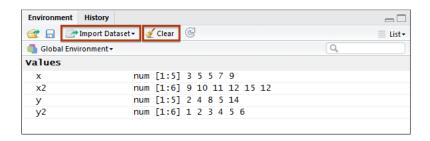
- What you see in the file pane shows R's working directory.
- This is very important to keep track of, as this is the directory R will look in when you specify a filename!

Tips and Tricks:

- If you open RStudio using the default shortcut, the working directory is set under Tools -> Global Options
- This usually is something like:C:/Users/your.name/Documents/R/
- However, if you open RStudio by double clicking an .R or .Rmd file, the working directory will automatically be set to the same directory the file is in.
- The working directory can also be manually modified via setwd().

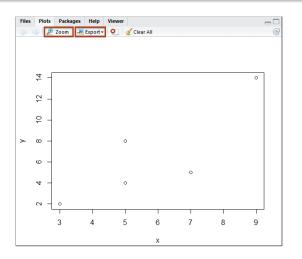
It is highly recommended to keep a project's datafiles and R scripts in the same directory, especially if you plan to knit a report.

Environment



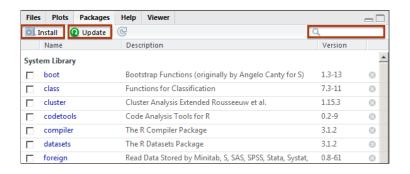
- Live updating, shows everything currently in the workspace.
- Headings for Data, Values, and Functions
- Same basic information as using 1s(), but also denotes variable type, length, and the first 10 elements.
- Can click on an object to see what it contains.
- Two very handy commands: Import Dataset and Clear to clean the workspace.

Plot Planel



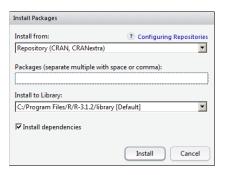
- Unlike base R, we can navigate through many plots via arrows.
- Use Zoom command to see plot in higher resolution.
- Use Export to save current plot in a variety of formats.

Packages



- Base functions and packages are shipped with R.
- Many packages exist and can be installed from:
 - CRAN or locally via Install
 - GitHub (e.g., Trending R Repositories)
 - Other Repositories (e.g., Bioconductor.org, rOpenSci.org)
- Use Update periodically to keep packages current.

Installing Packages



- Text input live updates with available CRAN packages
- Always leave Install dependencies checked
- Can also use: install.packages("NAME", dep = TRUE) in the console (my preferred method).

Installing Packages

install.packages("polycor", dep = T) installs: polycor,
mvtnorm, and sfsmisc.

Note: Just because a package has been installed, doesn't mean that it is available for use! We need to load packages that we want to use.



The package appears in our list, but is unchecked.

library(polycor) will make polycor's functions available for use.



Getting Help

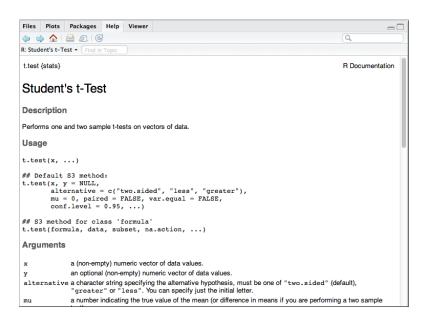
In the console, use ? and ?? to search for a particular function name, and for pattern matching, respectively. For example:

?t.test
??noise

Notes:

- The ? approach only searches loaded packages, and only for an exact match (?ttest gives an error)
- The ?? approach searches packages installed locally for your keyword, and they do not need to be loaded.

Getting Help



Getting Help

R help files generally follow a template:

- Function Name {Package}: e.g. t.test {stats}
- **Description**: Brief idea of what the function does.
- Usage: Function call, with defaults indicated
 - t.test(x, y = NULL, alternative = c("two.sided", "less",
 "greater"), mu = 0, paired = FALSE, var.equal = FALSE,
 conf.level = 0.95, ...)
- **Arguments**: Detailed list of function's parameters.
- Details: Generally technical details about function's behaviour.
- Value: Description of the output generated from the function.
- See Also: A list of other functions that you may find useful.
- Examples: The best part of a help file!

Further Documentation

- demo(): brings up a list of demonstrations for a set of tasks.
 - Base R only has 16, but many packages have additional demos.
 - All available: demo(package=.packages(all.available=TRUE))
 - Example: demo(lm.glm, package="stats")
- browseVignettes() brings up a list of vignettes.
 - These are short documents on a particular task, usually feature R syntax, output, and PDFs that include all output.
 - Browse from available demos from a particular package: browseVignettes(package="grid")
 - Example: vignette("rotated", package="grid")

R + RStudio + Markdown = Dynamic Documents

.R files (like SPSS syntax files) only ever have the commands and comments that you decide to include in them.

- No output!
- No plots!
- No narrative!

A substantially more powerful method of working is to work with **R Markdown (.Rmd) files** instead of R scripts (.R). The benefits of working with Markdown are:

- Flexibility in text formatting (titles, subtitles, quotes)
- The ability to include lists, links, and images
- The ability to output the document to HTML, PDF, even DOCX!
- But, most of all: the ability to include raw R code ("R chunks") and R output in the same document!

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Reproducable research is the best research!

Markdown Example: .Rmd Code

Create a new Markdown file via File -> New File -> R Markdown.

```
ABC Q ? + AF Knit HTML + @
                                                                                                 Run 🖦 🖸 Chunks 🕶
1 * # Project Markdown: Main Titles via Single Pound
   Name: Your Name
    Date: The Current Date
5 To force a line break, end the line with two spaces. Some simple formatting options are: "italic" and ""bold".
7 - ## Sub-Headings via Double Pound
9 Within a section of the document, we can use plain text to discuss our goals or plans for the analysis.
11 > To visually pop something, we can use a quotation environment via ">".
13 To insert R code, use three backticks followed by {r}. Anything that would work in an R script will work here!
14
15 - ```{r}
16 # Insert your R code inside a named chunk, like this:
17 summary(cars)
18 x \leftarrow c(1,2,3)
19 x
20 - ***
```

Markdown Example: Knit .html File

Project Markdown: Main Titles via Single Pound

Name: Your Name
Date: The Current Date

To force a line break, end the line with two spaces. Some simple formatting options are: italic and bold.

Sub-Headings via Double Pound

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To visually pop something, we can use a quotation environment via ">".

To insert R code, use three backticks followed by {r}. Anything that would work in an R script will work here!

```
# Insert your R code inside a named chunk, like this:
aummary(care)

## speed dist
## Min. : 4.0 Min. : 2.00
## lst Qu.:12.0 lst Qu.: 26.00
## Median :15.0 Median : 36.00
## Medn :15.4 Mean : 42.98
## 3rd Qu.:19.0 3rd Qu.: 56.00
## Max. :25.0 Max. :120.00

x <- c(1,2,3)
x

## [1] 1 2 3
```

Markdown Tips and Tricks



- May need to run: install.packages(knitr) before you are able to knit successfully.
- Knit to PDF requires a TeX distribution installed.
- Look at the Markdown Quick Reference page to see more functionality options, such as:
 - Lists (unordered via asterisks, ordered via numbers)
 - Incorporate Links, Images, LaTeX Equations, Page Breaks, Tables.
- Give chunks descriptive names for easier navigation, e.g.:



- Use the Chunks menu to run Previous/Current/Next/All Chunks.
- Incorporate Chunk options like: eval, echo, cache.

R Programming

Basic R

Working with the Fundamental Object Types

Data Types

When we created x earlier, we saw that it was "numeric". Objects in R have different types. It might seem cruel, but if you learn how to handle these types of objects you will be in a better position to correct problems with your code and manage data sets.

The most useful variable types to know about are:

- Numeric scalars/vectors
- Logical vectors
- Character vectors
- Matrices
- Lists
- Data frames

Numeric Scalers/Vectors

Basic unit of analysis. These objects are either a single element (a 'scalar'; e.g., $x \leftarrow 5$ or $x \leftarrow "Rrr"$) or a series of elements collected together inside a single object (a vector; e.g., x < c(5, 10, 15)).

We use the function c() which stands for 'combine' or 'concatenate' to create a vector.

```
x \leftarrow 5 # a scalar.

x \leftarrow c(3, 5, 5, 7, 9) # combine numbers into a vector

# via the function c().
```

We can access an entry in a vector using square brackets:

```
x[4] # Returns the 4th value of the vector x ## [1] 7 x[-4] # Returns all values of x but the 4th ## [1] 3 5 5 9
```

Numeric Scalers/Vectors

Shortcuts: Using :, seq(), and rep.

```
(one2ten <- c(1:10))
## [1] 1 2 3 4 5 6 7 8 9 10
```

```
(bytwos <- seq(from = 1, to = 10, by = 2))
## [1] 1 3 5 7 9
```

```
(repeats <- rep(1:4, times = 2))
## [1] 1 2 3 4 1 2 3 4
(repeats <- rep(1:4, each = 2))
## [1] 1 1 2 2 3 3 4 4</pre>
```

NOTE: Typically assignment is done *invisibly*. Surrounding an assignment with (...) is an R shorthand for print(...), which evaluates the line and also prints whatever is returned.

Only contain **TRUE** and **FALSE** values.

Logical Vector Operations

- == exactly equals (NOTE: this is not an assignment!)
- < less than</p>
- > greater than
- < = less than or equal to</pre>
- >= greater than or equal to
- ! = not equal to

For example:

```
x < 7 # Identifies positions in x less than 7
## [1] TRUE TRUE TRUE FALSE FALSE
x != 5 # Identifies positions in x not equal to 5
## [1] TRUE FALSE FALSE TRUE TRUE</pre>
```

Filtering: A logical vector can used to 'filter' or 'subset' a numeric vector based on some criteria. For example, if we wanted to select or filter all the cases from the vector x that are less than 7:

```
x # Print x ***
## [1] 3 5 5 7 9
x < 7 # Print positions less than 7 ***
## [1] TRUE TRUE TRUE FALSE FALSE
filter <- x < 7 # Save logical vector
x[filter] # Print values of x less than 7 ***
## [1] 3 5 5
x_less_7 <- x[filter]
x_less_7 # Print x_less_7 ***
## [1] 3 5 5
# Lines marked with `***` only print output to the console!</pre>
```

Filtering: A logical vector can used to 'filter' or 'subset' a numeric vector based on some criteria. For example, if we wanted to select or filter all the cases from the vector x that are less than 7:

```
x # Print x ***
## [1] 3 5 5 7 9
x < 7 # Print positions less than 7 ***
## [1] TRUE TRUE TRUE FALSE FALSE
filter <- x < 7 # Save logical vector
x[filter] # Print values of x less than 7 ***
## [1] 3 5 5
x_less_7 <- x[filter]
x_less_7 # Print x_less_7 ***
## [1] 3 5 5
# Lines marked with `***` only print output to the console!</pre>
```

One-liner:

```
(x_less_7 <- x[x < 7])
## [1] 3 5 5
```

We can also chain together such statements using:

- a vertical bar or pipe: | (not a capital i; found on backslash key) to signify OR
- an ampersand: & to signify AND
- an exclamation point: ! to signify NOT

This code will print ("return") the values of the vector 'x' that are not exactly equal to 5 AND are greater than 3.

```
x[x != 5 & x > 3]
## [1] 7 9
```

If we want the location of those values, we can use the function which():

```
which(x != 5 & x > 3)
## [1] 4 5
```

Character Vectors

Like numeric scalars/vectors but for non-numeric data (strings).

```
y <- "abc"
y
## [1] "abc"
```

```
class(y)
## [1] "character"
y[1] # the first element of y
## [1] "abc"
y[2] # the second element of y
## [1] NA
```

Character Vectors

Like numeric vectors, y can be longer than a single element:

```
y <- c("abc", "de", "123", "")
y
## [1] "abc" "de" "123" ""
y[1]
## [1] "abc"
y[2]
## [1] "de"
y[3]
## [1] "123"
class(y[3]) # even though this only has numbers it is not numeric!
## [1] "character"</pre>
```

Character Vectors

There are many ways we can manipulate character vectors in R. For example:

```
substring(y, first = 1, last = 2)
## [1] "ab" "de" "12" ""
# Returns the first two characters in each entry of 'y'

# Combine strings:
a <- 9
paste0("This is a number I like: ", a,". It's a good one, huh?")
## [1] "This is a number I like: 9. It's a good one, huh?"
paste0(y, " is the ", 1:4, " entry in character vector y.")
## [1] "abc is the 1 entry in character vector y."
## [2] "de is the 2 entry in character vector y."
## [3] "123 is the 3 entry in character vector y."
## [4] " is the 4 entry in character vector y."
## this even works on vectors!</pre>
```

Advanced: Character vectors can be manipulated with regular expressions using base functions grep() and sub(), or the functions in the stringr package.

Exercises, Pt. 1

Basic R Exercises, Pt. 1

We are now ready to attempt the Exercise questions pertaining to "Numeric and Logical Vectors".

Factors are a special type of character vector that are extremely useful in psychological research, and are used to code categorical data.

We could use a numeric variable for this purpose:

```
group <- c(1,1,1,2,2,2,3,3,3, sample(c(1:3), 11, replace=TRUE))
class(group)
## [1] "numeric"
str(group)
## num [1:20] 1 1 1 2 2 2 3 3 3 3 ...</pre>
```

... But it can lead to mistakes:

```
(group+2)
## [1] 3 3 3 4 4 4 5 5 5 5 5 4 5 5 4 5 5 4 3 5 3
```

Also, the labels are not useful (1, 2, and 3, rather than "Group1", "Group2", and "Control").

```
xtabs(~group)
## group
## 1 2 3
## 5 6 9
```

We can convert a numeric vector into a factor using the factor() command.

Note the levels argument for telling R which levels you want to keep (in most cases, you will keep all of them and this can be omitted), and the labels argument for specifying their names.

```
group1 <- factor(group)
class(group1)
## [1] "factor"
str(group1)
## Factor w/ 3 levels "1","2","3": 1 1 1 2 2 2 3 3 3 3 ...
xtabs(~group1)
## group1
## 1 2 3
## 5 6 9</pre>
```

When using levels and labels, order matters!

But besides making output easier to read and interpret, converting our variables to factors also protects us from silly mistakes, like:

If we already have a factor and want to rename its levels, it is pretty easy:

```
levels(group1)
## [1] "1" "2" "3"
levels(group1) <- c("Group1", "Group2", "Control")
group1
## [1] Group1 Group1 Group1 Group2 Group2 Control Control
## [9] Control Control Group2 Control Group2 Control Control
## [17] Group2 Group1 Control Group1
## Levels: Group1 Group2 Control</pre>
```

```
xtabs(-group1)
## group1
## Group1 Group2 Control
## 5 6 9
```

Factors typically are sorted alphabetically by functions (with the first group often used as the reference group) since factor levels are considered nominal.

One way of circumventing that behaviour is to make the factored "ordered". This can be done either by setting ordered = TRUE when calling factor, or calling ordered() on the factor variable directly.

Two-dimensional vectors: all elements must of the same type.

Matrices can be constructed directly via matrix(), which has the following defaults: matrix(data, nrow, ncol, byrow=FALSE) - data is filled in column-wise!

```
mat <- matrix(data = c(1,2,3,4,5,6,7,8,9,10,11,12), nrow = 3)
mat

## [,1] [,2] [,3] [,4]

## [1,] 1 4 7 10

## [2,] 2 5 8 11

## [3,] 3 6 9 12
```

We can also stack vectors via rbind() (which binds vectors together by row) or cbind() (bind by column).

Use double subscripts to navigate matrices.

```
mat1[1, 3] # first row, third column
## [1] 3
mat1[2, ] # second row, all columns
## [1] 3 4 5 6
mat1[ , 3] # all rows, third column
## [1] 3 5 7
```

Use double subscripts to navigate matrices.

```
mat1[1, 3] # first row, third column
## [1] 3
mat1[2, ] # second row, all columns
## [1] 3 4 5 6
mat1[, 3] # all rows, third column
## [1] 3 5 7
```

This can also be used to change values:

```
mat1[2, ] <- mat1[2, ] + 4
```

```
mat1
## [,1] [,2] [,3] [,4]
## [1,] 1 2 3 4
## [2,] 7 8 9 10
## [3,] 5 6 7 8
```

Labeling a Matrix: For aesthetic purposes, we can label our rows and columns via rownames() and colnames(). This brings a matrix closer to a data frame where each row might pertain to a particular person, and each column a variable we have measured.

```
rownames(mat1) <- c("Row1", "Row2", "Row3")
colnames(mat1) <- c("Col1", "Col2", "Col3", "Col4")
mat1

## Col1 Col2 Col3 Col4

## Row1 1 2 3 4

## Row2 7 8 9 10

## Row3 5 6 7 8
```

Exercises, Pt. 2

Basic R Exercises, Pt. 2

We are now ready to attempt the Exercise questions pertaining to "Matrices".

Lists are like a vector but each of its contents can be items of different data types. Many R functions return lists, so familiarity with them is important!

```
a <- 5
y <- c("abc", "de", "123", "")
iamalist <- list(a = a, yvar = y, mat = matrix(1:6, 2, 3))</pre>
iamalist.
## $a
## [1] 5
##
## $vvar
## [1] "abc" "de" "123" ""
##
## $mat
##
       [,1] [,2] [,3]
## [1,] 1 3
                    5
## [2,] 2 4 6
```

We can look at their structure to get a sense of what they contain:

```
str(iamalist)
## List of 3
## $ a : num 5
## $ yvar: chr [1:4] "abc" "de" "123" ""
## $ mat : int [1:2, 1:3] 1 2 3 4 5 6
```

Navigate lists either with square brackets and an index location, or using a dollar sign and the vector's name:

```
iamalist[[1]]
## [1] 5

iamalist$yvar
## [1] "abc" "de" "123" ""

iamalist[[3]][1, 2]
## [1] 3
```

NOTE: When we navigate lists with square brackets, we use the double square bracket notation: [[...]]. This is because using [...] returns a list object, while [[...]] returns the vector in its original type.

List output from t.test():

```
t.test(iamalist$mat[1, ], iamalist$mat[2, ])
##
## Welch Two Sample t-test
##
## data: iamalist$mat[1, ] and iamalist$mat[2, ]
## t = -0.6124, df = 4, p-value = 0.5734
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -5.533916 3.533916
## sample estimates:
## mean of x mean of y
## 3 4
```

Saved as an object:

```
out <- t.test(iamalist$mat[1, ], iamalist$mat[2, ])
str(out)
## List of 9
## $ statistic : Named num -0.612
## ..- attr(*, "names")= chr "t"
## $ parameter : Named num 4
## ..- attr(*, "names")= chr "df"
## $ p.value : num 0.573
## $ conf.int : atomic [1:2] -5.53 3.53
## ..- attr(*, "conf.level")= num 0.95
##
   $ estimate : Named num [1:2] 3 4
## ..- attr(*, "names")= chr [1:2] "mean of x" "mean of y"
##
   $ null.value : Named num O
##
   ..- attr(*, "names")= chr "difference in means"
##
   $ alternative: chr "two.sided"
   $ method : chr "Welch Two Sample t-test"
##
##
   $ data.name : chr "iamalist$mat[1, ] and iamalist$mat[2, ]"
   - attr(*, "class")= chr "htest"
##
```

If our output is saved as a list object, we can access elements of it just like we did before:

Data Frames are the most common object type for data analysis in R. They are similar to matrices in many ways except for one large difference: each column can hold a different object type.

For example, our first column may pertain to an identification code (character), followed by age (numeric), followed by gender (character, factor), and so on.

To combine vectors of the **same length** into a dataframe use data.frame(), e.g.: data.frame(var1, var2, var3)

More typically, we obtain data directly by it in from an external file. Data can be read in various ways, but the most reliable is to convert to CSV and then use read.csv(). For example:

```
dat <- read.csv("exerciseDat.csv")
head(dat)

## id gender mathPre readPre group score1 score2 pass

## 1 1 0 55.15 56.31 mnemonic 48.33 68.74 1

## 2 2 1 53.30 30.45 mnemonic 45.91 43.89 0

## 3 3 1 56.96 69.53 cram 55.98 84.43 1

## 4 4 1 55.64 38.25 mnemonic 52.81 38.24 0

## 5 5 1 48.14 60.25 mnemonic 53.70 54.46 1

## 6 6 0 49.07 61.91 cram 46.30 61.31 1
```

SPSS, SAS, and STATA datafiles are best loaded via the haven library:

```
#install.packages("haven")
library("haven")
dat1 <- read_dta("filename.dta")
dat2 <- read_sas("filename.sas7bdat")
dat3 <- read_spss("filename.sav")</pre>
```

Excel files can be read via read.xlsx() from the xlsx library:

```
dat4 <- read.xlsx("myfile.xlsx", sheetName = "Sheet1")</pre>
```

R objects can be saved as .RData files using save(), and are brought back using load(). These do not require assignment!

```
load("myfile.RData")
```

However, base ${\tt R}$ and many packages also come with datasets built-in to use as examples.

```
data() # lists all datasets from base R
data(package=.packages(all.available = TRUE)
data(package="ggplot2") # only datasets from ggplot2
```

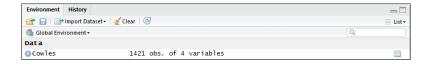
The following example uses data from Cowles and Davis' survey on Volunteering for Psychological Research, published in the *British Journal of Social Psychology* in 1987, and included in John Fox's car package.

NOTE: If the car package is not installed, you must install it!

```
#install.packages("car", dep=T)
library(car)
```

```
data(Cowles) # loads Cowles
names(Cowles)
## [1] "neuroticism" "extraversion" "sex" "volunteer"
```

- Neuroticism: scale from Eysenck personality inventory
- Extroversion: scale from Eysenck personality inventory
- Sex: factor with 2 levels (female, male)
- Volunteer: factor with 2 levels (yes, no)

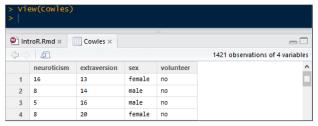


```
Data

Ocoles

1421 obs. of 4 variables
neuroticism : int 16 8 5 8 9 6 8 12 15 18 ...
extraversion: int 13 14 16 20 19 15 10 11 16 7 ...
sex : Factor w/ 2 levels "female", "male": 1 2 2 1 2 2 2 2 2 ...
volunteer : Factor w/ 2 levels "no", "yes": 1 1 1 1 1 1 1 1 1 1 ...
```

Variable list in RStudio's Environment Window



Using View(Cowles) opens a new tab to see data

After importing a dataset, it is good to make sure it was imported correctly:

- Use head(), tail(), and str() to understand dataset's structure
- Use summary() to get an overview of the descriptive statistics across all variables
- The describe() function from the psych library provides a more detailed summary
- Use complete.cases() if you want to filter out cases with missingness

```
head(Cowles)
    neuroticism extraversion sex volunteer
## 1
              16
                          13 female
                                            nο
## 2
               8
                          14 male
                                            no
## 3
               5
                           16 male
                                            no
## 4
                          20 female
                                           no
## 5
                          19 male
                                            no
## 6
                           15 male
                                           no
tail(Cowles, n = 8)
##
       neuroticism extraversion sex volunteer
## 1414
                              20 male
                                             yes
## 1415
                                  male
                                              ves
## 1416
                              10 female
                                              ves
## 1417
                              10
                                  male
                                              yes
## 1418
                 8
                              4 female
                                              ves
## 1419
                              8 male
                                             yes
## 1420
                 19
                              20 female
                                              yes
## 1421
                 15
                              20
                                  male
                                              ves
```

```
str(Cowles)
## 'data.frame': 1421 obs. of 4 variables:
## $ neuroticism : int 16 8 5 8 9 6 8 12 15 18 ...
## $ extraversion: int 13 14 16 20 19 15 10 11 16 7 ...
## $ sex
          : Factor w/ 2 levels "female", "male": 1 2 2 1 2 2 1 2 2 2 ...
## $ volunteer : Factor w/ 2 levels "no". "ves": 1 1 1 1 1 1 1 1 1 1 1 ...
mean (CowlesSextraversion)
## [1] 12.37298
var(Cowles$extraversion)
## [1] 15.16079
median(CowlesSextraversion)
## [1] 13
summary(Cowles)
## neuroticism extraversion
                                    sex volunteer
## Min. : 0.00 Min. : 2.00 female:780 no:824
## 1st Qu.: 8.00 1st Qu.:10.00 male :641 yes:597
## Median:11.00 Median:13.00
## Mean :11.47 Mean :12.37
## 3rd Qu.:15.00 3rd Qu.:15.00
## Max. :24.00
                  Max. :23.00
```

```
table(Cowles$sex)
##
## female male
## 780 641
table(Cowles$sex, Cowles$volunteer)
##
## no yes
## female 431 349
## male 393 248
```

How does R think about data?

```
dim(Cowles)
## [1] 1421     4
Cowles[40, 1] # 40th Case from "neuroticism"
## [1] 5
Cowles$neuroticism[40]
## [1] 5
Cowles$[1:8, 2] # First 8 cases from "extraversion"
## [1] 13 14 16 20 19 15 10 11
Cowles$extraversion[1:8]
## [1] 13 14 16 20 19 15 10 11
```

Manipulating Variables: Data frames are basically lists with an additional restriction (each variable has the same length), and so we can access elements of the dataframe using \$, as we did with lists.

To overwrite a variable, simply use its name on the lefthand side of an assignment operator:

```
Cowles$extraversion <- NULL
# this deletes extraversion from our dataset!
```

More often, we want to add a variable to our data frame. So long as we use a name not in use, it will be appended appropriately:

```
Cowles$sqrt.neuro <- sqrt(Cowles$neuroticism)
names(Cowles)
## [1] "neuroticism" "extraversion" "sex" "volunteer"
## [5] "sqrt.neuro"</pre>
```

Subsetting: Can use conditional expressions, like we did with vectors.

```
the.females <- Cowles[Cowles$sex == "female", ]
dim(the.females)
## [1] 780    5

the.extraverted <- Cowles[Cowles$extraversion > 18, ]
dim(the.extraverted)
## [1] 73    5
```

In real applications, it is better done via subset() because the basic logical operators treat missing values in a somewhat surprising way (they are included in the subset!).

```
Cowles[1,3] <- NA # set first value of Gender column to missing
the.females1 <- Cowles[Cowles$sex == "female", ]
the.females2 <- subset(Cowles, sex == "female")

dim(the.females1) # with conditional expression
## [1] 780 5
dim(the.females2) # using subset()
## [1] 779 5
identical(the.females1, the.females2)
## [1] FALSE</pre>
```

Converting a character variable into a factor:

```
test.factor <- sample(c("Cat","Dog","Alligator"), size = 30, replace = TRUE)</pre>
test.factor
## [1] "Dog"
                                                   "Cat"
                  "Dog"
                             "Dog"
                                        "Dog"
## [6] "Alligator" "Cat"
                             "Dog"
                                        "Cat"
                                                   "Alligator"
## [11] "Dog" "Cat"
                             "Alligator" "Dog"
                                                   "Cat."
## [16] "Cat" "Dog"
                             "Dog"
                                        "Alligator" "Alligator"
## [21] "Alligator" "Alligator" "Cat"
                                                   "Dog"
## [26] "Cat" "Dog"
                         "Alligator" "Alligator" "Dog"
str(test.factor)
## chr [1:30] "Dog" "Dog" "Dog" "Dog" "Cat" "Alligator" ...
xtabs(~test.factor)
## test.factor
## Alligator
                 Cat
                          Dog
        10
##
                           12
```

```
test.factor1 <- factor(test.factor,</pre>
                       levels = c("Alligator", "Cat", "Dog"),
                       labels = c("Alligator", "Cat", "Dog"))
str(test.factor1)
## Factor w/ 3 levels "Alligator", "Cat", ...: 3 3 3 3 2 1 2 3 2 1 ...
xtabs(~test.factor1)
## test.factor1
## Alligator
                   Cat
                             Dog
##
          10
                              12
test.factor2 <- factor(test.factor)
str(test.factor2)
## Factor w/ 3 levels "Alligator", "Cat", ...: 3 3 3 3 2 1 2 3 2 1 ...
xtabs(~test.factor2)
## test.factor2
## Alligator
                   Cat
                             Dog
##
          10
                              12
```

Converting a continuous variable into a categorical factor:

```
Cowles$ext.factor <- cut(Cowles$extraversion, 3)
str(Cowles$ext.factor)
## Factor w/ 3 levels "(1.98,9]","(9,16]",...: 2 2 2 3 3 2 2 2 2 1 ...
levels(Cowles$ext.factor)
## [1] "(1.98,9]" "(9,16]" "(16,23]"
levels(Cowles$ext.factor) <- c("Low", "Medium", "High")
str(Cowles$ext.factor)
## Factor w/ 3 levels "Low","Medium",...: 2 2 2 3 3 2 2 2 2 1 ...</pre>
```

Tips and Tricks:

- When importing with read.csv(), the following flags may be useful:
 - header = TRUE (variable names included in first row)
 - sep = "" (useful if dataset uses odd delimiter)
 - na.strings = c("", " ", ".", "NA"))
 - stringsAsFactors = FALSE
 - fileEncoding = "UTF-8-BOM"
- If you have transformed variables or created new ones, you might want to save the final datafile.
 - Use: write.csv(dataframe, "dataframe.csv", row.names = FALSE)
 - Use getwd() to see where R is saving output to.

Exercises, Pt. 3

Basic R Exercises, Pt. 3

We are now ready to attempt the Exercise questions pertaining to "Factors and Data Frames".

Intermediate R

Intermediate R

Programming Basics

IF statements evaluate a conditional statement, and return an expression if the value is true.

```
if (cond) expr
if (cond) expr1 else expr2

if (cond) {
    expr1
} else {
    expr2
}
```

```
x <- 5
if (x == 5) print("X is equal to 5")
## [1] "X is equal to 5"
if (x == 6) print("X is 6") else print("X is not 6")
## [1] "X is not 6"</pre>
```

FOR statements: Loop over a vector and perform an expression for each value.

```
for (var in seq) expr
```

```
for (i in 1:5) print(i)
## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
```

WHILE statements: Loop over elements so long as the condition is TRUE.

```
while (cond) expr
```

```
x <- 1
while(x < 8) {
    print(paste("The value of X is:", x))
    x <- x + 1
}
## [1] "The value of X is: 1"
## [1] "The value of X is: 2"
## [1] "The value of X is: 3"
## [1] "The value of X is: 4"
## [1] "The value of X is: 5"
## [1] "The value of X is: 6"
## [1] "The value of X is: 6"
## [1] "The value of X is: 7"</pre>
```

SWITCH statements: Return different values depending on input.

```
switch(expr, ...)
```

IFELSE statements: Similar to basic if() but vectorized. These are very useful for recoding data, and can be nested.

```
ifelse
ifelse(test, yes, no)
```

Intermediate R

Intermediate R

Examples of Applications

Splitting Samples

Creating Exploratory vs. Confirmatory Samples

```
round(prop.table(table(Cowles$sex, Cowles$volunteer))*100, 1)
##
##
              no yes
##
    female 30.3 24.6
    male 27.7 17.5
##
library('devtools')
source gist("https://gist.github.com/mrdwab/6424112")
Cowles.EFA <- stratified(df = Cowles, group = c("sex", "volunteer"), size = .5)
Cowles.CFA <- Cowles[!rownames(Cowles) %in% rownames(Cowles.EFA). ]
round(prop.table(table(Cowles.EFA$sex, Cowles.EFA$volunteer))*100, 1)
##
##
              no ves
    female 30.3 24.5
##
##
    male 27.6 17.5
round(prop.table(table(Cowles.CFA$sex, Cowles.CFA$volunteer))*100, 1)
##
##
              no yes
    female 30.2 24.6
##
##
    male 27.7 17.4
```

Basic Statistical Analyses

The base **stats** package has many useful functions for basic statistical analyses, such as:

- prop.test(), xtabs() (cross-tabs and χ^2), and t.test()
- cor(), and cor.test()
- anova(), and manova()
- lm(), and glm() (linear and generalized linear models)
- influence() (calculate DFBETAS, DFFITS, covariance ratios, Cook's Distance, hat-values)
- cov2cor() (convert covariance matrix to correlation matrix)
- p.adjust() (adjust p-values for multiple comparisons)
- residuals() (extract model residuals)

Basic Statistical Analyses

Most statistical functions utilize the "formula interface", which refers to a structured string with the following properties:

- the response variable(s) appear on the left-hand side of the equation
- the predictors or indicator variable(s) appear on the right-hand side of the equation, separated by +
- the two sides are separated by a tilde (~)

All functions that use the formula interface have a data = argument, which is very useful. For example, compare:

```
response ~ var1 + var2, data = data.name
data.name$response ~ data.name$var1 + data.name$var2
```

Additionally, these characters have specific behaviours:

- * denotes all possible interactions (e.g., sex * volunteer includes main effects for sex, volunteer, as well as their crossing, while adding extraversion would include the three-way interaction, and all second order interactions)
- ∧ is used to incorporate variable crossings to the specified degree (e.g. extraversion∧2 includes extraversion and extraversion²)
- : is used to denote particular crossings. For example, sex:volunteer will add their interaction.
- is used to remove a particular effect from the model (e.g., sex * volunteer * extraversion sex:extraversion)

Advanced Tip: Adding Many Variables

```
xnam <- paste0("x", 1:25) # vector contains x1, ..., x25
(fmla <- as.formula(paste("y ~ ", paste(xnam, collapse= "+"))))
## y ~ x1 + x2 + x3 + x4 + x5 + x6 + x7 + x8 + x9 + x10 + x11 +
## x12 + x13 + x14 + x15 + x16 + x17 + x18 + x19 + x20 + x21 +
## x22 + x23 + x24 + x25

xnam <- paste0(names(Cowles[1:3]))
(fmla <- as.formula(paste("volunteer ~ ", paste(xnam, collapse= "+"))))
## volunteer ~ neuroticism + extraversion + sex</pre>
```

- pasteO() combines element-wise, but with no spaces between
- paste() combines element-wise, but with whitespace between elements
- collapse argument is used to define what separates pasted elements

Linear Regression:

```
out <- lm(neuroticism ~ sex * volunteer + extraversion, data = Cowles)
summary(out)
##
## Call:
## lm(formula = neuroticism ~ sex * volunteer + extraversion. data = Cowles)
##
## Residuals:
            10 Median 30
                                   Max
##
      Min
## -13.0378 -3.5584 -0.0378 3.7029 11.8950
##
## Coefficients:
                    Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                    13.84484 0.45817 30.217 < 2e-16 ***
## sexmale
            -1.72595 0.33474 -5.156 2.88e-07 ***
                  0.22142 0.34693 0.638
## volunteeryes
                                               0.523
                   ## extraversion
## sexmale:volunteeryes -0.15936   0.52032 -0.306   0.759
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 4.796 on 1415 degrees of freedom
    (1 observation deleted due to missingness)
## Multiple R-squared: 0.04429, Adjusted R-squared: 0.04159
## F-statistic: 16.39 on 4 and 1415 DF. p-value: 3.888e-13
```

The output from most analyses can be passed to other functions, and will return different things. For example, try:

```
print(out)  # brief summary
summary(out)  # expanded summary
anova(out)  # ANOVA table
coef(out)  # model coefficients
head(resid(out))  # model residuals
plot(out)  # regression diagnostic plots
```

Logistic Regression:

```
out <- glm(volunteer ~ sex + extraversion, data = Cowles, family = binomial)
summary(out)
##
## Call:
## glm(formula = volunteer ~ sex + extraversion, family = binomial,
##
      data = Cowles)
##
## Deviance Residuals:
      Min
                1Q Median
                                         Max
##
                                 3Q
## -1.3873 -1.0509 -0.9034 1.2522 1.6855
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -1.02592 0.19181 -5.349 8.87e-08 ***
## sexmale -0.24902 0.10932 -2.278 0.0227 *
## extraversion 0.06552 0.01417 4.622 3.79e-06 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 1932.4 on 1419 degrees of freedom
## Residual deviance: 1905.2 on 1417 degrees of freedom
     (1 observation deleted due to missingness)
## AIC: 1911.2
##
## Number of Fisher Scoring iterations: 4
```

Graphical Analyses

Let's use the exercise dataset, and make sure it is read in nicely:

```
dat <- read.csv("exerciseDat.csv", header=TRUE, stringsAsFactors=FALSE)
head(dat)
    id gender mathPre readPre group score1 score2 pass
## 1 1
              55.15 56.31 mnemonic 48.33 68.74
## 2 2
           1 53.30 30.45 mnemonic 45.91 43.89
## 3 3 1 56.96 69.53
                              cram 55.98 84.43
## 4 4 1 55.64 38.25 mnemonic 52.81 38.24
## 5 5
          1 48.14 60.25 mnemonic 53.70 54.46
## 6 6
           0 49.07 61.91 cram 46.30 61.31
str(dat)
## 'data.frame': 300 obs. of 8 variables:
## $ id
           : int 1 2 3 4 5 6 7 8 9 10 ...
## $ gender : int 0 1 1 1 1 0 1 0 0 0 ...
## $ mathPre: num 55.1 53.3 57 55.6 48.1 ...
## $ readPre: num 56.3 30.4 69.5 38.2 60.2 ...
## $ group : chr "mnemonic" "mnemonic" "cram" "mnemonic" ...
## $ score1 : num 48.3 45.9 56 52.8 53.7 ...
## $ score2 : num 68.7 43.9 84.4 38.2 54.5 ...
## $ pass : int 1010111100...
```

Graphical Analyses

```
dat$gender <- factor(dat$gender, levels = c(0,1), labels = c("Female", "Male"))</pre>
dat$group <- factor(dat$group, levels = c("cram", "mnemonic", "none"),</pre>
                   labels = c("cram", "mnemonic", "none"))
dat$pass <- factor(dat$pass, levels = c(0.1), labels = c("Fail", "Pass"))
str(dat)
## 'data.frame': 300 obs. of 8 variables:
## $ id : int 1 2 3 4 5 6 7 8 9 10 ...
## $ gender : Factor w/ 2 levels "Female". "Male": 1 2 2 2 2 1 2 1 1 1 ...
## $ mathPre: num 55.1 53.3 57 55.6 48.1 ...
## $ readPre: num 56.3 30.4 69.5 38.2 60.2 ...
## $ group : Factor w/ 3 levels "cram", "mnemonic",...: 2 2 1 2 2 1 1 2 2 1 ...
## $ score1 : num 48.3 45.9 56 52.8 53.7 ...
## $ score2 : num 68.7 43.9 84.4 38.2 54.5 ...
## $ pass : Factor w/ 2 levels "Fail", "Pass": 2 1 2 1 2 2 2 2 1 1 ...
xtabs(~ gender + group, data = dat)
##
          group
## gender cram mnemonic none
## Female 36
                      32 36
## Male 69
                     72 55
```

Graphical Analyses

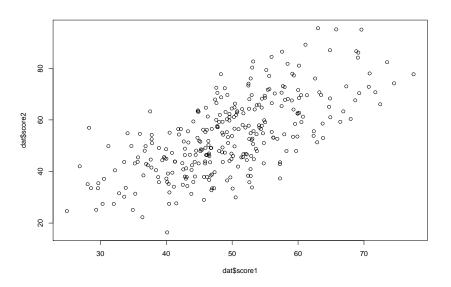
In R there are multiple ways to produce graphics. Two major approaches are using built-in functions (base), or using additional packages.

Advantages of base graphics:

- No need to include library() in your code
- Very quick to produce a plot (e.g., hist(dat\$mathPre))
- plot() is a generic function, meaning it will produce different plots depending on what it was passed.
 - For example: plot(dat\$score1, dat\$score2)
 - However, often this is not useful: plot(dat\$score1)

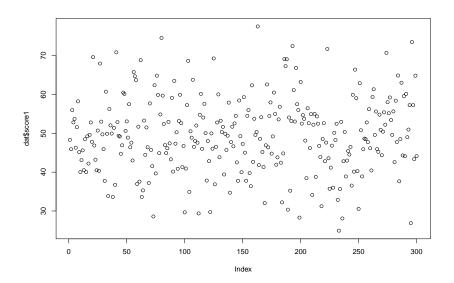
Base Graphics

plot(dat\$score1, dat\$score2)



Base Graphics

plot(dat\$score1)



ggplot2 Funamentals

A substantially more powerful package for plotting is Hadley Wickham's ggplot2.

General principals for using ggplot2:

- Define the data you want to plot and create a plot template object with ggplot()
- Specify the aesthetics of the shapes that will be used to represet the data with aes()
- Specify the graphical shapes (geoms) that will be used to view the data
 - Add them with the appropriate function
 - e.g., geom_point() or geom_line()
- Call the object to render and view it

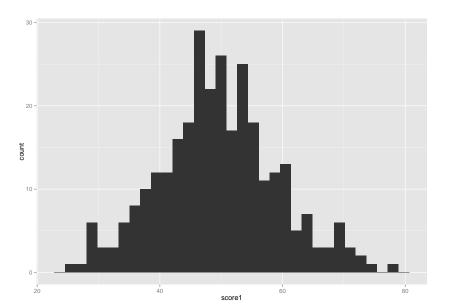
```
#install.packages("ggplot2", dep=T)
library(ggplot2)
```

For starting out with ggplot2, we will use qplot(), which is a convenience function that stands for *quick* plot.

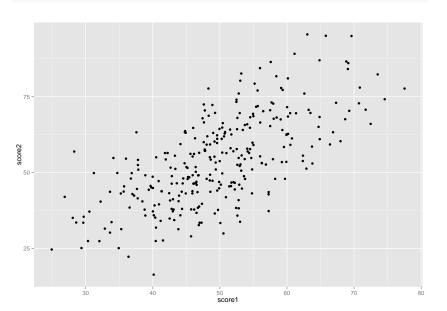
The primary arguments for qplot() are:

- x values we want on the x-axis
- y values we want on the y-axius (not necessary, depending on plot)
- data dataset
- geom geometric object to use ("point" if x and y are given, "histogram" if not)

qplot(score1, data = dat, geom = "histogram")



```
qplot(score1, score2, data = dat, geom = "point")
```



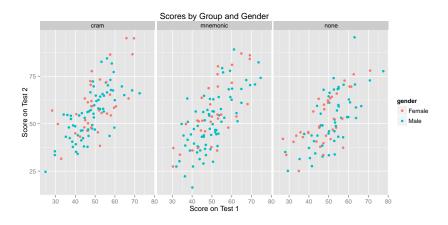
Graphic Manipulation:

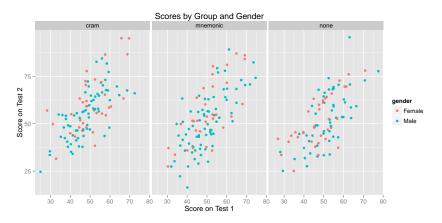
- xlim and ylim: define limits for x and y axis
 - e.g., $\times \lim = c(0, 100)$
- binwidth when creating histograms
- log: log transform any of the variables?
 - Can be x, y, or xy for both
 - However, I usually prefer to transform my variables myself
 - e.g., dat\$score1.log <- log(dat\$score1)

Graphic Manipulation:

- main, xlab, and ylab: title sections of graphic
- colour, size: Differentiate points by grouping variable
- facets: Create separate plots by grouping variable
 - e.g., one variable: facets = ~ gender
 - e.g., two variables: factets = group ∼ gender
- theme: allows other customizations
 - try adding + theme_bw() for black and white graphics
 - see this link for additional details

```
qplot(score1, score2, data = dat, colour = gender,
    facets = ~ group, main = "Scores by Group and Gender",
    xlab = "Score on Test 1", ylab = "Score on Test 2")
```

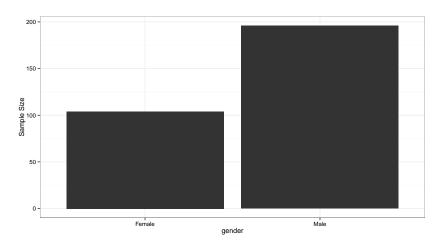




Of course, there are additional geoms we can use!

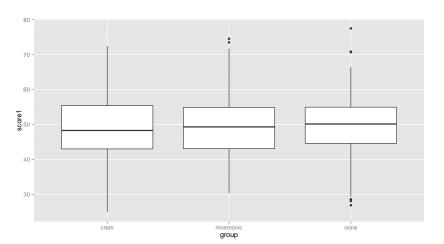
Bargaphs:

```
qplot(gender, data = dat, geom = "bar", ylab = "Sample Size") + theme_bw()
```



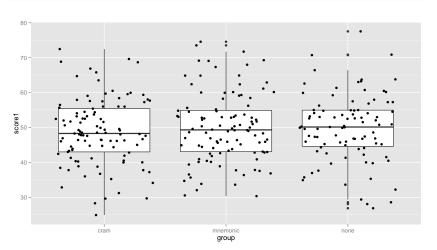
Boxplots:

```
qplot(group, score1, data = dat, geom="boxplot")
```



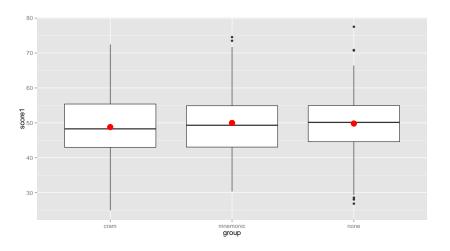
Geoms can be overlapped:

```
qplot(group, score1, data = dat, geom=c("boxplot", "jitter"))
```



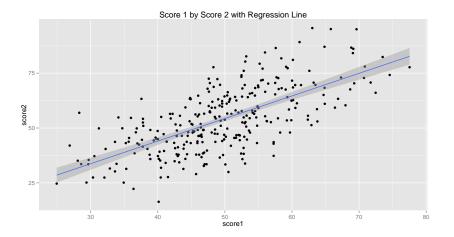
Geoms can be overlapped:

```
qplot(group, score1, data=dat, geom = 'boxplot') +
    stat_summary(fun.y=mean, colour="red", size=5, geom="point")
```



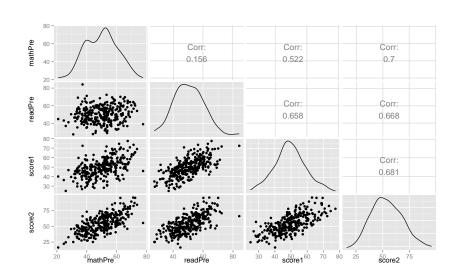
Geoms can be overlapped:

```
qplot(score1, score2, data = dat, geom = c("point", "smooth"), method = "lm",
    main = "Score 1 by Score 2 with Regression Line")
```



Again, with many numeric variables, we are often interested in seeing all combinations at once:

```
#install.packages("GGally")
library(GGally)
ggpairs(na.omit(dat[c(3,4,6,7)]))
```

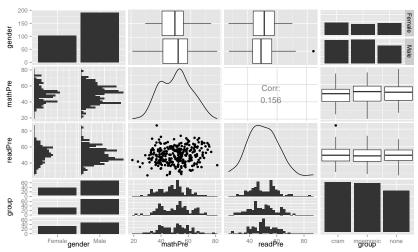


This actually works on practically all variable types!

```
library(GGally)
ggpairs(na.omit(dat[2:5]))
# Ignore id, but use first four variables in
# the dataset as a demonstration
```

- gender (factor)
- mathPre (numeric)
- readPre (numeric)
- group (factor)

This actually works on practically all variable types!



ggplot2 Proper

Again, qplot() is a convenience function for learning the ggplot2 basics.

It should suffice for the purposes of this workshop, but if you want more control over your graphics (or to use more esoteric geoms), using the primary ggplot() function allows for that.

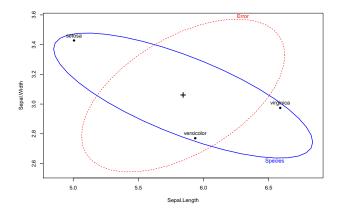
Exercises, Pt. 4

Basic R Exercises, Pt. 4

We are now ready to attempt the Exercise questions pertaining to "Graphing and Data Frames".

Alternate Packages for Graphical Output

Categorical data: vcd (mosaic(), agreementplot(), fourfold()). Multivariate models (e.g., MANOVA/multivariate regression): heplots, uses ellipses to represent hypothesis and error SSCP matrices.



Polychoric Correlations

Variety of packages:

- John Fox's polycor:::polychor()
 - Only calculates pairwise correlations one at a time
 - Can use polycor:::hetcor() (all categorical variables must be declared as factors, or else will be treated as continuous)
- William Revelle's psych:::polychoric()
 - If not all variables are categorical, can also use psych:::mixed.cor()

Exploratory Factor Analysis

Two choices for packages:

- Base R's stats:::factanal() (maximum-likelihood only)
- William Revelle's psych:::fa() (use fa.poly() for dictotomous/polytomous items)
 - The psych functions have additional support for multiple rotation, scoring, and factoring methods.

```
fa(r, nfactors=1, n.obs = NA, n.iter=1, rotate="oblimin",
scores="regression", residuals=FALSE, SMC=TRUE, covar=FALSE,
missing=FALSE, impute="median", min.err = 0.001, max.iter = 50,
symmetric=TRUE, warnings=TRUE, fm="minres", alpha=.1,p=.05,
oblique.scores=FALSE, np.obs, use="pairwise", ...)
```

- Rotation: 8 orthogonal, 7 oblique rotation options
- Factoring Method: minres (minimum residual, OLS), wls (weighted least squares), gls (generalized weighted least squares), pa (principal factors), ml (maximum likelihood), minchi (minimize sample size weight chi square).
- Factor Scores: regression, correlation preserving (tenBerge), Anderson, and Bartlett.

Exploratory Factor Analysis

Supplementary:

- faoutlier is a great package for running influential case diagnostics for factor analysis and SEM.
 - gCD(): Generalized Cook's Distance.
 - LD(): Likelihood Distance.
 - robustMD(): Robust Mahalanobis Distances.
 - obs.resid(): Model predicted residual outliers.
 - Works with fa, sem, and lavaan models.
- nFactors is useful for determining appropriate number of factors.
 - Implements parallel analysis, as well as 11 other criterions.

Confirmatory Factor Analysis and SEM

Recommended package: lavaan, whose primary functions are:

- lavaan() general framework for all latent variable models
- sem() defaults changed to accommodate SEM models
- cfa() defaults changed to accommodate CFA models

...and are supplemented by:

- fitMeasures() return a variety of fit measures
- coef() return coefficients from model
- residuals() return covariance/mean residuals from model
- parameterEstimates() view paramater estimates
- parTable() view model parameters as a dataframe
- modificationIndices() calculate modification indices
- inspect() useful for extracting particular output elements
- anova() likelihood ratio test for nested models

lavaan Model Syntax

Models in lavaan are plain text strings, but follow particular conventions.

- Use =~ to define latent varianbles
- Use ~ to define regressions
- Use ~~ to define (co)variances
- Use ~ 1 to define intercepts
- Use X* to fix a parameter to X
- Use equal() to contrain parameters to be equal

lavaan Model Syntax

Basic Template for lavaan model syntax:

```
myModel <- '
  # latent variable definitions
    f1 = v1 + v2 + v3
    f2 = y4 + y5 + y6
    f3 = v7 + v8 +
         y9 + y10
  # regressions
    f1 \sim f3 + f4
    f2 ~ f4
    y1 + y2 \sim x1 + x2 + x3
  # (co)variances
    y1 ~~ y1
    y2 \sim y4 + y5
    f1 ~~ f2
  # intercepts
    f1 ~ 1; y5 ~ 1 '
```

The Holzinger and Swineford (1939) Example:

```
summary(fit)
## lavaan (0.5-17) converged normally after 35 iterations
##
##
     Number of observations
                                                        301
##
##
     Estimator
                                                         ML.
##
     Minimum Function Test Statistic
                                                    85.306
##
     Degrees of freedom
                                                         24
##
     P-value (Chi-square)
                                                     0.000
##
## Parameter estimates:
##
##
     Information
                                                  Expected
##
     Standard Errors
                                                  Standard
##
##
                      Estimate Std.err Z-value P(>|z|)
## Latent variables:
##
     visual =~
##
       x1
                          1,000
##
       x2
                         0.554
                                   0.100
                                            5.554
                                                     0.000
##
       x3
                         0.729
                                   0.109
                                            6.685
                                                     0.000
##
     textual =~
##
       ×4
                          1.000
##
       x5
                         1.113
                                   0.065
                                           17.014
                                                     0.000
##
       x6
                         0.926
                                   0.055
                                           16.703
                                                     0.000
##
     speed =~
##
       x7
                         1.000
##
       x8
                          1.180
                                   0.165
                                            7.152
                                                     0.000
##
       x9
                          1.082
                                   0.151
                                            7.155
                                                     0.000
##
## Covariances:
```

Some general starting parameters:

- Correlated Factors: use cfa(), with: std.lv = TRUE, std.ov = TRUE.
- Hierarchical Model: use cfa(), with: 'std.lv = TRUE, std.ov = TRUE, and orthogonal = TRUE.
- **Bifactor Model**: use cfa(), with: 'std.lv = TRUE, std.ov = TRUE, and orthogonal = TRUE.

Reference Guide:

Beaujean, A.A. (2014). Latent Variable Modeling using R: A Step-By-Step Guide. Routledge Publishing.

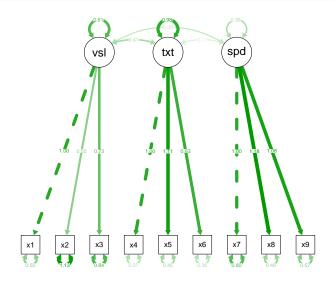
Supplementary:

- **lavaan.survey** has additional functions for dealing with complex sampling design.
- John Fox wrote the sem package, which is computationally faster although less user friendly

Plotting CFA/SEM Path Diagrams

 $Recommendation: \ \textbf{semPlot}, \ using \ function \ \texttt{semPaths()} \ to \ plot \ model.$

```
library(semPlot)
semPaths(fit, what = "est")
```



(Multidimensional) Item Response Theory

Recommended package: mirt

- mirt() primary function
- bfactor() shorthand for setting up bifactor models
- multipleGroup() shorthand for MG models

Supplemented by:

- mirt.model() create mirt model object
- testinfo() calculate test information/plot
- itemplot() obtain item characteristic curves
- iteminfo() calculate item information
- itemfit() calculate item fit statistics
- fscores() calculate factor scores (MAP, EAP, ML, or WLE)
- wald() calculate Wald log-likelihood tests across models
- DIF() investigate differential item functioning for MG models
- itemplot(shiny=TRUE) interactive visualization of MIRT models

(Multidimensional) Item Response Theory

MIRT Model Syntax

- Simply entering a number will estimate an exploratory IRT model with that many factors.
 - mod1 <- mirt(data, 1)</pre>
- Otherwise, model is specified by item number or variable name:

MIRT Parameters

- itemtype = Rasch, 2-4PL, graded, grsm, gpcm/nominal, ideal, partcomp, 2-4PLNRM
- method = for unidimensional: EM, for MIRT: QMCEM, MHRM
- see ?mirt for more details

General Psychometrics

The psych library

- William Revelle's psych library has a many useful functions, primarily for scale construction using factor analysis, cluster analysis and reliability analysis, although others provide basic descriptive statistics.
- Some support for Item Response Theory is done using factor analysis of tetrachoric and polychoric correlations.
- Functions for simulating particular item and test structures are included.
- More details at the Personality Project website

General Psychometrics

CRAN's psychometric Task View

- Maintained by Patrick Mair (Harvard), last updated February 25th, 2015.
- Subheadings for: Item Response Theory, Correspondence Analysis, SEM/FA/PCA, Multidimensional Scaling, Classical Test Theory, Knowledge Structure Analysis, and Other Related Packages
- Link: Psychometric Models and Methods

Life on the Edge

Most R packages have two lives: the extensively tested "stable" build that lives on CRAN, and the cutting edge "developmental" build that may live outside of CRAN. Why go developmental?

- Cutting edge features
- Recent bug fixes

mirt in particular uses the developmental branch of their GitHub repository frequently. To install a package from source you need:

- On Windows, you must install RTools.exe; on Mac OS X you may need to install XCode's Command Line Tools.
- Run install.packages("devtools")
- load.packages("devtools")
- install_github('philchalmers/mirt')

Some packages **only** live on the fringes (insert grumble about draconian CRAN policies), so knowing how to install them can be very important!

Write It Yourself!

One of the best parts about R is how easy it is to create your own functions and packages. These can become community-wide staples (for example, the package Hmisc is literally a collection of Frank Harrell's miscellaneous functions.).

For instance: Manolo wrote a script for calculating empirical percentiles.

Emprical Percentiles

SET-UP

- The Hmisc packange needs to be loaded
- Variables should be imported as INTEGERS
- Variables must be defined as FACTORS
- Must source() in Manolo's functions
 - source("G:/R&D/_PRIVATE SHARED/R/EmPeRnd.R")

Documentation

The documentation for EmPeRnd() and its helper functions is in the .R file: G:/R&D/ PRIVATE SHARED/R/EmPeRnd.R

Source script and datafile:

```
library(Hmisc)
source("G:/R&D/_PRIVATE SHARED/R/EmPeRnd.R")
mp <- spss.get("G:/R&D/_PRIVATE SHARED/R/MASC2.sav")</pre>
```

Round variables to integers:

```
mp$ROUND.TOT <- round(mp$MASC.TOTAL.50)
mp$ROUND.SEP <- round(mp$Raw.Sep)
mp$ROUND.GAD <- round(mp$CAD.INDEX.SEP07)
mp$ROUND.SANX <- round(mp$Raw.Sanx)
mp$ROUND.HR <- round(mp$Raw.HR)
mp$ROUND.PF <- round(mp$Raw.PP)
mp$ROUND.OCD <- round(mp$CD.AUG15)
mp$ROUND.PHYS <- round(mp$Raw.Phys)
mp$ROUND.PANIC <- round(mp$PANIC)
mp$ROUND.TR <- round(mp$PANIC)
mp$ROUND.TR <- round(mp$TANIC)
mp$ROUND.TR <- round(mp$TANIC)
mp$ROUND.HA <- round(mp$TANIC)
```

Convert to Factors:

```
mp$ROUND.TOT <- factor(mp$ROUND.TOT, levels=150:0)
mp$ROUND.SEP <- factor(mp$ROUND.SEP, levels=27:0)
mp$ROUND.GAD <- factor(mp$ROUND.GAD, levels=30:0)
mp$ROUND.SANX <- factor(mp$ROUND.SANX, levels=27:0)
mp$ROUND.HR <- factor(mp$ROUND.HR, levels=15:0)
mp$ROUND.PF <- factor(mp$ROUND.PF, levels=12:0)
mp$ROUND.OCD <- factor(mp$ROUND.PF, levels=30:0)
mp$ROUND.PHYS <- factor(mp$ROUND.PHYS, levels=36:0)
mp$ROUND.PANIC <- factor(mp$ROUND.PANIC, levels=21:0)
mp$ROUND.TR <- factor(mp$ROUND.HR, levels=15:0)
mp$ROUND.TA <- factor(mp$ROUND.HA, levels=15:0)
```

Create loop to calculate the percentiles (with EmPeRnd()):

In this example, the scores we want to create percentiles for are in columns 80:90, and the weighting values are in the variable WT.ALL.FINAL.

What does our output look like?

```
head(QA.emptile.p) # Full output, inc. weighted frequency & raw percentiles.
                 rawper per per.txt scl
## 0 0 0.0000000 0.0000000 1 1st TOT
## 1 1 0.0000000 0.0000000 1 1st TOT
## 2 2 0.0000000 0.0000000 1 1st TOT
## 3 3 2.8880424 0.0712042 1 1st TOT
## 4 4 0.4615546 0.1537880
                          1 1st TOT
## 5 5 0.3653974 0.1741763 1 1st TOT
head(emptile.p) # Truncated output.
    scl l per per.txt
## 0 TOT 0 1
                 1st
## 1 TOT 1 1
                1st
## 2 TOT 2 1
             1st
## 3 TOT 3 1
             1st
## 4 TOT 4 1
             1st
## 5 TOT 5
                1st
```

Write output to file:

```
write.csv(emptile.p, file="emptile.p.csv", row.names=F)
write.csv(QA.emptile.p, file="QA.emptile.p.csv", row.names=F)
```

What does a custom function look like?

```
EmPeRnd <- function(wth, fact) {</pre>
  #EmPeRnd : calculates rounded empirical percentiles
  #*Requires* functions WtdFreq and stndrdth
  #'wth' is the weight variable
  #'fact' is the score or factor for the rows
  # Returns a dataframe with values:
  # 'l': the "levels" of 'fact' (it includes all levels including with 0 freq)
  # 'f' : the weighted frequencies of the levels/values of 'fact'
  # 'rawper' the empirical percentile
  # 'per' the truncated empirical percentile (with the plus.one)
  # 'per.txt' the truncated empirical percentile in ordinal text format
wf <- WtdFreq(wth, fact)
tot <- sum(wf$f)
ln \leftarrow dim(wf)[1]
for (i in 1:ln) {
 if(i==1) {
    wf[i,"rawper"] <- wf[i,"f"]/2/tot*100
    } else {
      wf[i,"rawper"] \leftarrow ((wf[i,"f"]/2) + sum(wf[1:i-1,"f"]))/tot*100
wf$per <- round(wf$rawper)</pre>
wf$per <- ifelse(wf$per < 1, 1, ifelse(wf$per >99, 99, wf$per))
wf$per.txt <- stndrdth(wf$per)</pre>
return(wf)
```

Magic Pipes

A recent addition to R is the magrittr package, whose main contribution is the addition of the pipe operator: %>%.

In basic R code, functions are evaluated from the inside out. For example:

```
library(dplyr)
head(arrange(group_by(select(mutate(iris,
Sepal.Size=Sepal.Length*Sepal.Width,
Petal.Size=Petal.Length*Petal.Width),
Sepal.Size, Petal.Size, Species), Species),
desc(Sepal.Size+Petal.Size)), n = 3)
```

What is this doing?

Magic Pipes

With some indenting, it gets a bit better:

```
## Source: local data frame [3 x 3]
## Groups: Species
##
## Sepal.Size Petal.Size Species
## 1 25.08 0.60 setosa
## 2 23.20 0.24 setosa
## 3 23.10 0.28 setosa
```

Magic Pipes

The pipe, on the other hand, takes what is on the left and "pipes" its output to the next function. This can greatly enhance the readability of your code.

```
library(magrittr)
iris %>%
 mutate(Sepal.Size=Sepal.Length*Sepal.Width,
         Petal.Size=Petal.Length*Petal.Width) %>%
 select(Sepal.Size, Petal.Size, Species) %>%
 group_by(Species) %>%
 arrange(., desc(Sepal.Size+Petal.Size)) %>%
 head(3)
## Source: local data frame [3 x 3]
## Groups: Species
##
##
    Sepal.Size Petal.Size Species
## 1
         25.08
                     0.60 setosa
## 2
         23.20
                   0.24 setosa
## 3
         23.10
                     0.28 setosa
```

Summary!

- Base R is extremely powerful (and free)
- RStudio improves the usability of R immenseley (and is also free)
- Additional packages can extend the software in many directions, with techniques appearing in R in advance of other software programs.
- The R language has some fundamental learning blocks, but will tend to "click" with perserverance.
- Once you have a solid repository of scripts, it is easy to adapt code from one project to another.
- R help is fairly accessible, and there are many (free) resources available.
- Knit reports provide a succinct summary of what you coded, and how to interpret it.

References

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