#### R Basics Fundamental Techniques in Data Science



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The R Statistical Programming Language

Data I/O

Functions

Iteration



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This course was originally developed by Gerko Vink. You can access the original version of these materials on Dr. Vink's GitHub page: <a href="https://github.com/gerkovink/fundamentals">https://github.com/gerkovink/fundamentals</a>. The course materials

have been (extensively) modified. Any errors or inaccuracies introduced via these modifications are fully my own responsibility and shall not be taken as representing the views and/or beliefs of Dr. Vink. You can see

Gerko's version of the course on his personal website: https://www.gerkovink.com/fundamentals.



# What is "Open-Source"?

R is an open-source software project, but what does that mean?

- Source code is freely available to anyone who wants it.
  - Free Speech, not necessarily Free Beer
- Anyone can edit the original source code to suit their needs.
  - Ego-less programming
- Many open source programs are also "freeware" that are available free of charge.
  - R is both open-source and freeware

# What is R?

I prefer to think about R as a *statistical programming language*, rather than as a data analysis program.

- R IS NOT its GUI (no matter which GUI you use).
- You can write R code in whatever program you like (e.g., RStudio, EMACS, VIM, Notepad, directly in the console/shell/command line).
- R can be used for basic (or advanced) data analysis, but its real strength is its flexible programming framework.
  - Tedious tasks can be automated.
  - Computationally demanding jobs can be run in parallel.
  - R-based research *wants* to be reproducible.
  - Analyses are automatically documented via their scripts.

# What is RStudio?

RStudio is an integrated development environment (IDE) for R.

- Adds a bunch of window dressing to R
- Also open-source
- Both free and paid versions

R and RStudio are independent entities.

- You do not need RStudio to work with R.
- You are analyzing your data with R, not RStudio
  - RStudio is just the interface through which you interact with R.



You can download R, for free, from the following web page:

https://www.r-project.org/

Likewise, you can freely download RStudio via the following page:

• https://posit.co/downloads/



R is an interpreted programming language.

- The commands you enter into the R *Console* are executed immediately.
- You don't need to compile your code before running it.
- In this sense, interacting with R is similar to interacting with other syntax-based statistical packages (e.g., SAS, STATA, Mplus).

# Interacting with R

When working with R, you will write *scripts* that contain all of the commands you want to execute.

- There is no "clicky-box" Tom-foolery in R.
- Your script can be run interactively or in "batch-mode", as a self-contained program.

The primary purpose of the commands in your script will be to create and modify various objects (e.g., datasets, variables, function calls, graphical devices).

# Getting Help

Everything published on the Comprehensive R Archive Network (CRAN), and intended for R users, must be accompanied by a help file.

• If you know the name of the function (e.g., <a>anova()</a> ), then execute

```
?anova Or help(anova) .
```

- If you do not know the name of the function, type ?? followed by your search criterion.
  - For example, **??anova** returns a list of all help pages that contain the word "anova".

The internet can also tell you almost everything you'd like to know.

- Sites such as http://www.stackoverflow.com and http://www.stackexchange.com can be very helpful.
- If you google R-related issues, include "R" somewhere in your search string.



Packages give R additional functionality.

- By default, some packages are included when you install R.
- These packages allow you to do common statistical analyses and data manipulation.
- Installing additional packages allows you to perform state-of-the-art statistical analyses.





These packages are all developed by R users, so the throughput process is very timely.

- Newly developed functions and software are readily available
- Software implementations of new methods can be quickly disseminated
- This efficiency differs from other mainstream software (e.g., SPSS, SAS, MPlus) where new methodology may take years to be implemented.

A list of available packages can be found on CRAN.



# Installing & Loading Packages

Install a package (e.g., mice):

install.packages("mice")

There are two ways to load a package into R

library(stats)
require(stats)



#### **Project Management**

Getting a handle on three key concepts will dramatically improve your data analytic life.

- 1. Working directories
- 2. Directory structures and file paths
- 3. RStudio projects



# DATA I/O



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#### R Data & Workspaces

#### R has two native data formats.

```
## Load the built-in 'bfi' data from the 'psychTools' package
data(bfi, package = "psychTools")
```

## Access the documentation for the 'bfi' data
?psychTools::bfi

```
## Define the directory holding our data
dataDir <- "../../data/"</pre>
```

## Load the 'boys' data from the R workspace
## '../../ata/boys.RData'
load(paste0(dataDir, "boys.RData"))

## Load the 'titanic' data stored in R data set
## '../../data/titanic.rds'
titanic <- readRDS(paste0(dataDir, "titanic.rds"))</pre>

# **Delimited Data Types**

```
## Load the 2017 UTMB data from the comma-separated file
## '../../ata/utmb_2017.csv'
utmb1 <- read.csv(paste0(dataDir, "utmb_2017.csv"))</pre>
```

NOTES:

- The read.csv() function assumes the values are separated by commas.
- For EU-formatted CSV files—with values delimited by semicolons—we can use the read.csv2() function.

Reading data in from other stats packages can be a bit tricky. If we want to read SAV files, there are two popular options:

- foreign::read.spss()
- haven::read\_spss()

```
## Load the foreign package:
library(foreign)
```

```
## Use foreign::read.spss() to read '../../../data/mtcars.sav' into a list
mtcars1 <- read.spss(paste0(dataDir, "mtcars.sav"))</pre>
```

```
## Read '../../ata/mtcars.sav' as a data frame
mtcars2 <- read.spss(paste0(dataDir, "mtcars.sav"), to.data.frame = TRUE)</pre>
```

## View the results: mtcars1[1:3]

\$mpg

[1] 21.0 21.0 22.8 21.4 18.7 18.1 14.3 24.4 22.8 19.2 17.8 [12] 16.4 17.3 15.2 10.4 10.4 14.7 32.4 30.4 33.9 21.5 15.5 [23] 15.2 13.3 19.2 27.3 26.0 30.4 15.8 19.7 15.0 21.4

\$cyl

\$disp

[1] 160.0 160.0 108.0 258.0 360.0 225.0 360.0 146.7 140.8
[10] 167.6 167.6 275.8 275.8 275.8 472.0 460.0 440.0 78.7
[19] 75.7 71.1 120.1 318.0 304.0 350.0 400.0 79.0 120.3
[28] 95.1 351.0 145.0 301.0 121.0

head(mtcars2)

	mpg	cyl	disp	hp	drat	wt	qsec	vs	am
1	21.0	6	160	110	3.90	2.620	16.46	V-Shaped	Manual
2	21.0	6	160	110	3.90	2.875	17.02	V-Shaped	Manual
3	22.8	4	108	93	3.85	2.320	18.61	Straight	Manual
4	21.4	6	258	110	3.08	3.215	19.44	Straight	Automatic
5	18.7	8	360	175	3.15	3.440	17.02	V-Shaped	Automatic
6	18.1	6	225	105	2.76	3.460	20.22	Straight	Automatic
	gear	carb	)						
1	4	4	ł						
2	4	4	ł						
3	4	1							
4	3	1							
5	3	2	2						
6	3	1							

head(mtcars3)

	mpg	cyl	disp	hp	drat	wt	qsec	vs	$\mathtt{am}$	gear	carb
1	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
2	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
3	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
4	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
5	18.7	8	360	175	3.15	3.440	17.02	0	0	3	2
6	18.1	6	225	105	2.76	3.460	20.22	1	0	3	1

## Load the packages: library(haven) library(labelled)

## Use haven::read\_spss() to read '../../ata/mtcars.sav' into a tibble
mtcars4 <- read\_spss(paste0(dataDir, "mtcars.sav"))</pre>

head(mtcars4)

#	A tibb	A tibble: 6 x 11									
	mpg	cyl	disp	hp	drat	wt	qsec	vs	5	an	1
	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<0	lbl+lb>	<c< th=""><th>lbl+l&gt;</th></c<>	lbl+l>
1	21	6	160	110	3.9	2.62	16.5	0	[V-Sh~	1	[Man~
2	21	6	160	110	3.9	2.88	17.0	0	[V-Sh~	1	[Man~
3	22.8	4	108	93	3.85	2.32	18.6	1	[Stra~	1	[Man~
4	21.4	6	258	110	3.08	3.22	19.4	1	[Stra~	0	[Aut~
5	18.7	8	360	175	3.15	3.44	17.0	0	[V-Sh~	0	[Aut~
6	18.1	6	225	105	2.76	3.46	20.2	1	[Stra~	0	[Aut~
#	i 2 mo	ore va	riables	s: gear	<dbl;< th=""><th>&gt;, carb</th><th><dbl></dbl></th><th>&gt;</th><th></th><th></th><th></th></dbl;<>	>, carb	<dbl></dbl>	>			

haven::read\_spss() converts any SPSS variables with labels into labelled vectors.

• We can use the labelled::unlabelled() function to remove the value labels.

```
mtcars5 <- unlabelled(mtcars4)</pre>
```

```
head(mtcars5)
```

```
# A tibble: 6 \times 11
   mpg
        cyl disp hp drat wt qsec vs
                                           am
 <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <fct><fct>
 21
          6 160 110 3.9 2.62 16.5 V-Shaped Manual
1
2 21
          6 160 110 3.9 2.88 17.0 V-Shaped Manual
3 22.8 4 108 93 3.85 2.32 18.6 Straight Manual
4 21.4 6 258 110 3.08 3.22 19.4 Straight Automa~
 18.7 8 360 175 3.15 3.44 17.0 V-Shaped Automa~
5
          6
6
 18.1
             225 105 2.76 3.46 20.2 Straight Automa~
 i 2 more variables: gear <dbl>, carb <dbl>
```

mtcars4\$am[1:20]

```
<labelled<double>[20]>: Transmission type
 [1] 1 1 1 0 0 0 0 0 0 0 0 0 0 0 0 0 1 1 1
Labels:
 value
        label
    0 Automatic
         Manual
    1
mtcars5$am[1:20]
 [1] Manual Manual Manual Automatic Automatic
 [6] Automatic Automatic Automatic Automatic
[11] Automatic Automatic Automatic Automatic Automatic
[16] Automatic Automatic Manual Manual
                                          Manual
Levels: Automatic Manual
```

We have two good options for loading data from Excel spreadsheets:

- readxl::read\_excel()
- openxlsx::read.xlsx()

```
## Load the packages:
library(readxl)
library(openxlsx)
```

```
## Check the results from read_excel():
str(titanic2)
tibble [887 x 8] (S3: tbl_df/tbl/data.frame)
$ survived : chr [1:887] "no" "yes" "yes" "yes" ...
$ class : chr [1:887] "3rd" "1st" "1st" ...
$ name : chr [1:887] "Mr. Owen Harris Braund" "Mrs. John Bradley (Flow
$ sex : chr [1:887] "male" "female" "female" ...
$ age : num [1:887] 22 38 26 35 35 27 54 2 27 14 ...
$ siblings_spouses: num [1:887] 1 1 0 1 0 0 0 3 0 1 ...
$ parents_children: num [1:887] 7.25 71.28 7.92 53.1 8.05 ...
```

## Check the results from read.xlsx():
str(titanic3)

'data.frame': 887 obs. of	8 variables:
<pre>\$ survived : chr</pre>	"no" "yes" "yes"
\$ class : chr	"3rd" "1st" "3rd" "1st"
\$ name : chr	"Mr. Owen Harris Braund" "Mrs. John Bradley (Florence B
\$ sex : chr	"male" "female" "female"
\$ age : num	22 38 26 35 35 27 54 2 27 14
<pre>\$ siblings_spouses: num</pre>	1 1 0 1 0 0 0 3 0 1
<pre>\$ parents_children: num</pre>	0 0 0 0 0 0 1 2 0
\$ fare : num	7.25 71.28 7.92 53.1 8.05

## Compare:

all.equal(as.data.frame(titanic2), titanic3)

[1] TRUE

# Workspaces & Delimited Data

All of the data reading functions we saw earlier have complementary data writing versions.



To write SPSS data, the best option is the haven::write\_sav()
function.

write\_sav(mtcars2, paste0(dataDir, "mctars2.sav"))

write\_sav() will preserve label information provided by factor variables and the 'haven\_labelled' class.



The **openxlsx** package provides a powerful toolkit for programmatically building Excel workbooks in R and saving the results.

• Of course, it also works for simple data writing tasks.

# **FUNCTIONS**

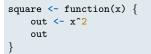
### **R** Functions

Functions are the foundation of R programming.

- Other than data objects, almost everything else that you interact with when using R is a function.
- Any R command written as a word followed by parentheses, () , is a function.
  - mean()
  - o library()
  - o mutate()
- Infix operators are aliased functions.



We can define our own functions using the function() function.



After defining a function, we call it in the usual way.

square(5)

[1] 25

One-line functions don't need braces.

```
square <- function(x) x<sup>2</sup>
square(5)
[1] 25
```

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Function arguments are not strictly typed.

square(1:5)
[1] 1 4 9 16 25
square(pi)
[1] 9.869604
square(TRUE)
[1] 1

But there are limits.

square("bob") # But one can only try so hard Error in x^2: non-numeric argument to binary operator

Functions can take multiple arguments.

```
mod <- function(x, y) x %% y
mod(10, 3)
[1] 1</pre>
```

Sometimes it's useful to specify a list of arguments.

```
getLsBeta <- function(datList) {
   X <- datList$X
   y <- datList$y
   solve(crossprod(X)) %*% t(X) %*% y
}</pre>
```

```
X <- matrix(runif(500), ncol = 5)
datList <- list(y = X %*% rep(0.5, 5), X = X)
getLsBeta(datList = datList)
       [,1]
[1,] 0.5
[2,] 0.5
[3,] 0.5
[4,] 0.5
[5,] 0.5
```

#### **User-Defined Functions**

Functions are first-class objects in R.

• We can treat functions like any other R object.

R views an unevaluated function as an object with type "closure".

```
class(getLsBeta)
```

[1] "function"

typeof(getLsBeta)

[1] "closure"

An evaluated functions is equivalent to the objects it returns.

```
class(getLsBeta(datList))
```

[1] "matrix" "array"

```
typeof(getLsBeta(datList))
```

[1] "double"

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#### **Nested Functions**

We can use functions as arguments to other operations and functions.

```
fun1 <- function(x, y) x + y
```

```
## What will this command return?
fun1(1, fun1(1, 1))
```

[1] 3

Why would we care?

```
s2 <- var(runif(100))
x <- rnorm(100, 0, sqrt(s2))</pre>
```

#### **Nested Functions**

#### X[1:8, ]

[,1][,2][,3][,4][,5][1,]0.524313820.671364470.282287260.71483830.54204681[2,]0.019267420.116937620.091485020.69291710.88371944[3,]0.051007350.184320740.435477990.60974620.09026598[4,]0.605669720.129441270.210001430.24419170.68141473[5,]0.487373030.940304050.239886190.49159100.36353771[6,]0.199419580.966706780.114558200.12439470.24253273[7,]0.955078040.387058290.497335350.29684700.81001800[8,]0.110931970.077317570.849230060.86539870.61914193

c(1, 3, 6:9, 12)

[1] 1 3 6 7 8 9 12

# **ITERATION**



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There are three types of loops in R: for, while, and until.

- You'll rarely use anything but the for loop.
- So, we won't discuss while or until loops.

A for loop is defined as follows.

for(INDEX in RANGE) { Stuff To Do with the Current INDEX Value }



For example, the following loop will sum the numbers from 1 to 100.

val <- 0
for(i in 1:100) {
 val <- val + i
}
val
[1] 5050</pre>

This loop will compute the mean of every column in the mtcars data.

```
means <- rep(0, ncol(mtcars))
for(j in 1:ncol(mtcars)) {
    means[j] <- mean(mtcars[ , j])
}</pre>
```

means

[1] 20.090625 6.187500 230.721875 146.687500 3.596563 [6] 3.217250 17.848750 0.437500 0.406250 3.687500 [11] 2.812500 Loops are often one of the least efficient solutions in R.

```
n <- 1e8
t0 <- system.time({
    val0 <- 0
    for(i in 1:n) val0 <- val0 + i
})
t1 <- system.time(
    val1 <- sum(1:n)
)</pre>
```

#### Loops

Both approaches produce the same answer.

val0 - val1

[1] 0

But the loop is many times slower.

t0 user system elapsed 1.479 0.001 1.480 t1 user system elapsed 0 0 0 There is often a built in routine for what you are trying to accomplish with the loop.

## The appropriate way to get variable means: colMeans(mtcars)

mpg	cyl	disp	hp	drat
20.090625	6.187500	230.721875	146.687500	3.596563
wt	qsec	vs	am	gear
3.217250	17.848750	0.437500	0.406250	3.687500
carb				
2.812500				

# **Apply Statements**

In R, some flavor of *apply statement* is often preferred to a loop.

- Apply statements broadcast some operation across the elements of a data object.
- Apply statements can take advantage of internal optimizations that loops can't use.

There are many flavors of apply statement in R, but the three most common are:

- apply()
- lapply()
- sapply()



## **Apply Statements**

Apply statements generally take one of two forms:

apply(DATA, MARGIN, FUNCTION, ...)

apply(DATA, FUNCTION, ...)



```
## Load some example data:
data(mtcars)
```

```
## Subset the data:
dat1 <- mtcars[1:5, 1:3]</pre>
```

```
## Find the range of each row:
apply(dat1, 1, range)
```

	Mazda RX4 Mazda	RX4 Wag	Datsun 710	Hornet 4 Drive
[1,]	6	6	4	6
[2,]	160	160	108	258
Hornet Sportabout				
[1,]		8		
[2,]	36	50		

## Find the maximum value in each column: apply(dat1, 2, max) mpg cyl disp 22.8 8.0 360.0 ## Subtract 1 from every cell: apply(dat1, 1:2, function(x) x - 1) mpg cyl disp Mazda RX4 20.0 5 159 Mazda RX4 Wag 20.0 5 159 Datsun 710 21.8 3 107 Hornet 4 Drive 20.4 5 257 Hornet 5portabout 17.7 7 359

```
## Create a toy list:
11 <- list()
for(i in 1:3) l1[[i]] <- runif(10)</pre>
## Find the mean of each list entry:
lapply(l1, mean)
[[1]]
[1] 0.526697
[[2]]
[1] 0.4020885
[[3]]
[1] 0.607818
## Same as above, but return the result as a vector:
sapply(11, mean)
```

[1] 0.5266970 0.4020885 0.6078180

## Find the range of each list entry: lapply(l1, range)

[[1]] [1] 0.04395916 0.99350611

[[2]] [1] 0.002797563 0.821082495

[[3]] [1] 0.09926892 0.90430843

sapply(11, range)

[,1] [,2] [,3] [1,] 0.04395916 0.002797563 0.09926892 [2,] 0.99350611 0.821082495 0.90430843

We can add additional arguments needed by the function.

• These arguments must be named.

apply(dat1, 2, mean, trim = 0.1)
 mpg cyl disp
 20.98 6.00 209.20
 sapply(dat1, mean, trim = 0.1)
 mpg cyl disp
 20.98 6.00 209.20

#### Some Programming Tips

You can save yourself a great deal of heartache by following a few simple guidelines.

- Keep your code tidy.
- Use comments to clarify what you are doing.
- When working with functions in RStudio, use the TAB key to quickly access the documentation of the function's arguments.
- Give your R scripts and objects meaningful names.
- Use a consistent directory structure and RStudio projects.

#### **General Style Advice**

Use common sense and BE CONSISTENT.

- Browse the tidyverse style guide.
  - The point of style guidelines is to enforce a common vocabulary.
  - You want people to concentrate on *what* you're saying, not *how* you're saying it.
- If the code you add to a project/codebase looks drastically different from the extant code, the incongruity will confuse readers and collaborators.

Spacing and whitespace are your friends.

- a<-c(1,2,3,4,5)
- a <- c(1, 2, 3, 4, 5)
- At least put spaces around assignment operators and after every comma!

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