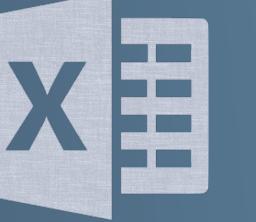


FROM  EXCEL TO 



WHO ARE WE?

1. Center for Health Data Science (HeaDS) - <https://heads.ku.dk/>

SUND Center, which includes a KU data lab

- Courses & Workshops, Seminars, etc.
- Health DS Consultations
- Commissioned Research
- Matchmaking
- Commissioned Supervision



Thilde Terkelsen ¹



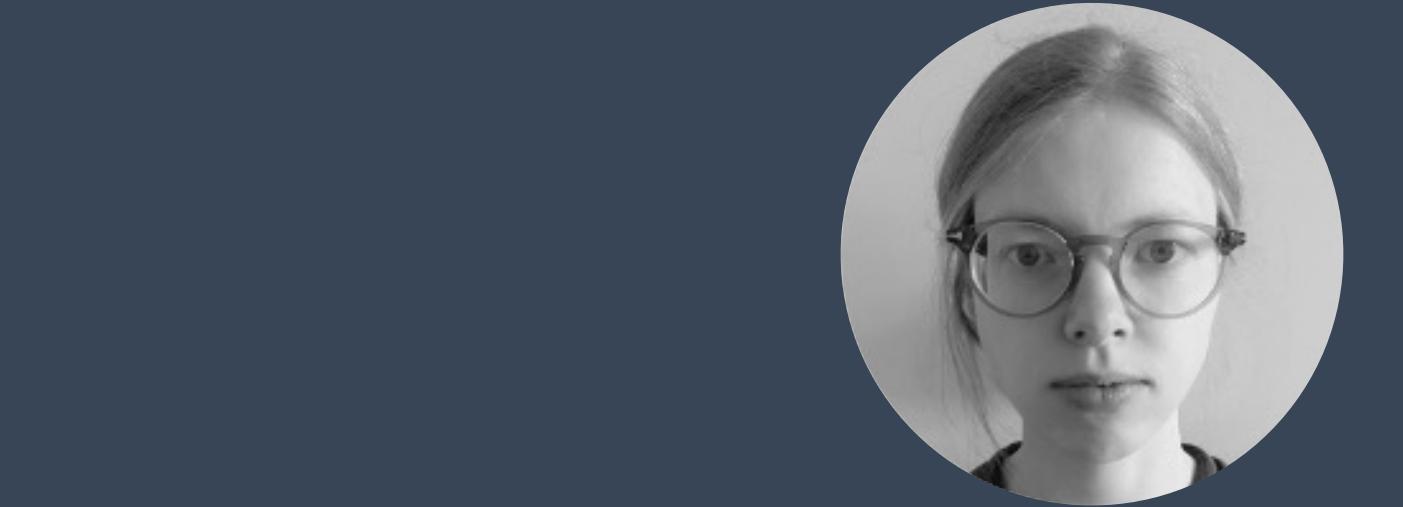
Diana Andrejeva ¹



Tugce Karaderi ¹



Henrike Zschach ¹



Adrija Kalvisa ²

2. ReNEW NNF Center for Stem Cell Medicine

3. Data Science Laboratory (DSL) - <https://datalab.science.ku.dk/>



— FROM EXCEL TO R

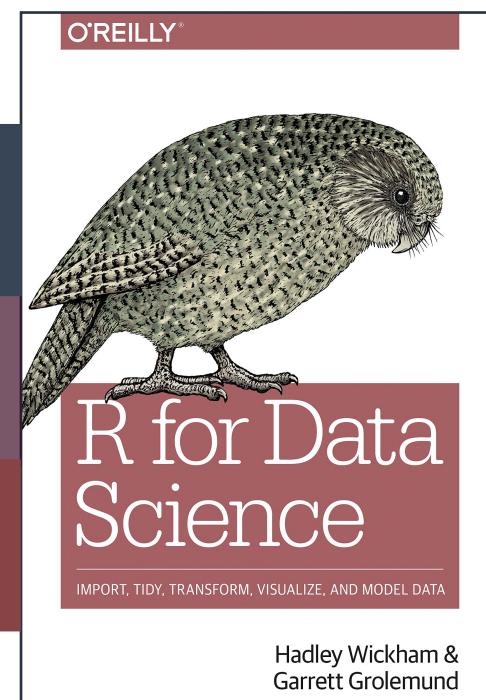
THE PRACTICALS



Two days: 9.00-16.30. There will be coffee breaks, we promise ☕

“R for Data Science” - a generally useful book on R, also for this course

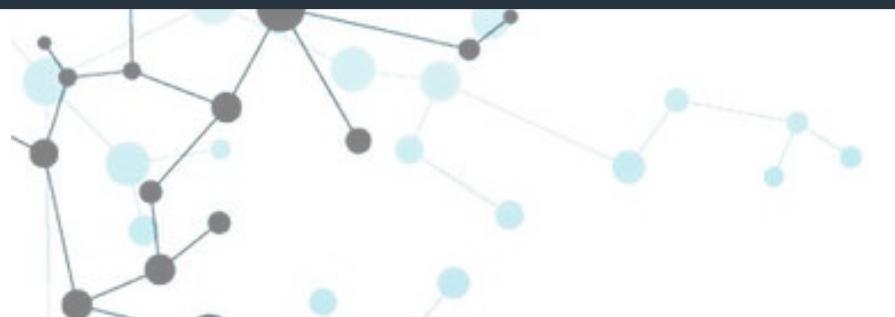
The course is build on hands-on presentations (.R, .Rmd) & exercises



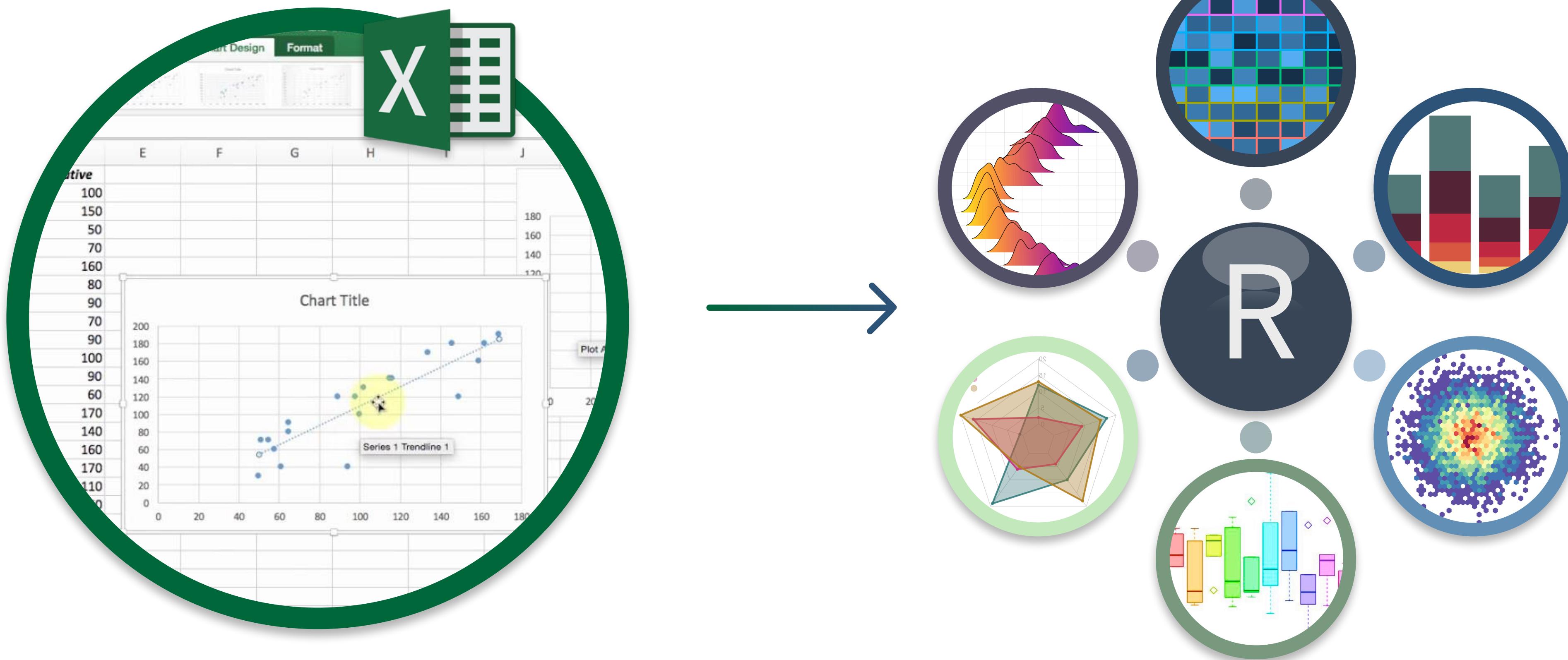
Download and install the newest version of R (<https://cran.r-project.org/>)

Download and install the newest version of R-studio (<http://www.rstudio.com/download>)

Download the course material and place it somewhere you can find it again!
<https://github.com/Center-for-Health-Data-Science/FromExceltoR>

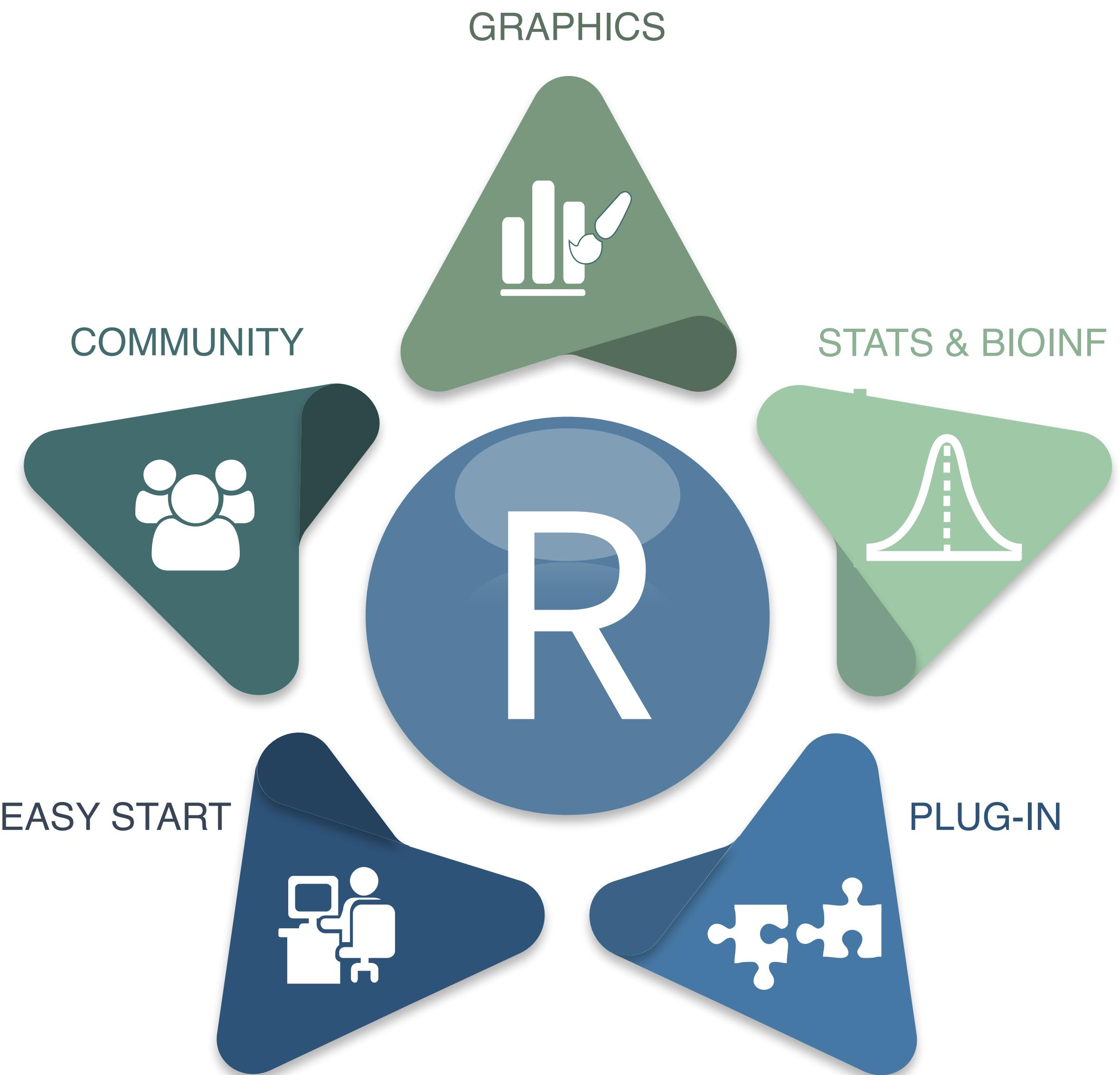


WELCOME TO FROM EXCEL TO R

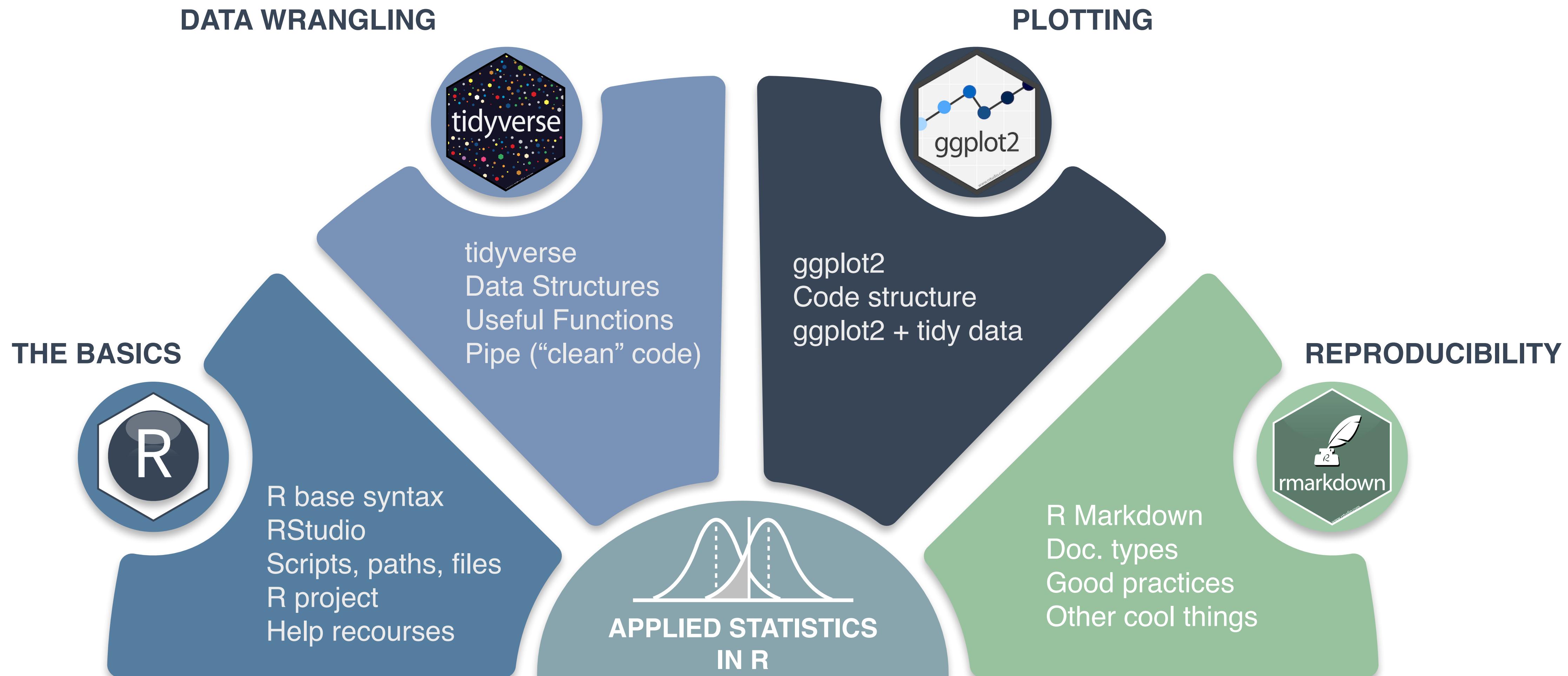


WHY R ?

- **Easy to get started with:**
Compatible with all systems, great support
- **Open Source - Community:**
R-packages, pipelines, tutorials, help pages
- **Flexible Language:**
Plugins, git/github, R Shiny, Rmarkdown, ...
- **Easy & Customisable Graphics**
- **Made for Statistical & Bioinformatics Analysis**



WHAT WILL YOU LEARN IN THIS COURSE?



PROGRAM

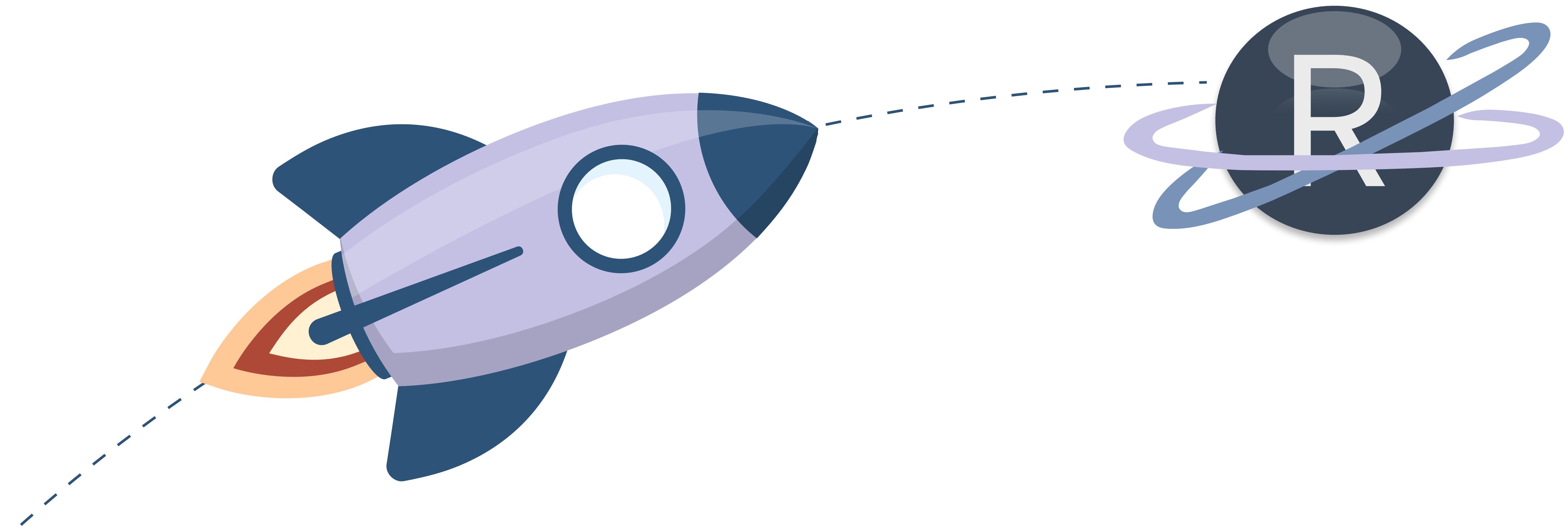
DAY 1

08:30 - Installation Issues & Coffee
09:00 - Introduction to R Basics
10:00 - Rstudio Exercise
10:45 - Break
11:00 - Tidyverse
12:00 - Lunch
13:00 - Tidyverse Exercise
14:30 - Break
14:45 - Rmarkdown
15:15 - Rmarkdown Exercise
16:00 - Q&A - See you tomorrow

DAY 2

08:30 - Coffee
09:00 - ggplot2
10:00 - ggplot2 Exercise
10:45 - Break
11:00 - Applied Statistics
12:00 - Lunch
13:00 - Applied Statistics
14:30 - Break
14:45 - Basic Data Analysis Exercise
15:45 - Cool things in R & Course Evaluation

— FROM EXCEL TO R
LET'S GET STARTED



R & FRIENDS



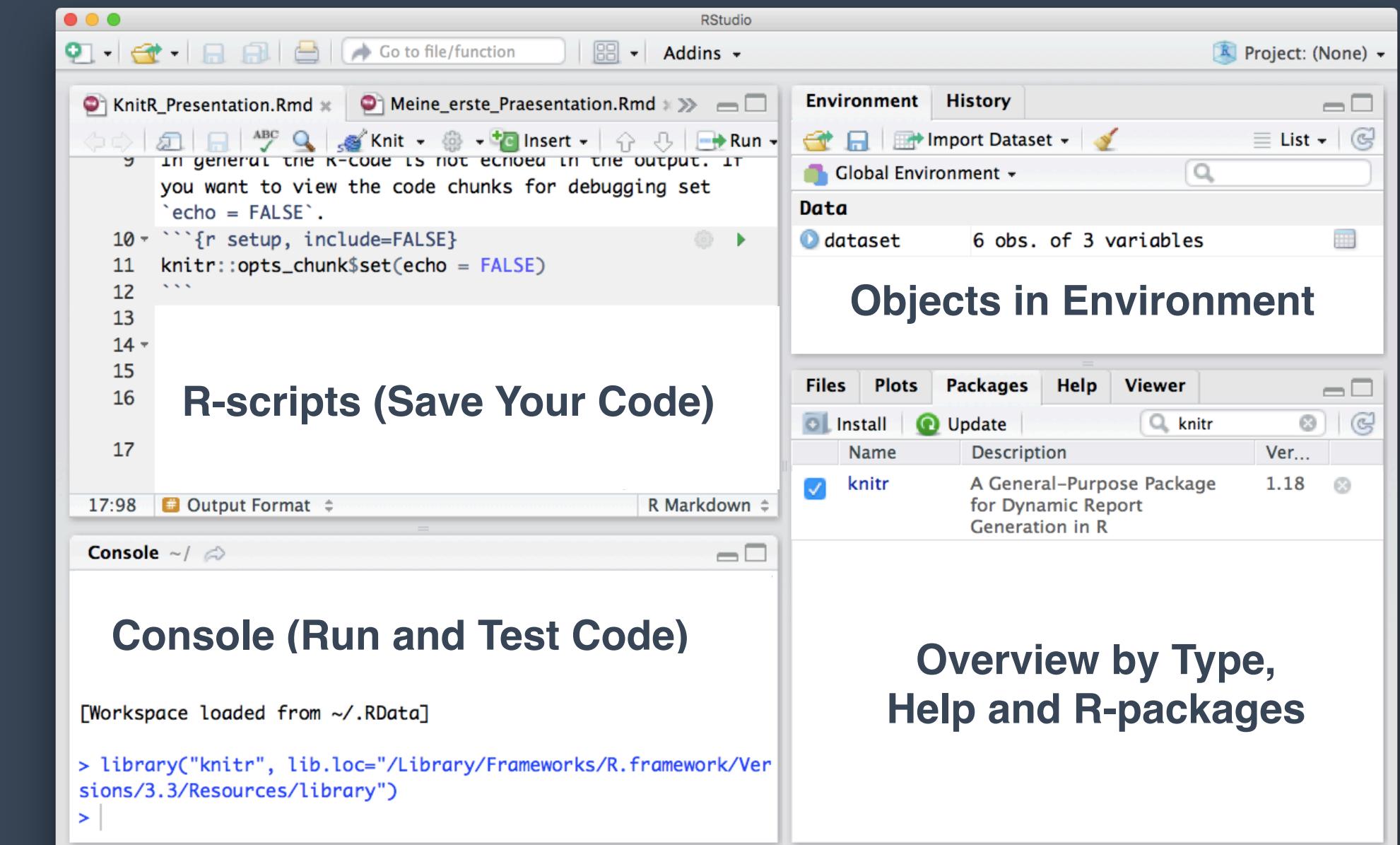
Scripting / Programming Language



Reports (html, pdf, latex)

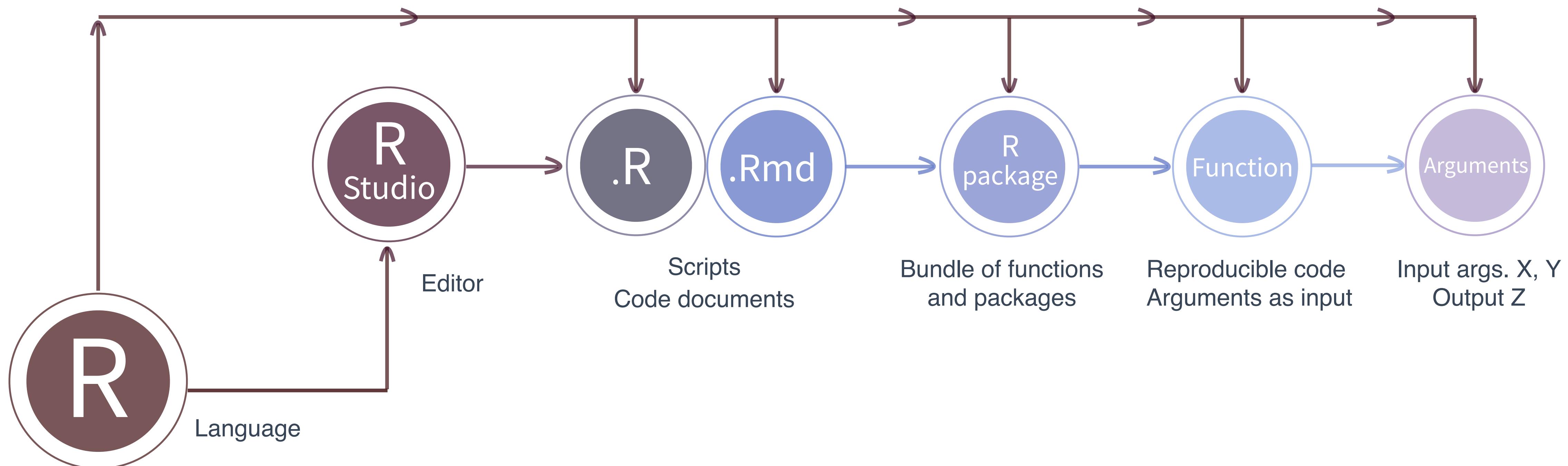


R Studio



R Code Interpreter and Editor

THE ANATOMY OF R



FIRST TIME IN R?

PACKAGES & FUNCTIONS

?*my.package*, ?*my.function*

What is it? Input?

install.packages(), *remove.packages()*

TIPS

Arrows↑↓ to find the code you ran

R studio tips: view, diagnostics

✖️⚠️ Auto-complete with tab

R-cheat sheets (<https://rstudio.com/resources/cheatsheets/>)



WORKING DIRECTORY

setwd(), *getwd()*, *list.files()*, *list.dirs()*

Where am I working from? Full/relative path.

SAVE YOUR WORK

.R, (or .Rmd)

The file with my code. Save it!

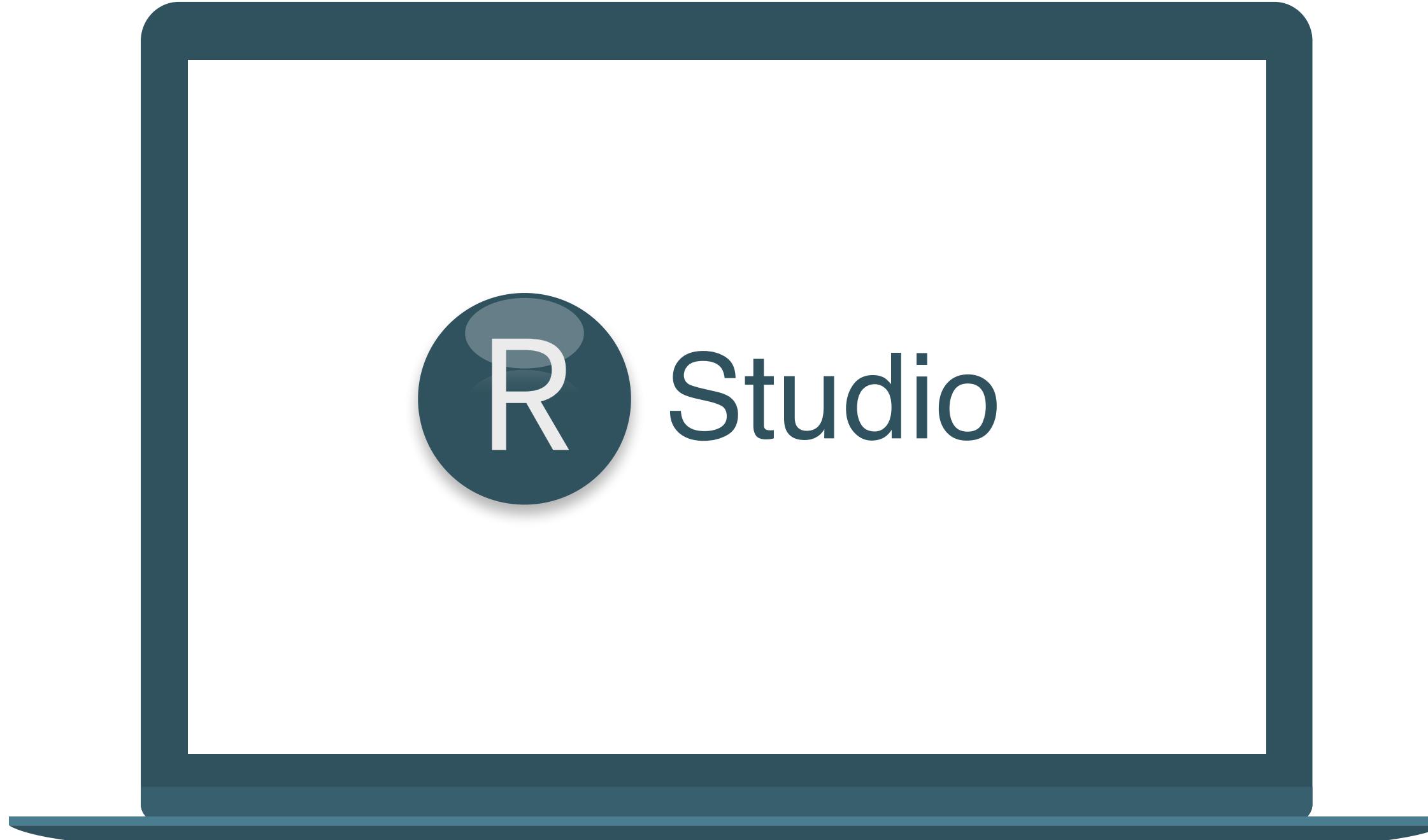
R project

Save Session, everything together

RUN CODE

Run button, highlight enter, short-cut

R STUDIO BASICS



1. R Project

File —> New Project —> New/Existing —> Create Project

2. Set Path

getwd() - Get directory
setwd() - Set directory

setwd("/Users/Tom/Rstuff") - Full path
setwd("./Rstuff") - Relative path

Session —> Set Working Directory —> Choose Directory

3. R Script

Script Icon —> R Script —> File —> Save as ...

4. Install and load a R package.

install.packages("my.package")
library (my.package) - Load package.

Tools —> Install packages —> my.package

ONLINE RESOURCES FOR R

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



GET STARTED

<https://rseek.org/>

<https://rstudio.com/resources/cheatsheets/>

<http://www.cookbook-r.com/>

<https://www.statmethods.net/r-tutorial/index.html>



GRAPHICS

<https://www.r-graph-gallery.com/>

<http://r-statistics.co/Top50-Ggplot2-Visualizations-MasterList-R-Code.html>



BOOKS & COURSES

<https://www.r-bloggers.com/best-books-to-learn-r-programming/>

<https://www.datacamp.com/>

<https://www.codecademy.com/>

<https://www.coursera.org/>



OTHER RESOURCES

<https://github.com/trending/r>

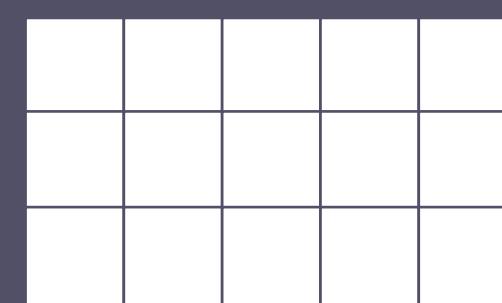
<https://blog.revolutionanalytics.com/>

<https://stackoverflow.com/questions/tagged/r>



R DATA TYPES & STRUCTURES

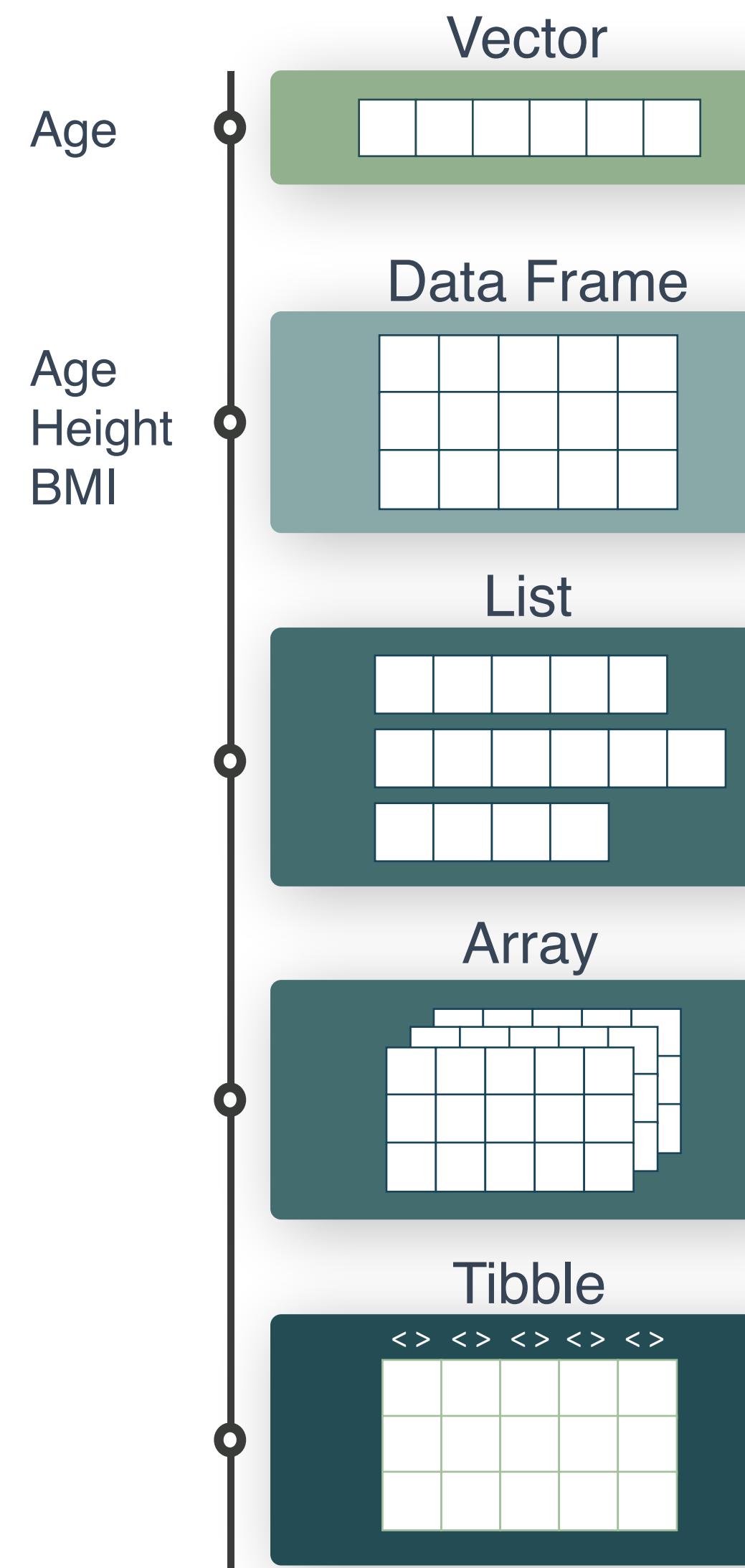
VARIABLES



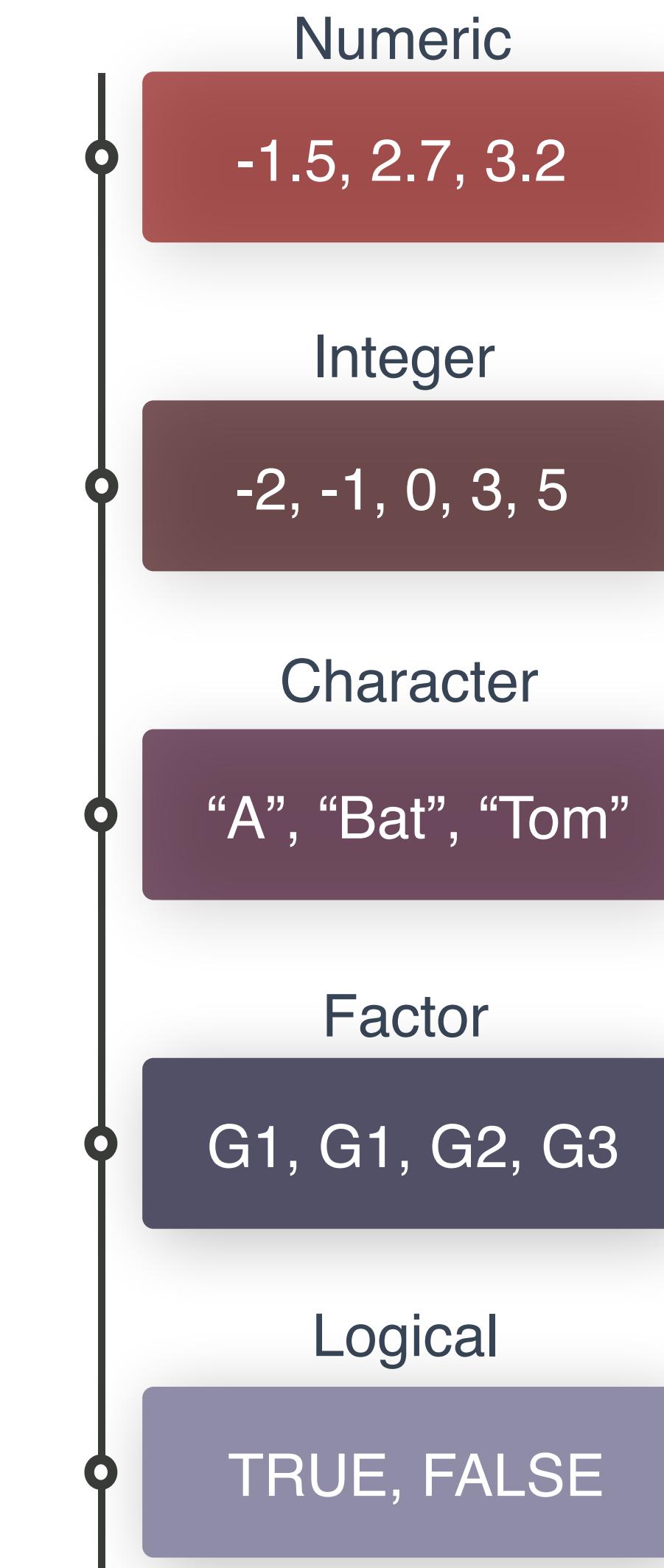
OBSERVATIONS

— FROM EXCEL TO R

DATA STRUCTURES



DATA TYPES



R BASE SYNTAX - RUN THROUGH

VARIABLE ASSIGNMENT

```
> a <- 'apple'  
> a  
[1] 'apple'
```

```
> x <- c(1.5, 2.6, 1.7, 3.2, 3.0, 2.9, ...)  
> x  
[1] 1.5 2.6 1.7 3.2 3.0 2.9 ...
```

READING AND WRITING DATA

Input	Output	Description
<pre>df <- read.table('file.txt')</pre>	<pre>write.table(df, 'file.txt')</pre>	Read and write a delimited text file.
<pre>df <- read.csv('file.csv')</pre>	<pre>write.csv(df, 'file.csv')</pre>	Read and write a comma separated value file. This is a special case of read.table/write.table.
<pre>load('file.RData')</pre>	<pre>save(df, file = 'file.Rdata')</pre>	Read and write an R data file, a file type special for R.

DON'T USE

Spaces in names

Special characters
% ? / | \ & \$ @

Unspecific names

Short/long names

R BASE SYNTAX - RUN THROUGH

SELECTING ELEMENTS

x[4] The fourth element.

x[-4] All but the fourth.

x[2:4] Elements two to four.

x[!(2:4)] All elements except two to four.

x[c(1, 5)] Elements one and five.

By Value

x[x == 10] Elements which are equal to 10.

x[x < 0] All elements less than zero.

x[x %in% c(1, 2, 5)] Elements in the set 1, 2, 5.

R-BASE FUNCTIONS

log(x) Natural log.

exp(x) Exponential.

max(x) Largest element.

min(x) Smallest element.

round(x, n) Round to n decimal places.

sig.fig(x, n) Round to n significant figures.

cor(x, y) Correlation.

sum(x) Sum.

mean(x) Mean.

median(x) Median.

quantile(x) Percentage quantiles.

rank(x) Rank of elements.

var(x) The variance.

sd(x) The standard deviation.

CONDITIONS

a == b	Are equal	a > b	Greater than	a >= b	Greater than or equal to	is.na(a)	Is missing
a != b	Not equal	a < b	Less than	a <= b	Less than or equal to	is.null(a)	Is null

BASE R CHEAT SHEET

Basics: `getwd()`, `setwd() # location`
`install.packages('pname')`, `library(pname)`
`ls()`, `rm() # list, remove objects`
`load()`, `data()`, `save() # load, save as .Rdata`

Overview: `head(df, n=10)`, `df[1:10,]` `tail(df, n=10)`
first or last 10 rows
`class()` # data structure
`unique()`, `table() # unique vals, count vals`

Is/As type:
`is.numeric(x)` (character, factor, integer, etc.)
`as.numeric(x)` (factor, matrix, data.frame, etc.)

Other: `seq(1, 10, by = 1.0) # sequence from-to`
`rep(x, times) # replicate n times`
`sort(), reverse() # sort or reverse vector`

Read in data:
`read.xlsx('name.xlsx')`,
`read.delim('name.txt', sep = '\t')`
`read.csv('name.csv', sep=';')`

Make Data: `c() # vector`
`data.frame(x=x, y=y)`
`matrix(x, nrow = 3, ncol = 3)`
`list(x=x, y=y)`

Strings:
`paste(x, y, sep = '')`
`grep('pattern', x) # find str pattern`
`gsub('pattern', 'replace', x) # replace with`

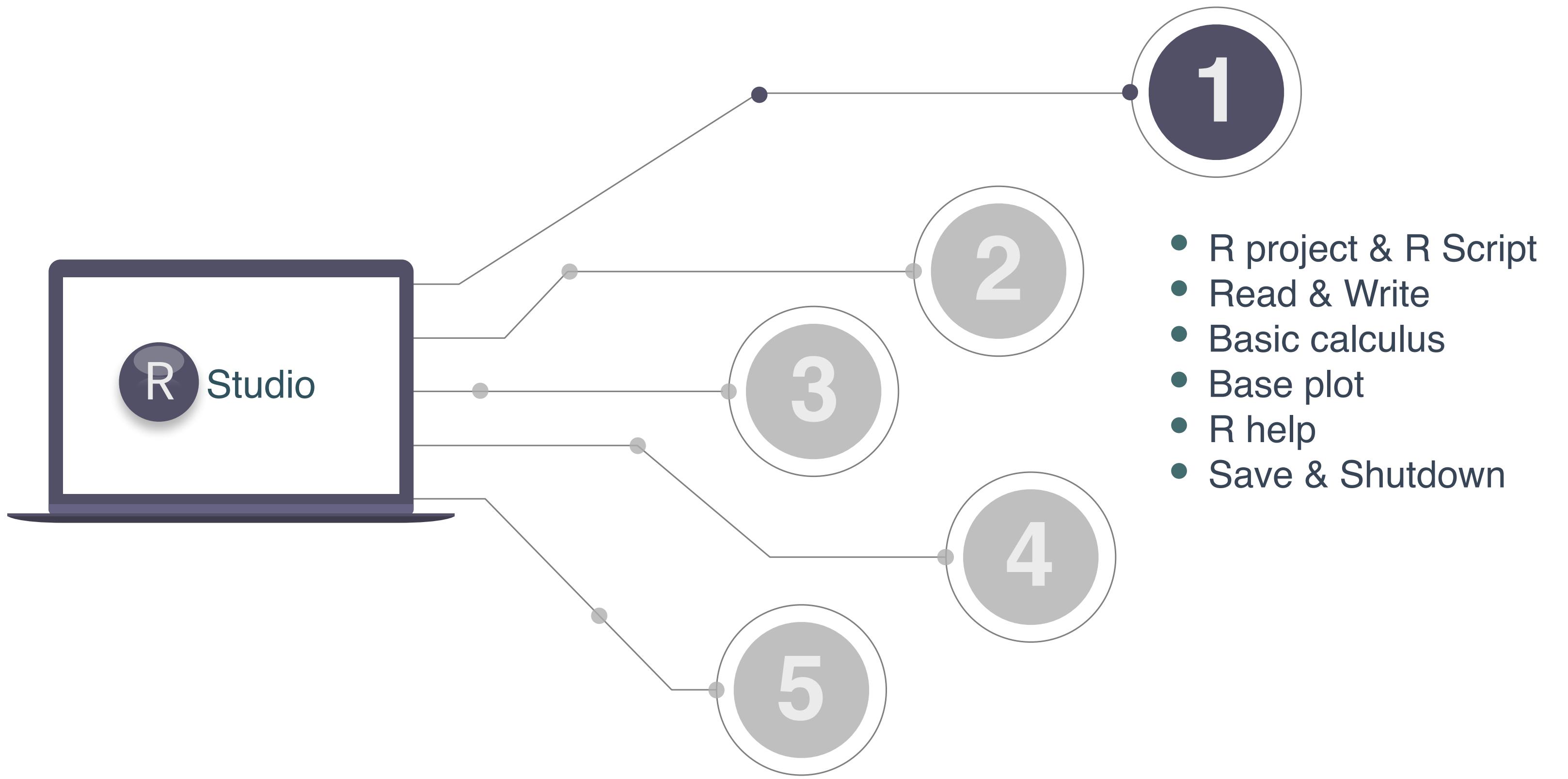
Plots: `plot(x)`
`plot(x,y) # scatter`
`hist(x) # histogram`

GETTING
STARTED

DATA STRUCTURES
& OVERVIEW

DATA TYPES &
STRINGS

VECTORS &
BASE PLOTS



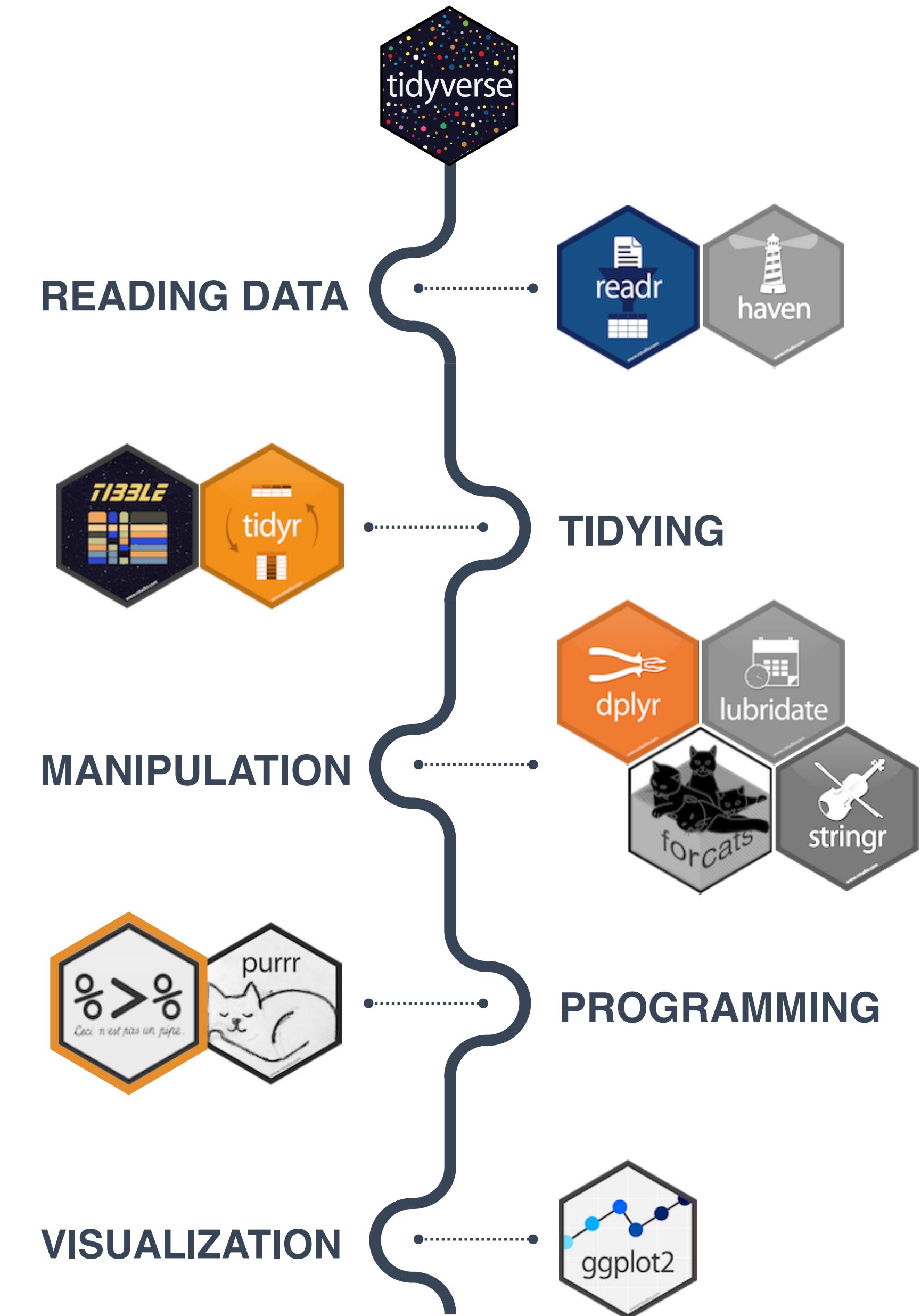
TIDYVERSE

<https://www.tidyverse.org/>

tidyverse is a collection of R packages for data science

“The packages share an underlying design philosophy, grammar, and data structures.” *Wickham and Grolemund*

tidyverse is used to “tidy up” your datasets, so they are easy to work with



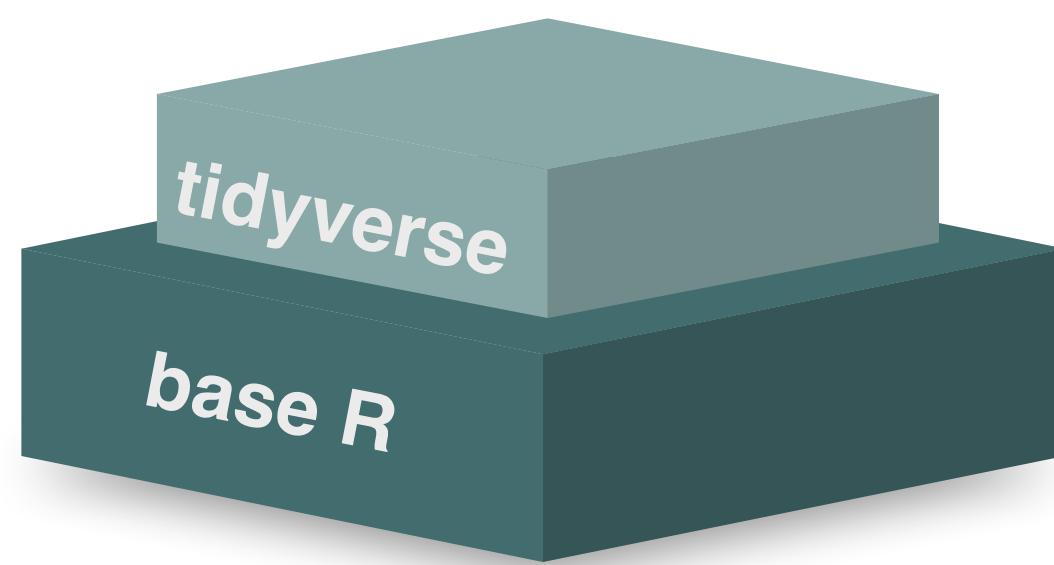
CECI N'EST PAS UNE PIPE

%>%

- You do NOT have to “choose” between tidyverse and base R

BENEFITS

- Short & well-organised code
- Tidy datasets, easy to work with
- Great documentation
- Functions with logical names & inputs



CONSIDERATIONS

- Can be less stable
- “Different syntax”
- Remember what tidyverse is made for!

base R

```
# think from the inside out  
g(f(x,y),z)
```

tidyverse

```
# no brain acrobatics  
x %>% f(y) %>% g(z)
```



pipe symbol

TIDYVERSE CHEAT SHEET

readr, tidyverse, dplyr, ...

Read Data (*readr*)

Reading tabular data

There are solutions for multiple data types
`read_excel()` # using *readxl* package
`read_table()`
`read_csv()`

Useful arguments

Skip lines: `read_csv(file, skip=1)`
Read subset: `read_csv(file, n_max=1)`

Data types

readr guesses the types of each column and tells you about it
("Parsed with column specifications: ...")

HELP

R Documentation (e.g. enter `?dplyr::filter` and see examples)

Much more info and detailed cheat sheets:

<https://brianward1428.medium.com/introduction-to-tidyverse-7b3dbf2337d5>

It also helps to google "tidyverse + whatever you want to do"

Data Tidying (*tidyr*)

Handle missing values

`drop_na()`
`fill()`
`replace_na()`

Subsetting

`tibble[:,1:5]` # returns a tibble
`tibble$colname` # returns a vector
(same as `tibble[[colname]]`)

Reorganize layout

Change between long and wide format
`gather()` # wide to long
`spread()` # long to wide

Data Manipulation (*dplyr*)

Summary

`summarise()`/`summarize()`
`count()`

Group

`group_by()`

Functions will manipulate each group separately and combine results.

Extract and sort observations # i.e. rows

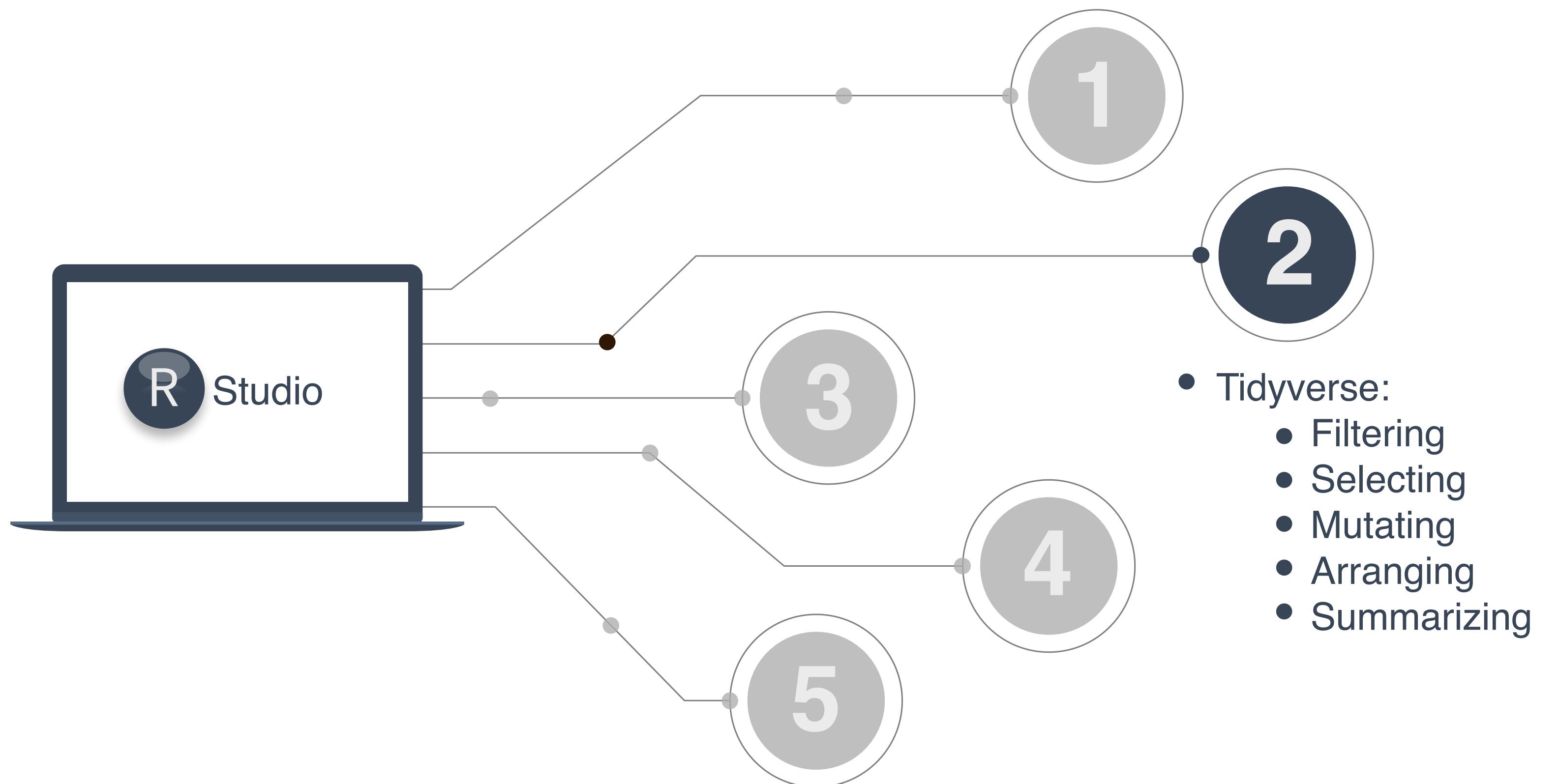
`filter()` # subset by condition
`distinct()` # subset to unique values
`top_n()` # subset by position
`arrange()` # sort low->high, other way with `desc()`

Manipulate variables # i.e. columns

`select()`
`mutate(new_name = f(column))`

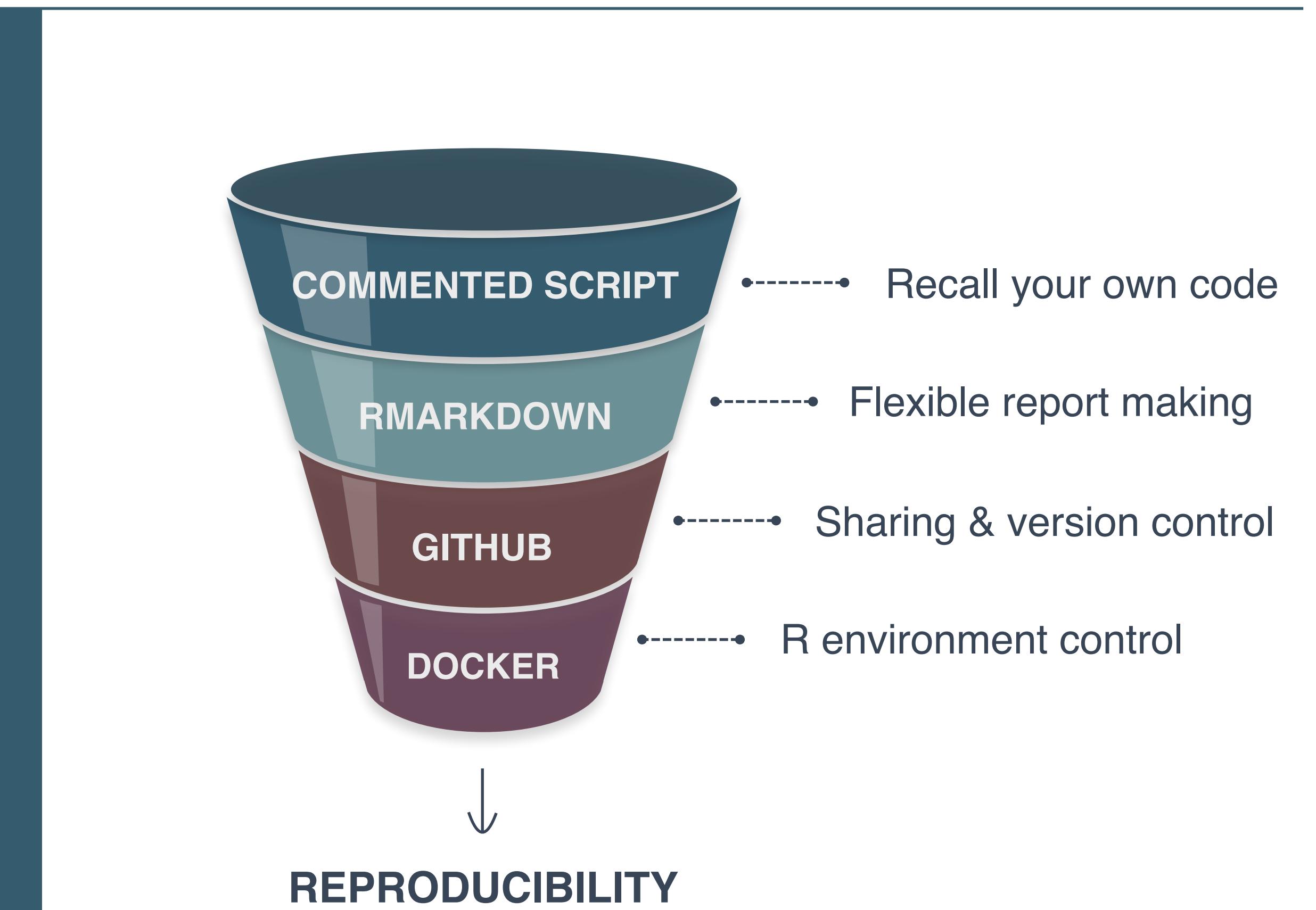
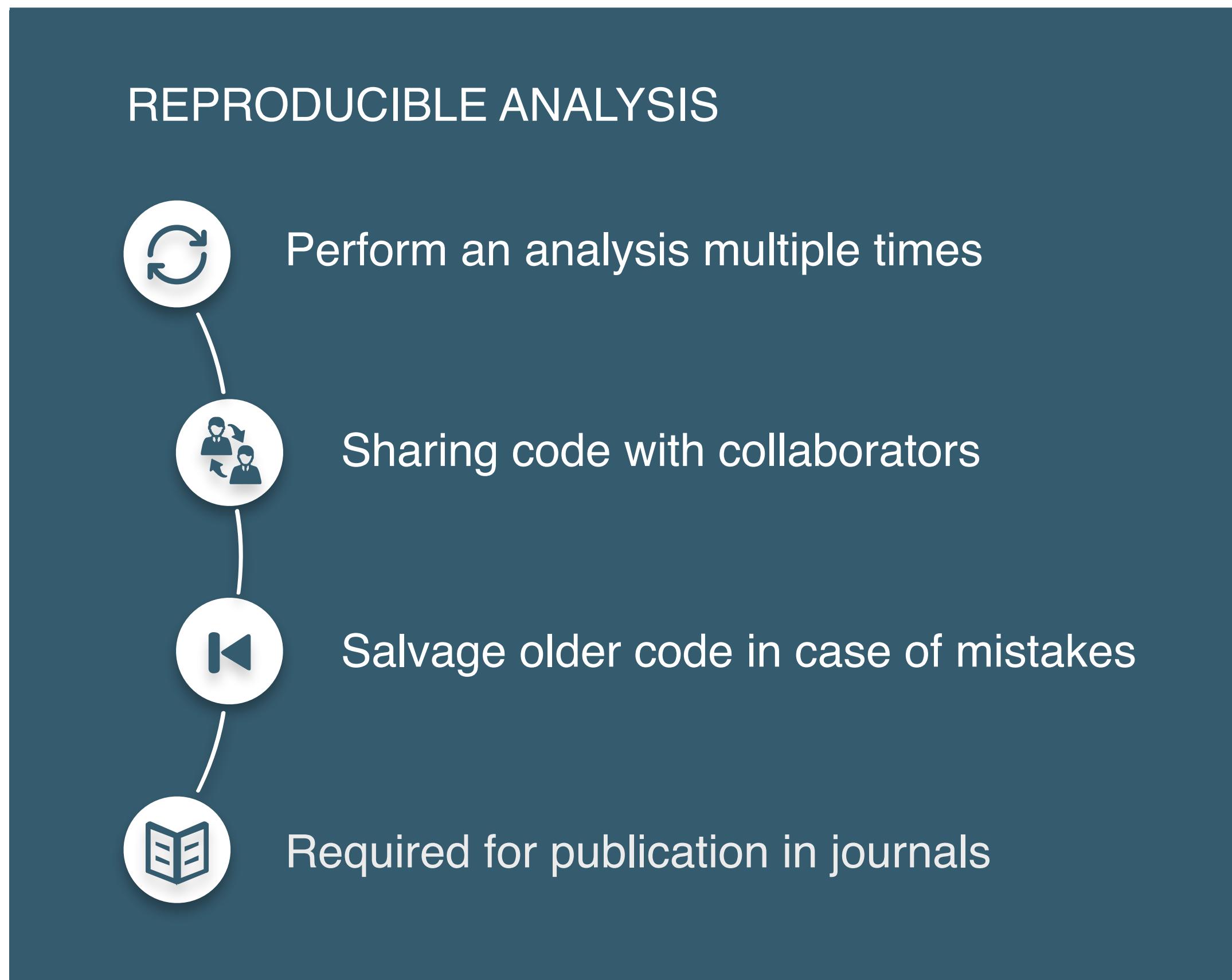
Vectorised functions

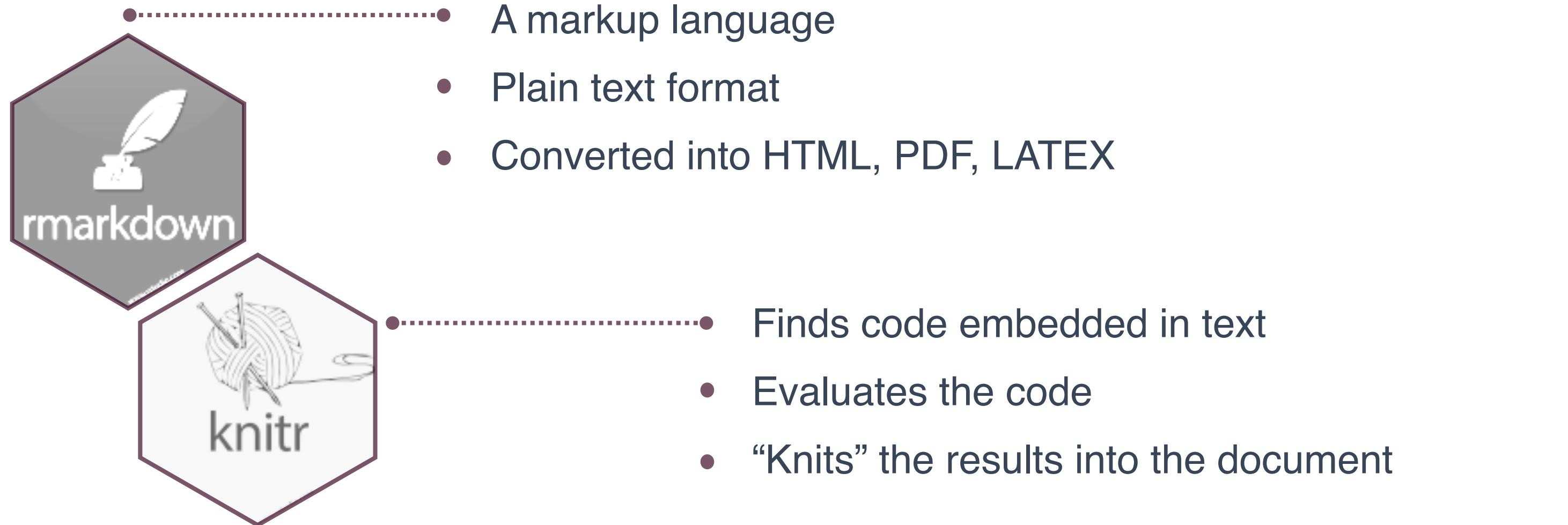
Ranking: `percent_rank()`
Math: Any arithmetic or logical operations, `between()`,
`near()`
`if_else()`



TIDYVERSE EXERCISE 2

REPRODUCIBILITY IN R





— <https://www.markdowntutorial.com/>

R SCRIPT & RMARKDOWN

HOW TO

Write code as normal •

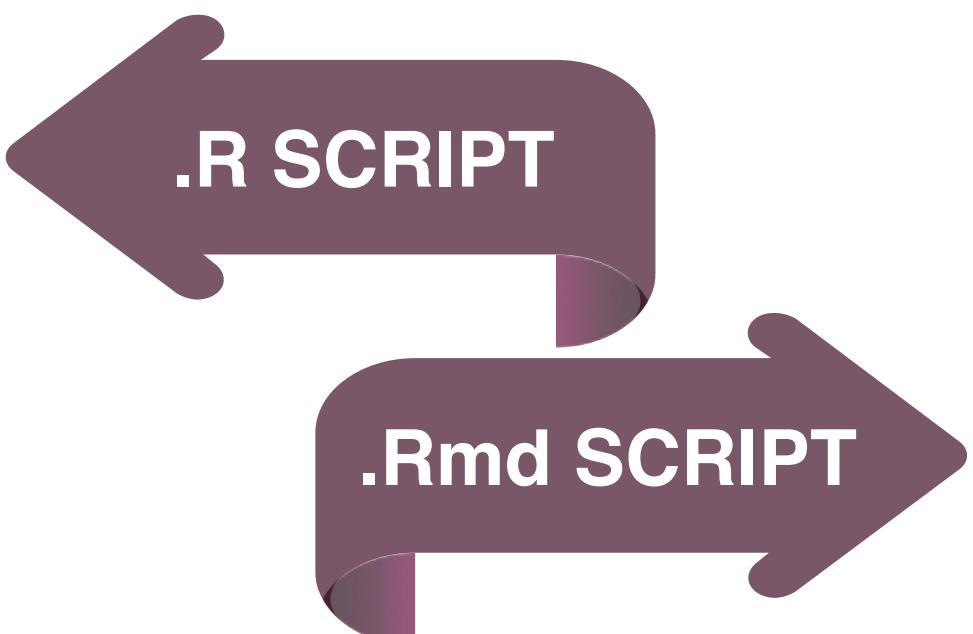
Comment text with # •

USE FOR

Testing new code •

Big data analysis •

Software development •



HOW TO

Write text as normal •

Embed code ``{r} my.code`` •

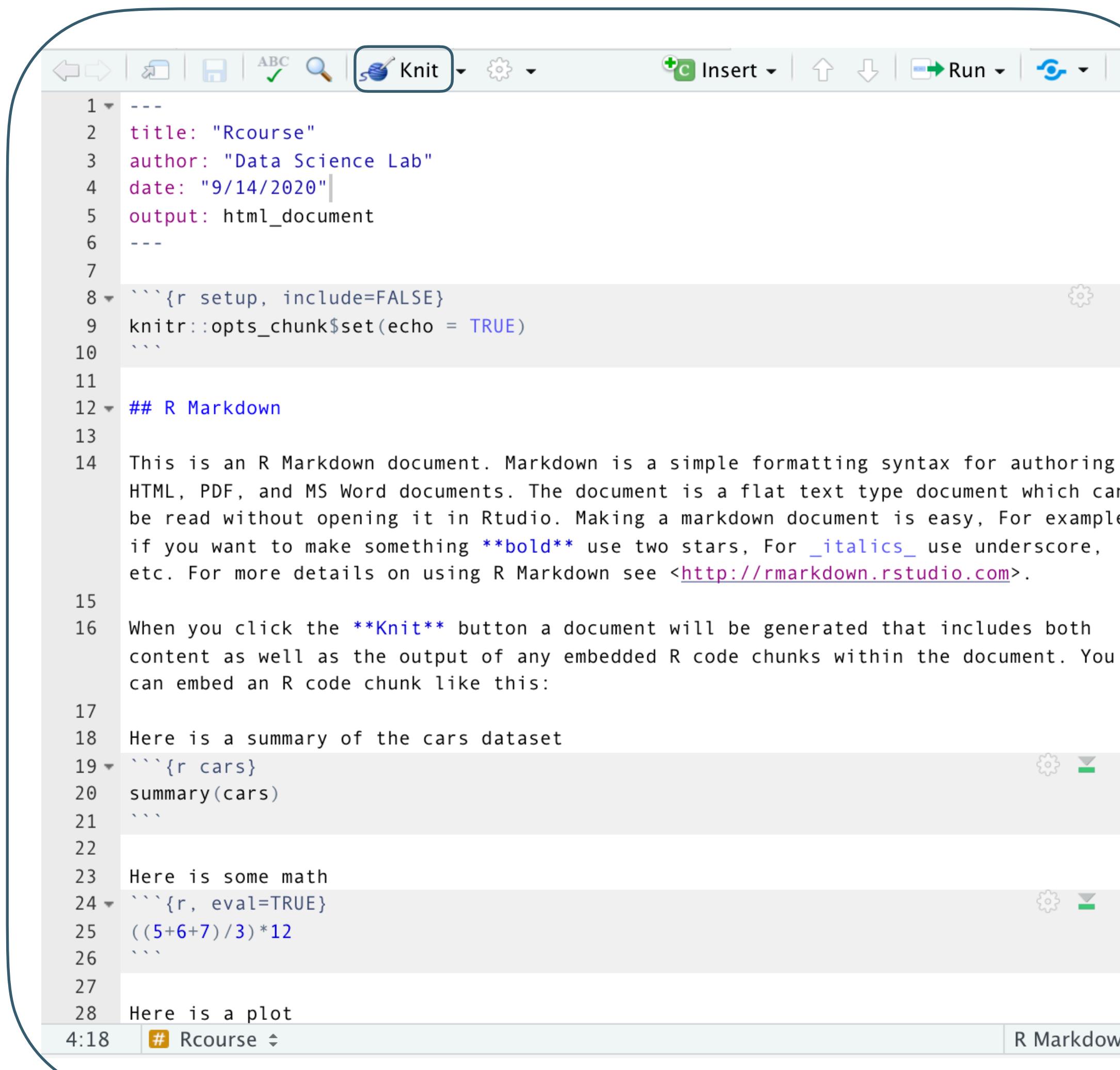
USE FOR

Reports for yourself •

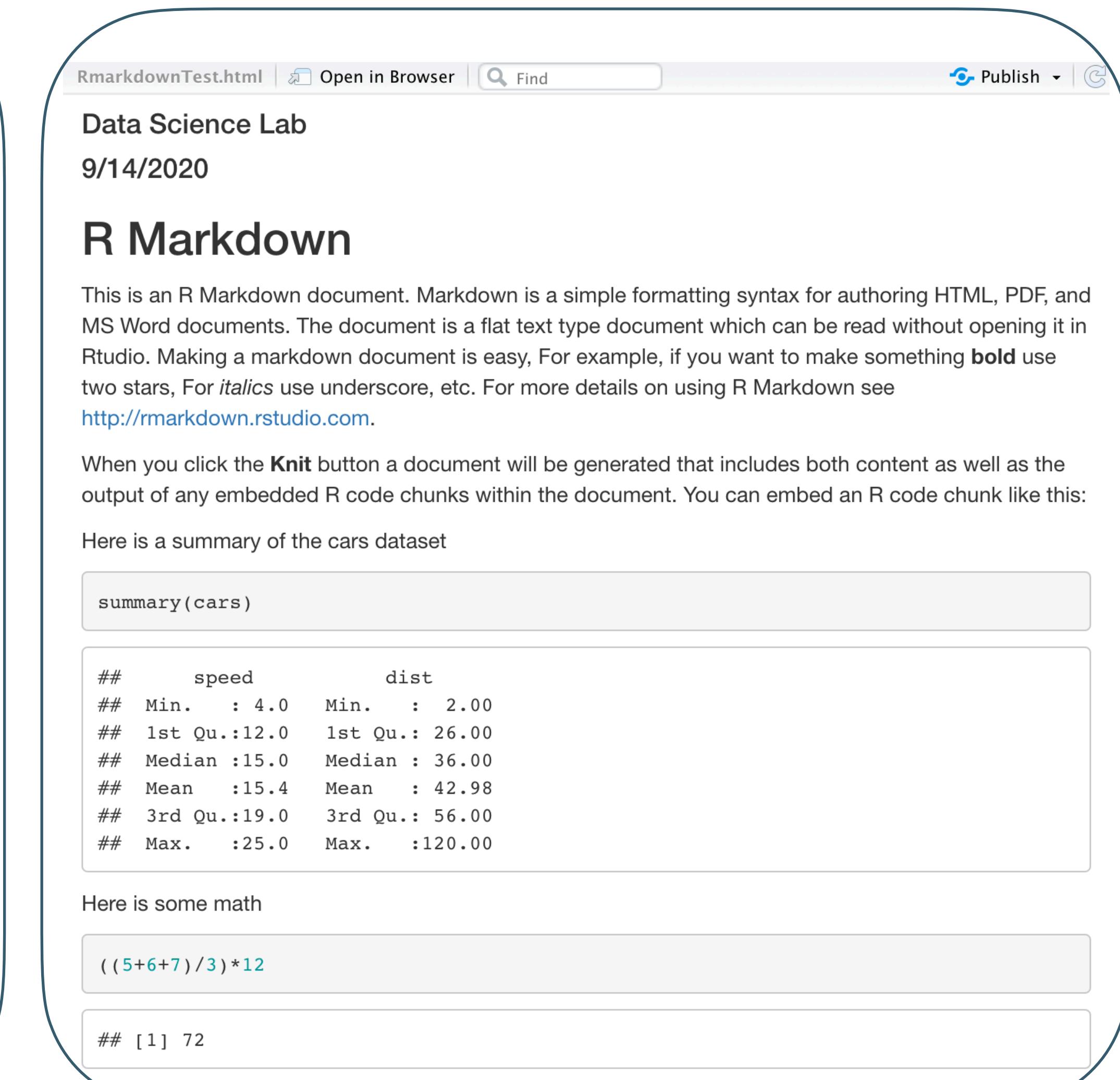
Reports for collaborators •

Tutorials •

RMarkdown



```
1 ---  
2 title: "Rcourse"  
3 author: "Data Science Lab"  
4 date: "9/14/2020"  
5 output: html_document  
---  
6  
7 ```{r setup, include=FALSE}  
8 knitr::opts_chunk$set(echo = TRUE)  
9 ```  
10  
11 ## R Markdown  
12  
13 This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. The document is a flat text type document which can be read without opening it in RStudio. Making a markdown document is easy, For example, if you want to make something bold use two stars, For italics use underscore, etc. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.  
14  
15 When you click the Knit button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:  
16  
17 Here is a summary of the cars dataset  
18 ```{r cars}  
19 summary(cars)  
20 ```  
21  
22  
23 Here is some math  
24 ```{r, eval=TRUE}  
25  $((5+6+7)/3)*12$   
26 ```  
27  
28 Here is a plot  
4:18 # Rcourse
```



Data Science Lab
9/14/2020

R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. The document is a flat text type document which can be read without opening it in RStudio. Making a markdown document is easy, For example, if you want to make something **bold** use two stars, For italics use underscore, etc. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

Here is a summary of the cars dataset

```
summary(cars)
```

	speed	dist
## Min.	4.0	Min. : 2.00
## 1st Qu.	12.0	1st Qu.: 26.00
## Median	15.0	Median : 36.00
## Mean	15.4	Mean : 42.98
## 3rd Qu.	19.0	3rd Qu.: 56.00
## Max.	25.0	Max. : 120.00

Here is some math

```
((5+6+7)/3)*12
```

```
## [1] 72
```

RMARKDOWN CHEAT

Begin .Rmd:

```
---
```

```
title: My Project Name
```

```
output:
```

```
    html_document (pdf_document, ...)
```

```
---
```

Code Chunk:

```
```{r}
```

```
some R code
```

```
```
```

Global Option:

```
```{r setup, include=FALSE}
```

```
knitr::opts_chunk$set(echo = TRUE)
```

```
```
```

GETTING
STARTED

Code Options:

```
echo (= TRUE or FALSE - print my code)
```

```
eval (= TRUE or FALSE - run my code)
```

```
warning (= TRUE or FALSE display warning messages)
```

Figure Options:

```
fig.align (= 'left', 'right', 'center')
```

```
fig.cap (= 'my figure caption')
```

```
fig.height (= n), fig.width (= n)
```

CHUNK
OPTIONS

Header:

Header size ranging from largest (one #)
to smallest (six #):
my.text, ## my.text, ### my.text, etc.

Text:

italics

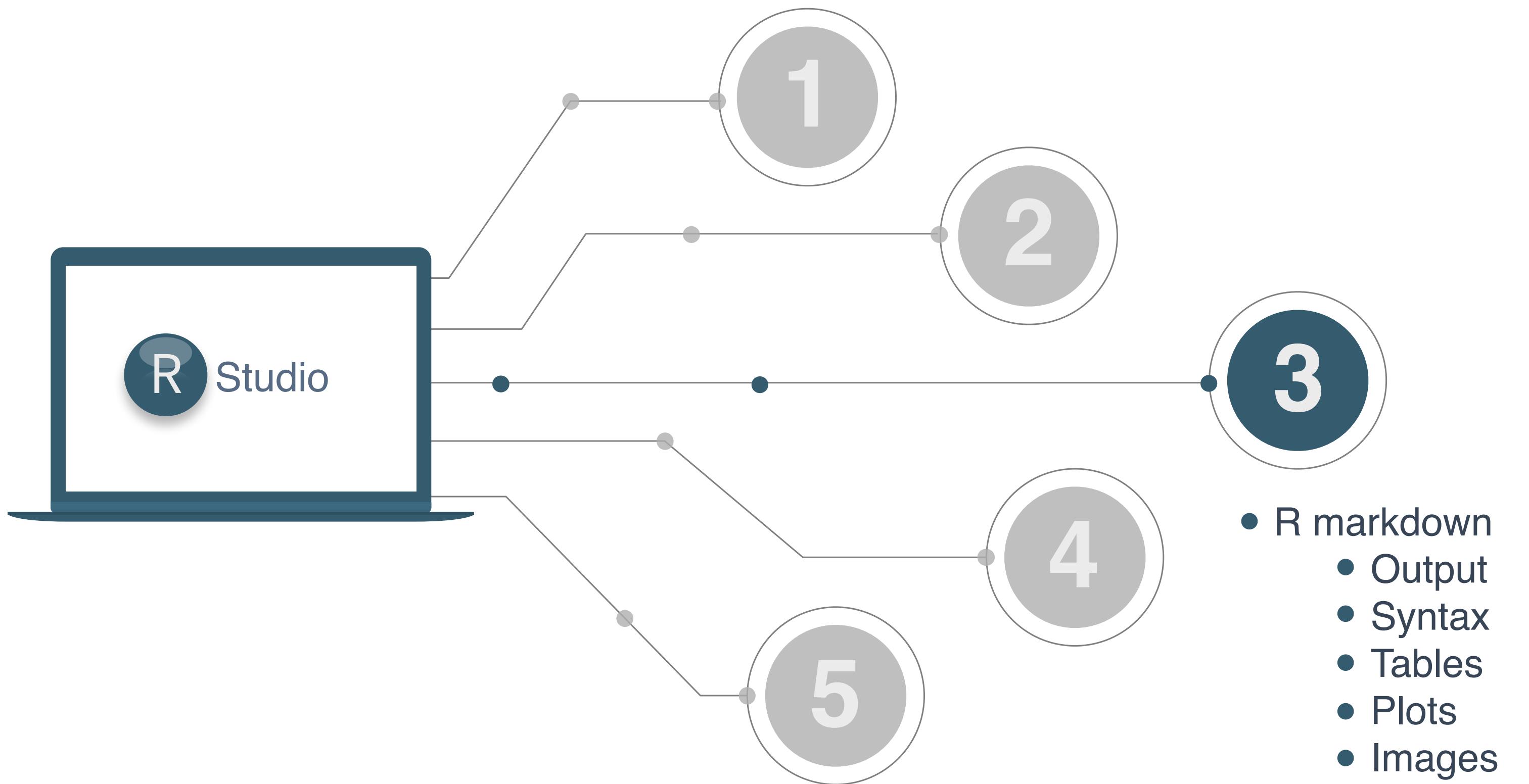
bold

`highlighted`

Lists:

- * List item1 (filled dot)
 - + sub-item1 (open dot)
- 1. List item1 (numbered)
 - i) sub-item1 (roman)

TEXT

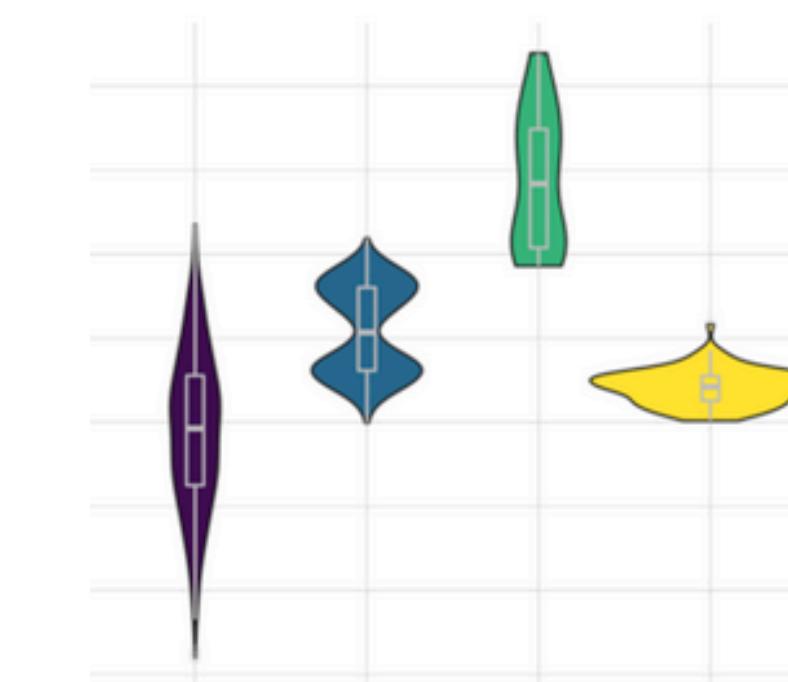


R MARKDOWN EXERCISE 3

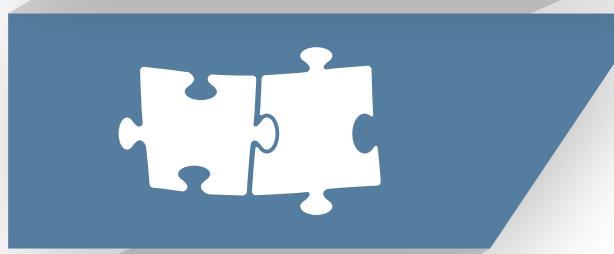
GGPLOT2 - EASY GRAPHICS



Aesthetically pleasing graphics.



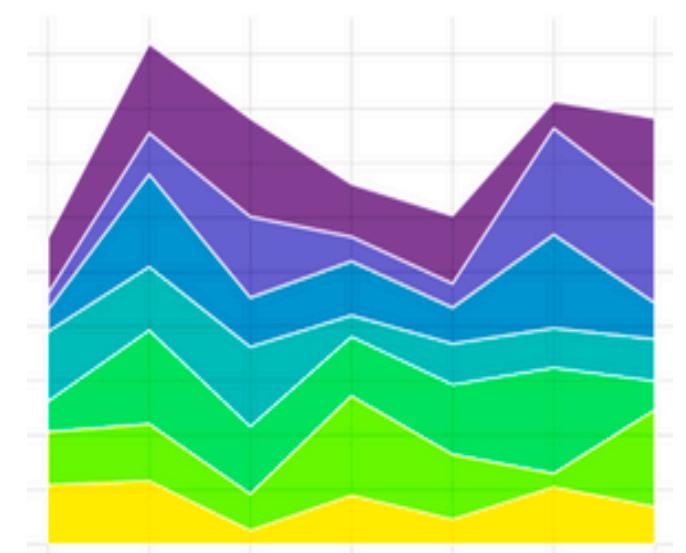
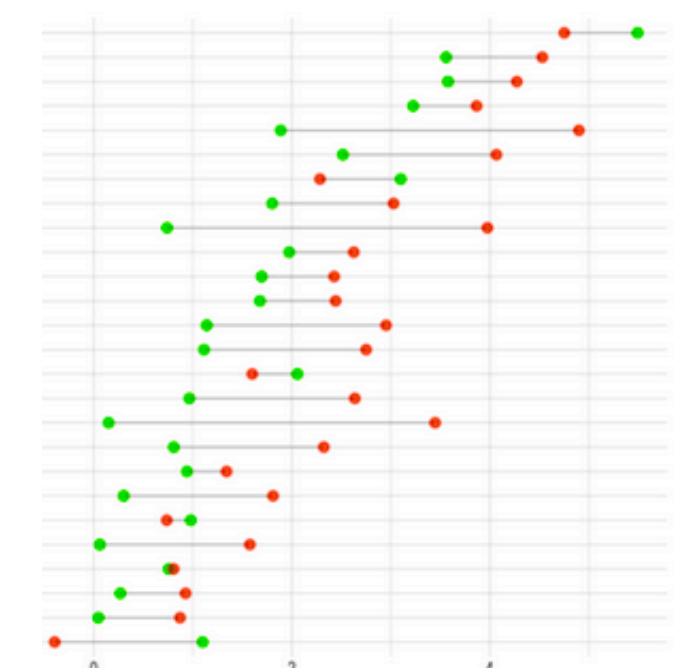
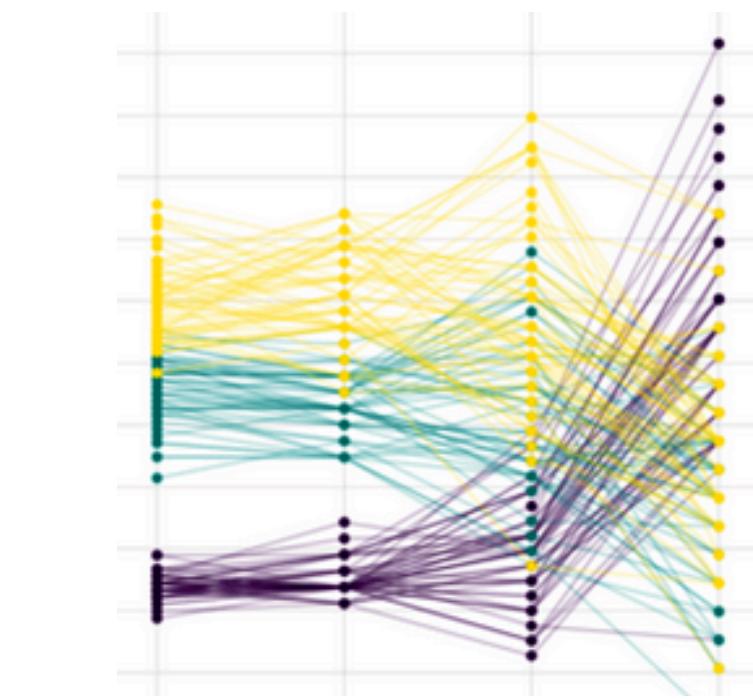
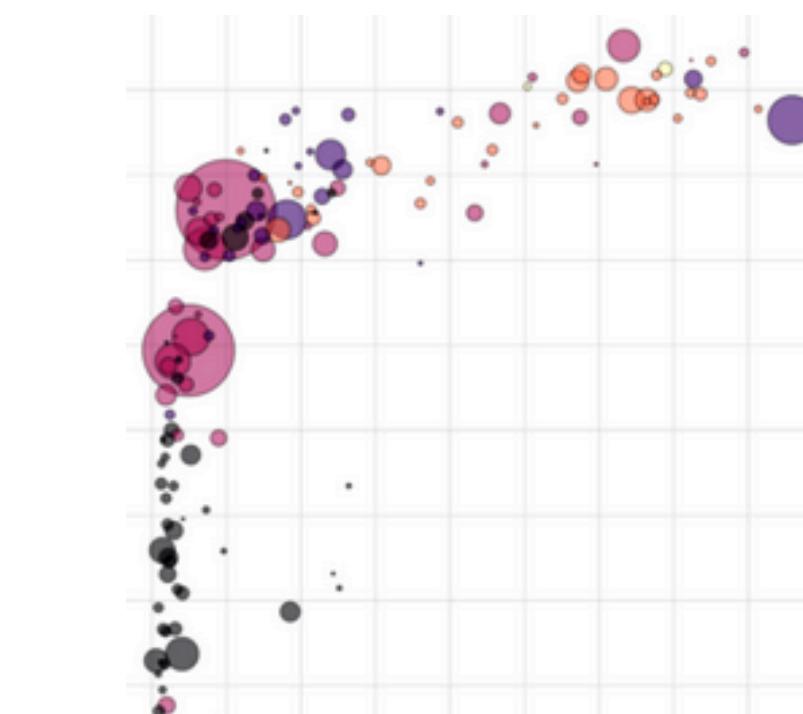
Well-defined “additive” (+) structure.



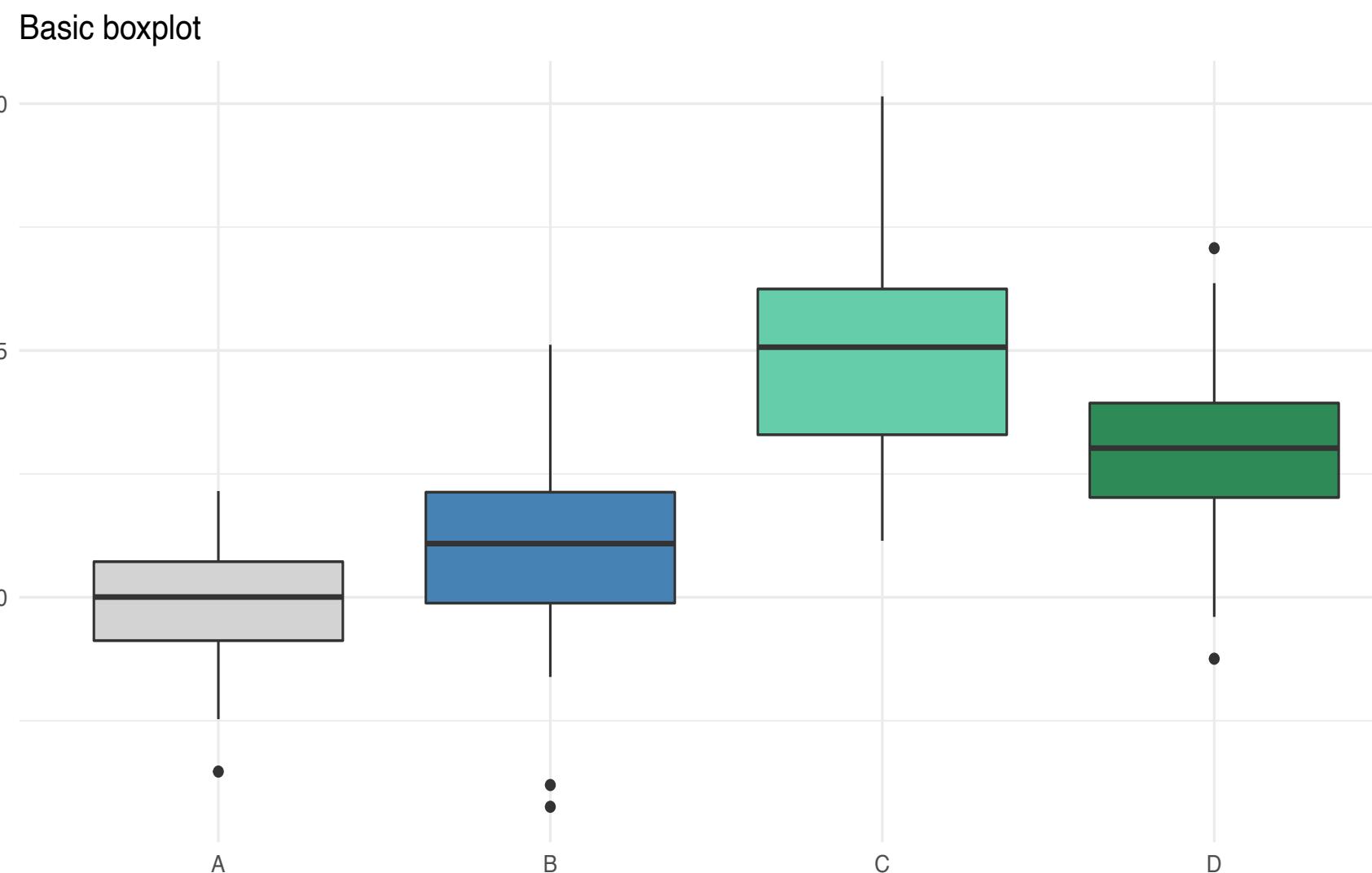
Integrates perfectly with tidy data.



Great documentation & community



GGPLOT2 ADDITIVE STRUCTURE



DATASET, SAMPLES &
OBSERVATIONS

```
ggplot(my.DS, aes(x=Sample,  
y=Measure))
```

DEFINE PLOT TYPE

```
ggplot(my.DS, aes(x=Sample,  
y=Measure))  
+ geom_boxplot()
```

COLOR BY GROUP

```
ggplot(my.DS, aes(x=Sample,  
y=Measure, fill=Sample))  
+ geom_boxplot()
```

TITLE AND LEGEND

```
... + ggtitle("Basic boxplot") +  
theme(legend.position="none",  
plot.title = element_text(size=11))
```

CUSTOM COLORS

```
... + scale_fill_manual(values =  
c("lightgray", "steelblue",  
"aquamarine3", "seagreen4"))
```

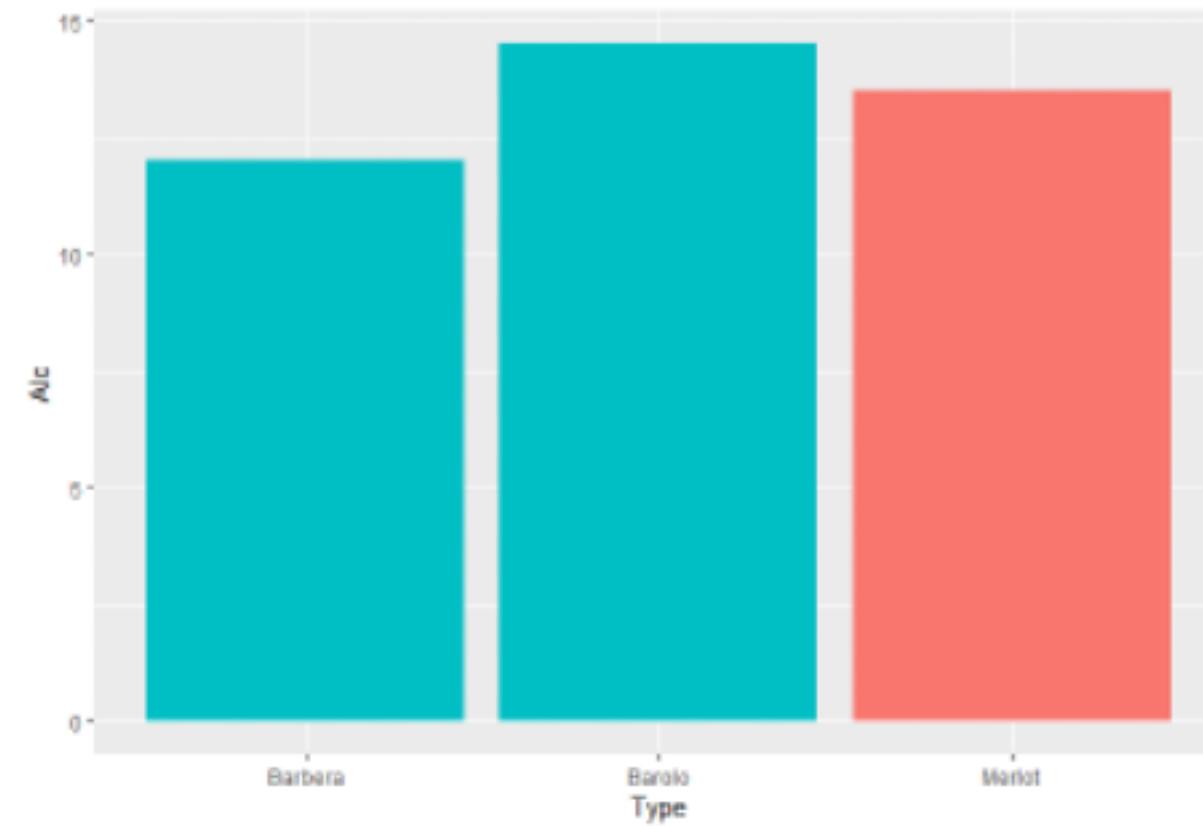
BACKGROUND

```
+ theme_minimal()
```

GGPLOT BASIC STRUCTURE

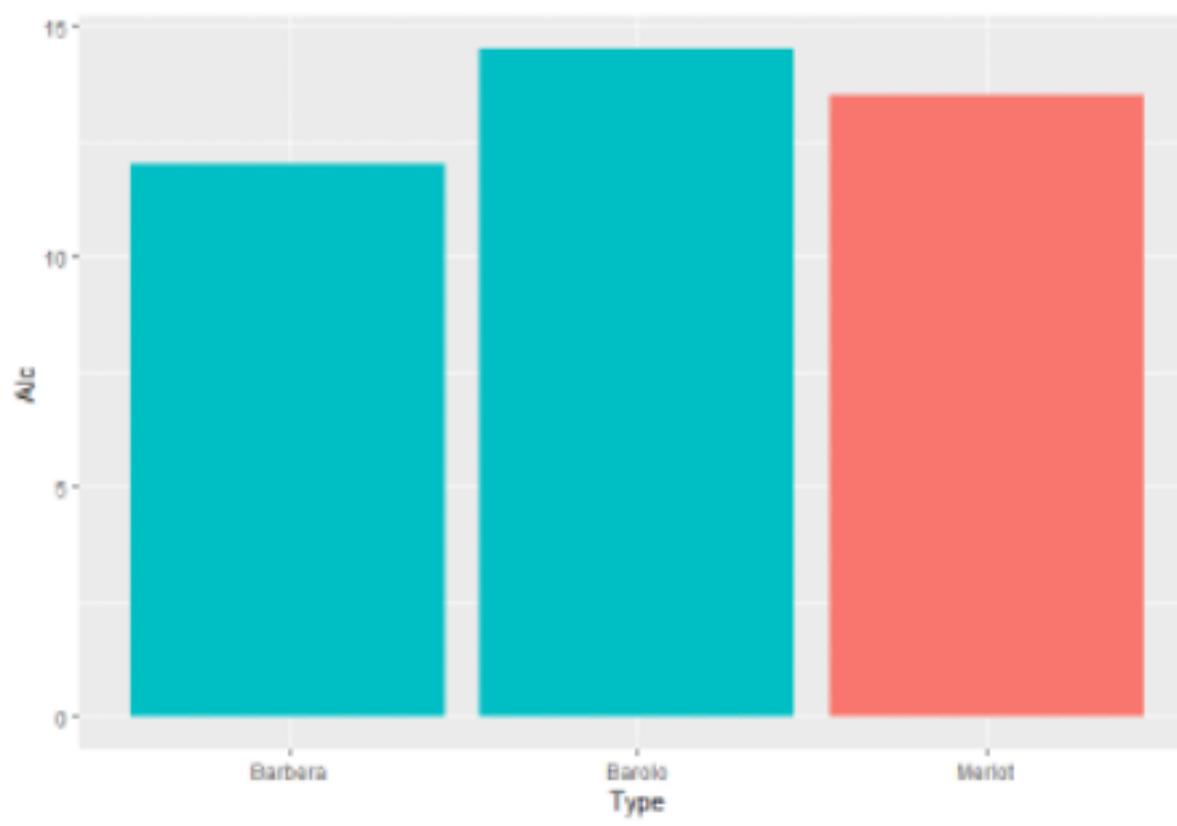
Geoms inherit the parameters from the ggplot they are added to:

```
ggplot(my_wine,
aes(x=Type, y=Alc, fill=Country))
+ geom_col()
```



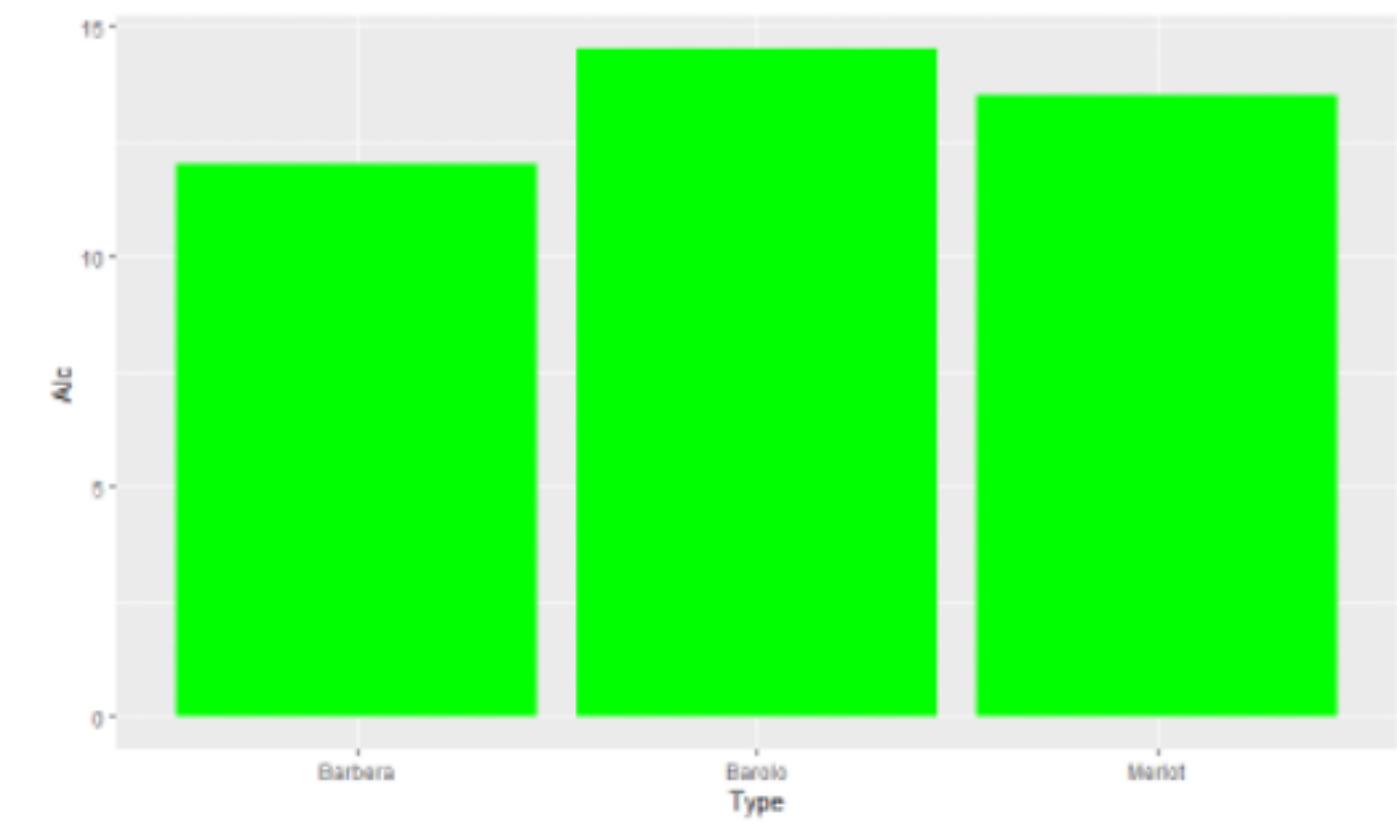
Alternatively, you can specify additional parameters directly in the geom:

```
ggplot(my_wine,
aes(x=Type, y=Alc))
+ geom_col(aes(fill=Country))
```



Things outside the aes are applied to everything!

```
ggplot(my_wine,
aes(x=Type, y=Alc))
+ geom_col(fill='green')
```



GGPLOT CHEAT SHEET

Define Plot:

```
ggplot(data = my.data,  
aes(x = x.var, y = y.var))
```

Add Plot Type:

- + geom_point()
- + geom_line()
- + geom_boxplot()
- + geom_col()
- + geom_density()
- + geom_histogram()

One Color:

```
ggplot(..., aes(...,  
color = "green"))
```

Color Fill by Group:

```
ggplot(..., aes(...,  
fill = group.var))
```

Custom Colors:

- + scale_*_manual(values = c())
- ex: scale_color_manual(values = c("blue", "pink"))

More Colors:

- + scale_fill_grey(start = 0.2, end = 0.8)
- + scale_fill_gradient(low="white", high="red")

Grid Theme:

- + theme_bw()
- + theme_minimal()
- + theme_dark()
- + theme_*

Labels:

- + ggtitle(...)
- + xlab(...)
- + ylab(...)

Text:

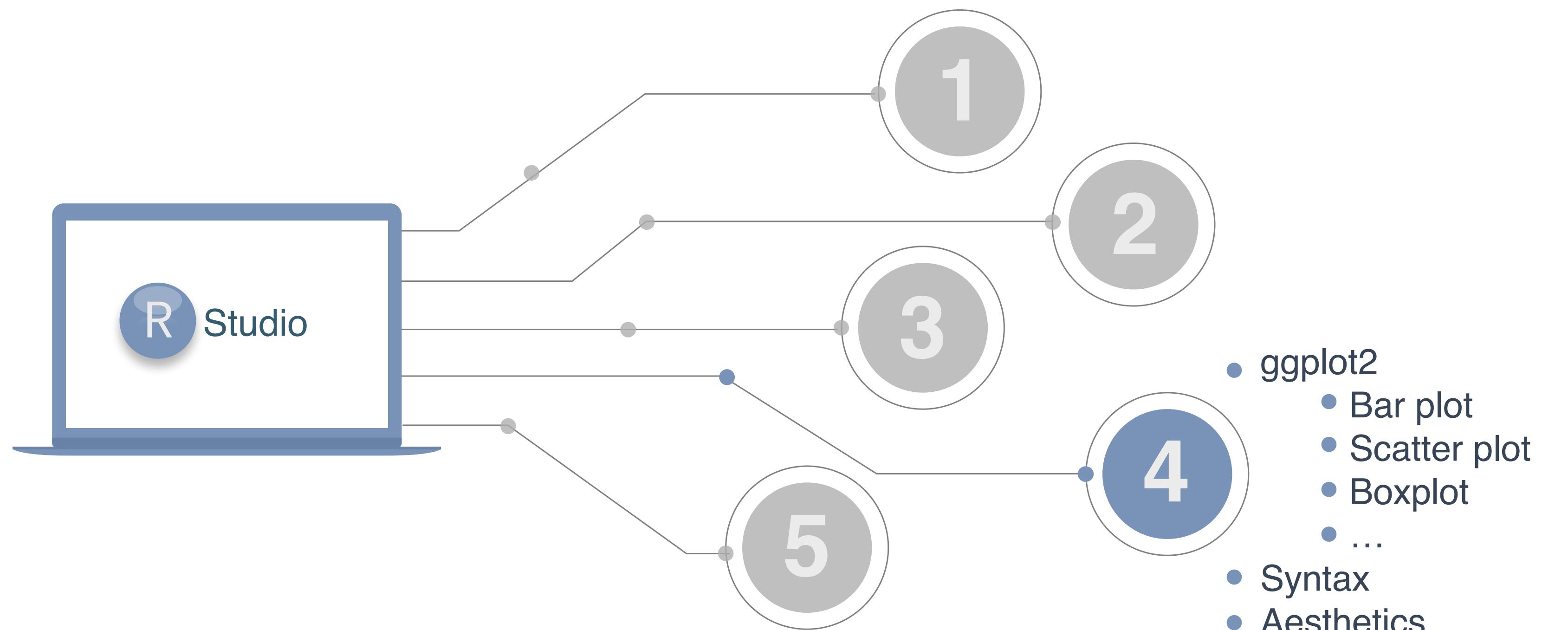
- + theme(* = element_text())
- + theme(axis.title = element_text(angle = 90, colour= "red"),
legend.text = element_text(size = 8, face = "bold"))

GET
STARTED

COLORS

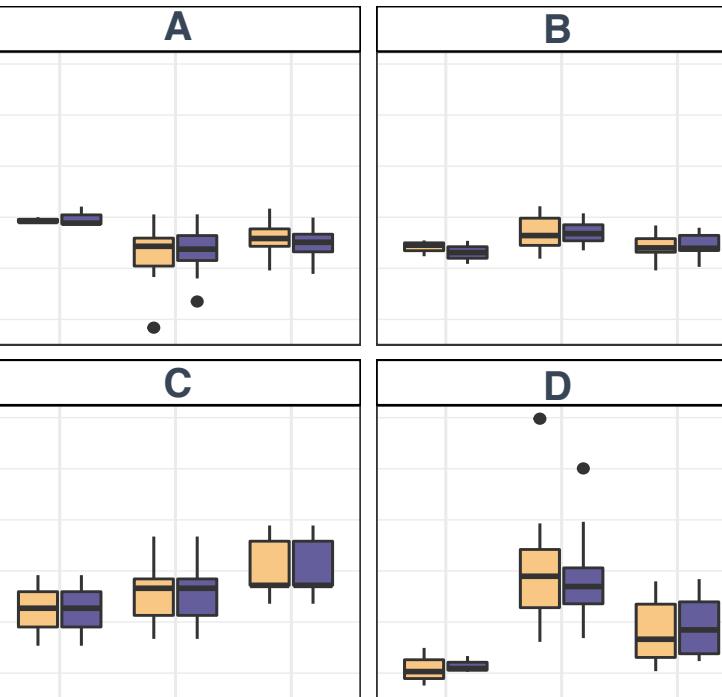
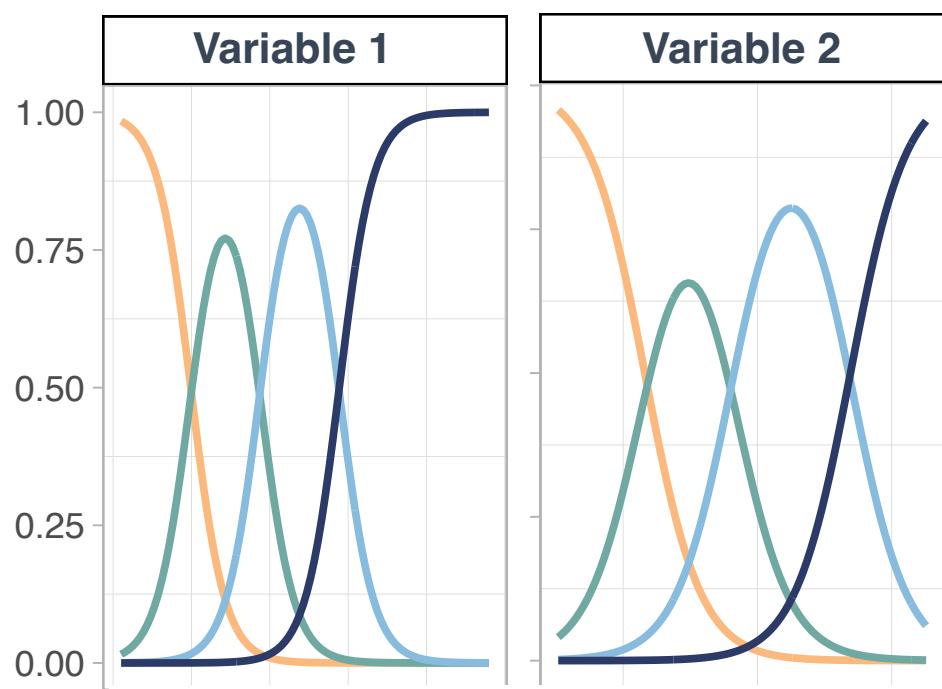
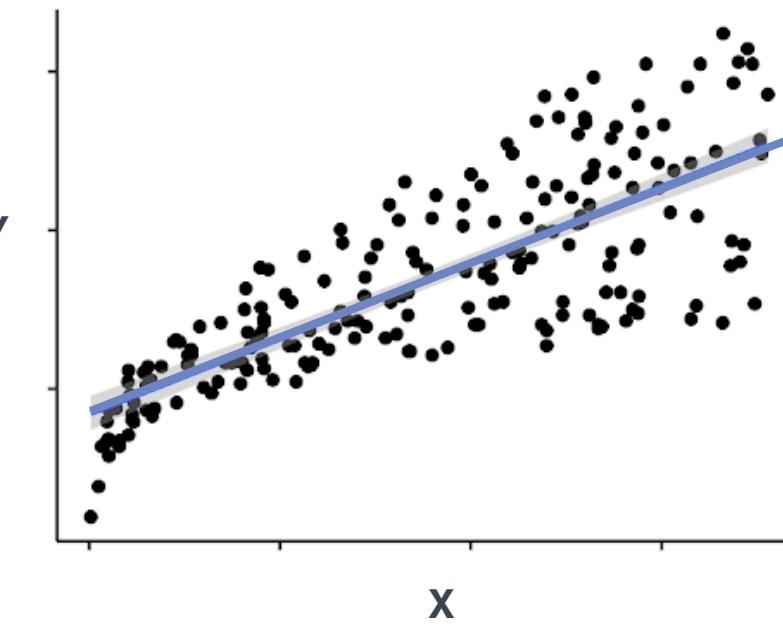
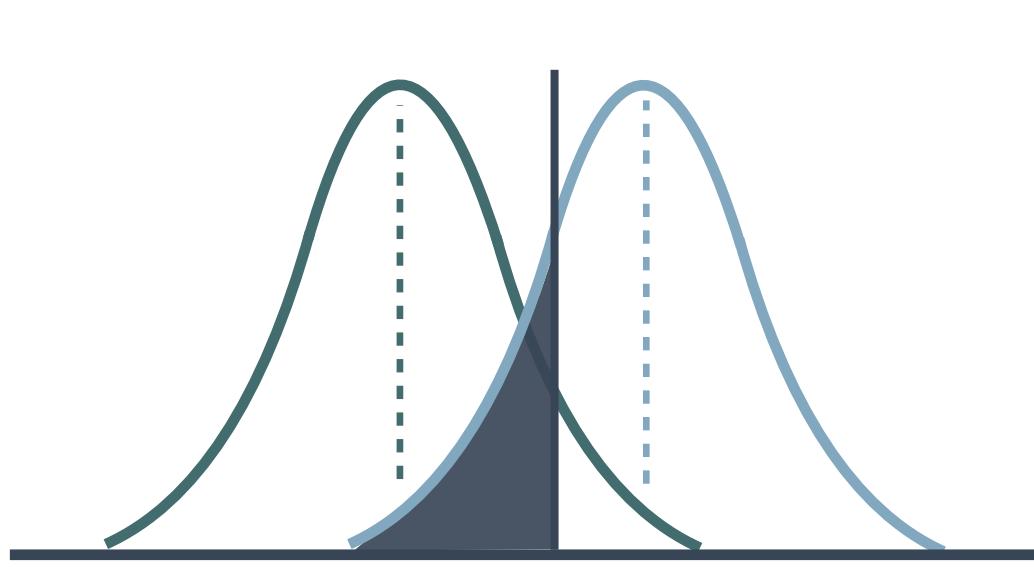
COLOR SCALES
& THEMES

TEXT



— GGPlot2 EXERCISE 4

R - A STATISTICAL SCRIPTING LANGUAGE



MODEL FUNCTIONS

`lm()`, `glm()`
`lmer()`, `glmer()`,
`nls()`, ...

EMMEANS PACKAGE

`emmeans()`,
`pairs()`, `cld()`

APPLY TO MODEL

`summary()`, `anova()`,
`confint()`, `predict()`,
`drop1()`, `update()`,
`step()`, ...

MORE FUNCTIONS

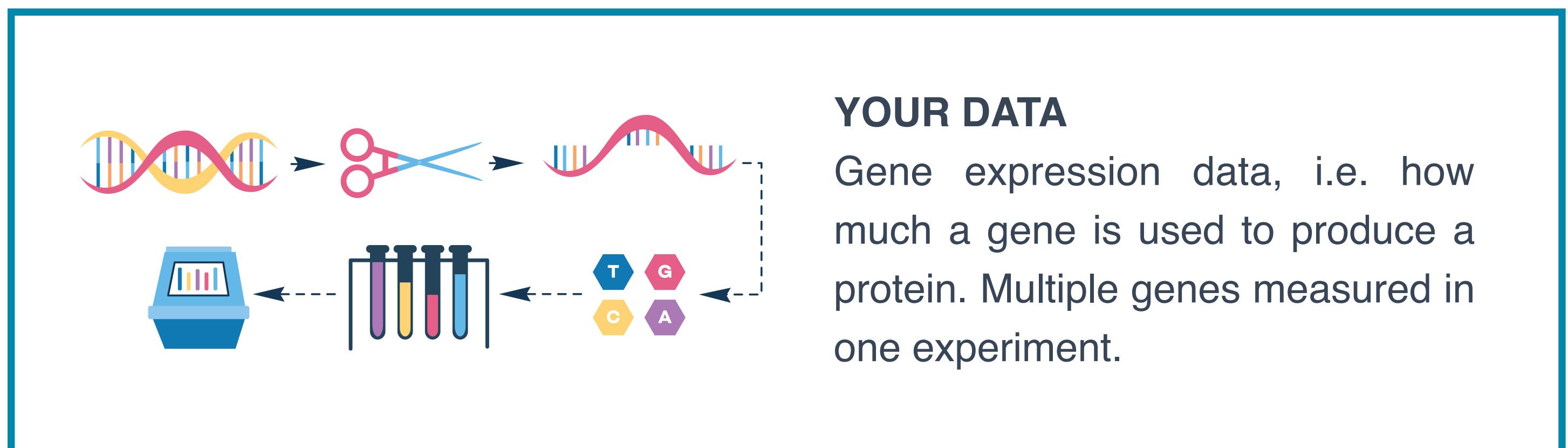
`t.test()`, `cor()`,
`cor.test()`, `aov()`,
`quantile()`,
`p.adjust()`,
`rank()`, ...

Let's use R in a statistical analysis



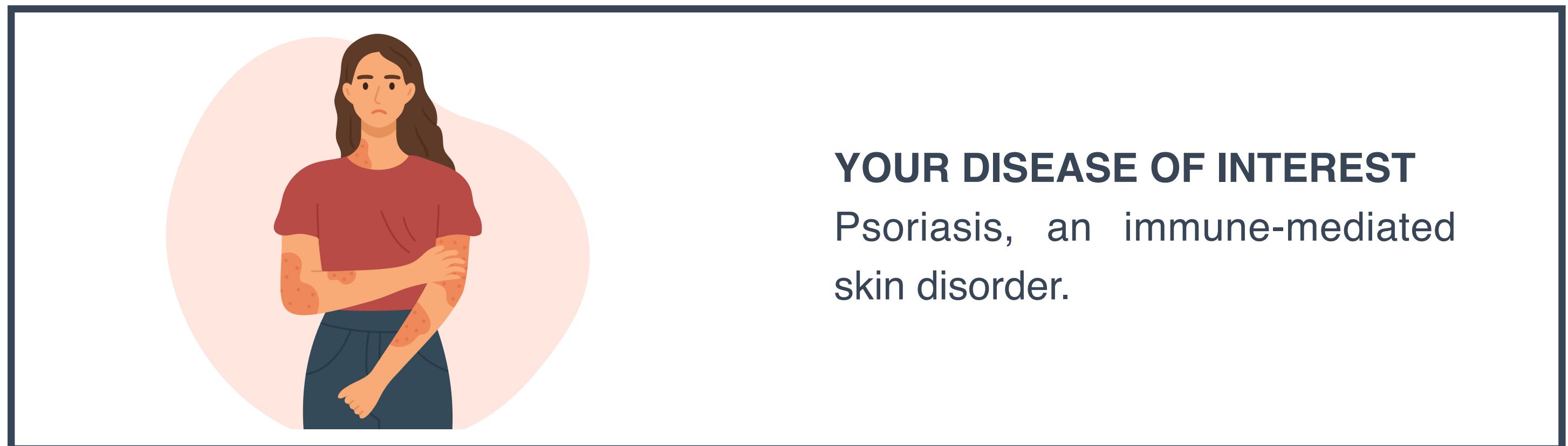
YOU

The researcher with R skills!



YOUR DATA

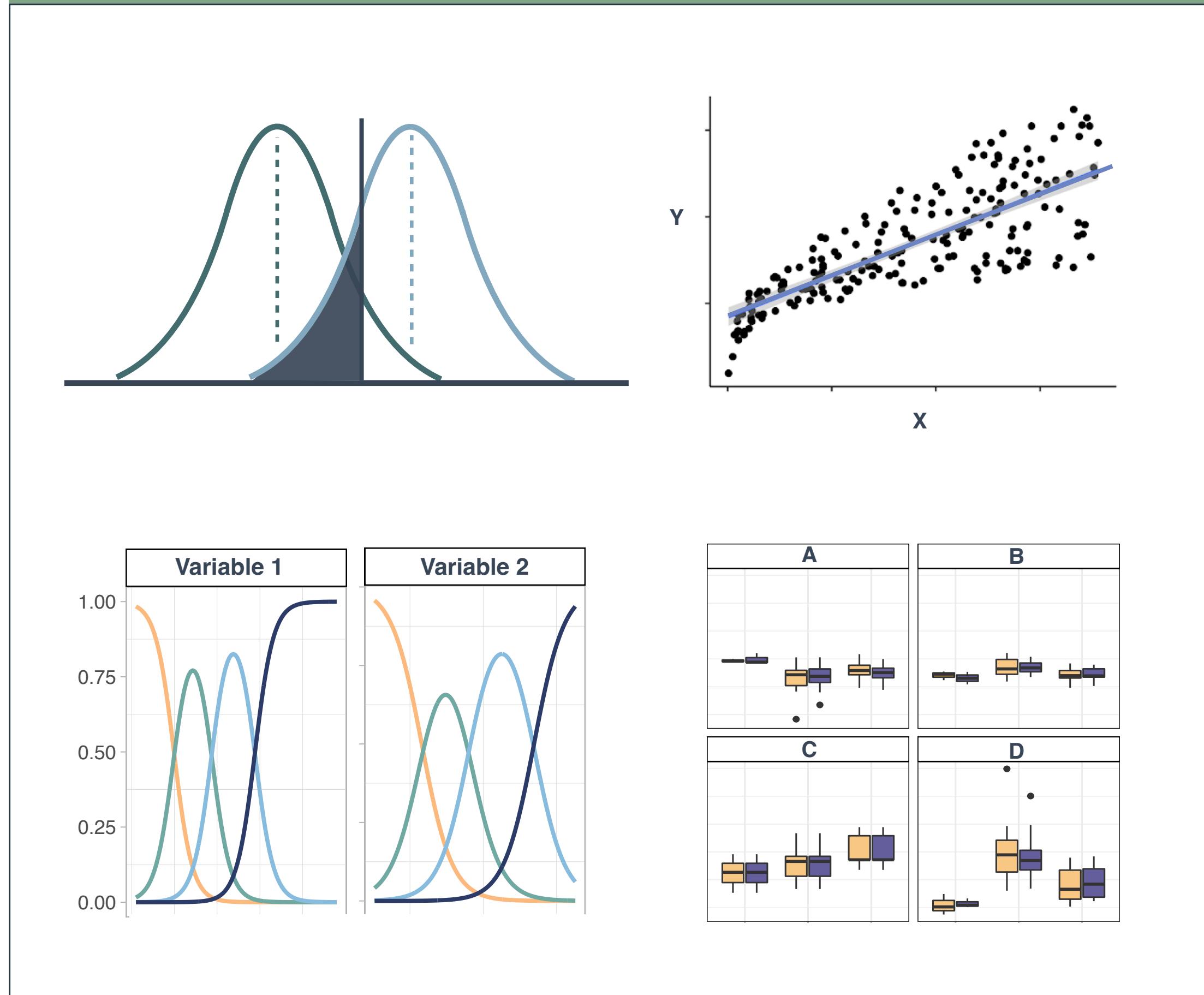
Gene expression data, i.e. how much a gene is used to produce a protein. Multiple genes measured in one experiment.



YOUR DISEASE OF INTEREST

Psoriasis, an immune-mediated skin disorder.

R - A STATISTICAL SCRIPTING LANGUAGE



During this session:

- *Cooperatively* discuss and share ideas about the data
- Apply steps of basic statistical analysis for hypothesis testing consistent with the given data
- Suggest conclusions based on your analysis, regarding the association between psoriasis and gene expression levels

STATS CHEAT SHEET

Import Data:

```
read_excel("my.data.xlsx")
```

Overview of Data:

```
summary(my.data)  
nrow(my.data)
```

```
length(my.data)  
names(my.data)
```

Linear:

```
lm(y~x, data=my.data)  
confint(model)
```

Logistic:

```
glm(y~x,  
data=my.data)
```

Linear Mixed:

```
lmer(y~x + (1|z),  
data=my.data)
```

Check Model:

```
summary(model)  
par(mfrow=c(2,2))  
plot(model)
```

ANOVA:

```
anova(model2, model1)
```

F-Test:

```
drop1(model, test="F")
```

Emmeans:

```
emmeans(model, ~x)  
pairs(emmeans(model, ~x))
```

Check Type:

```
table(my.data$x)  
is.numeric(my.data$x)  
is.factor(my.data$x)
```

Change Type:

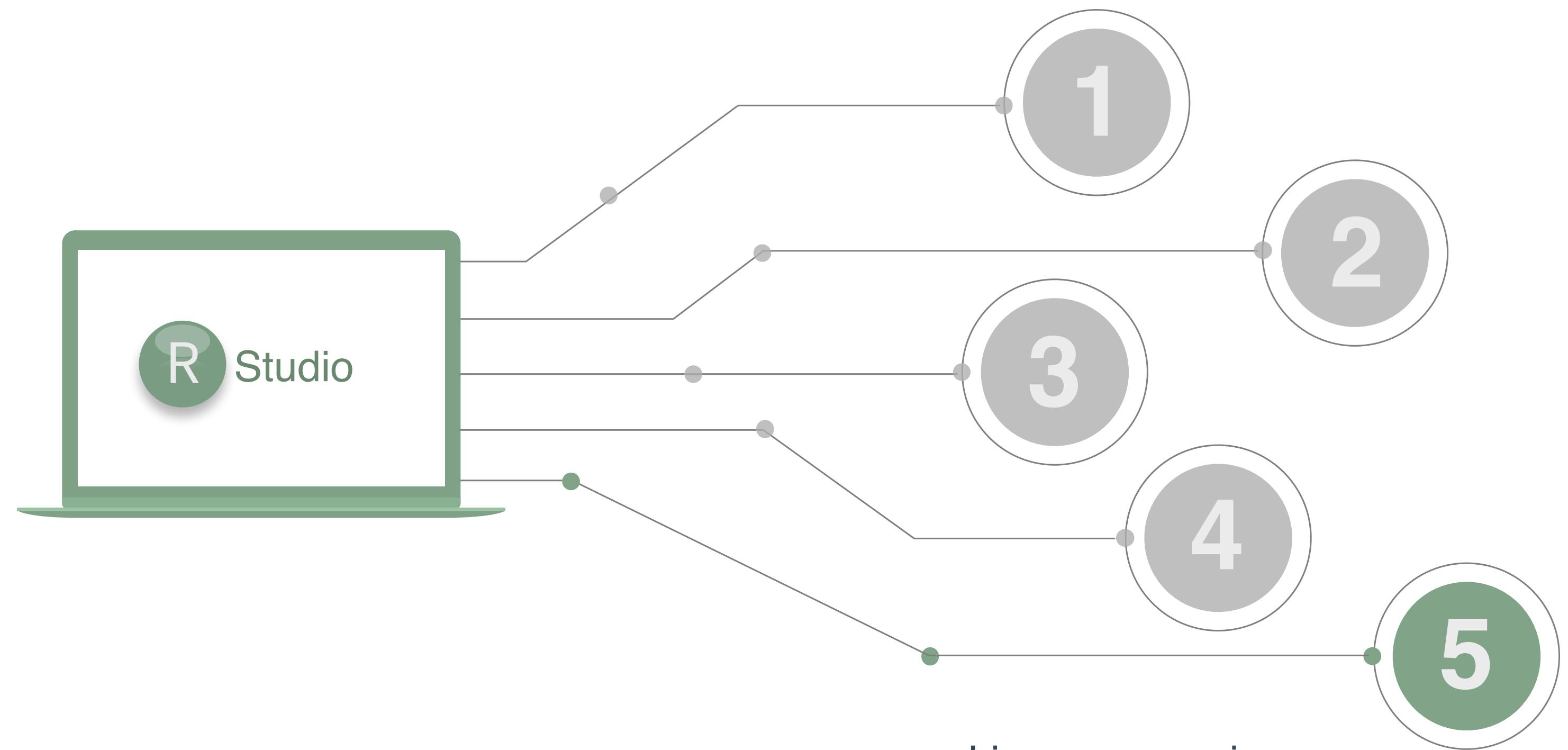
```
my.data <- mutate(my.data, x = factor(x))  
my.data$z <- as.numeric(my.data$z)
```

GET
STARTED

REGRESSION
MODELS

TESTS/
COMPARISONS

VARIABLES

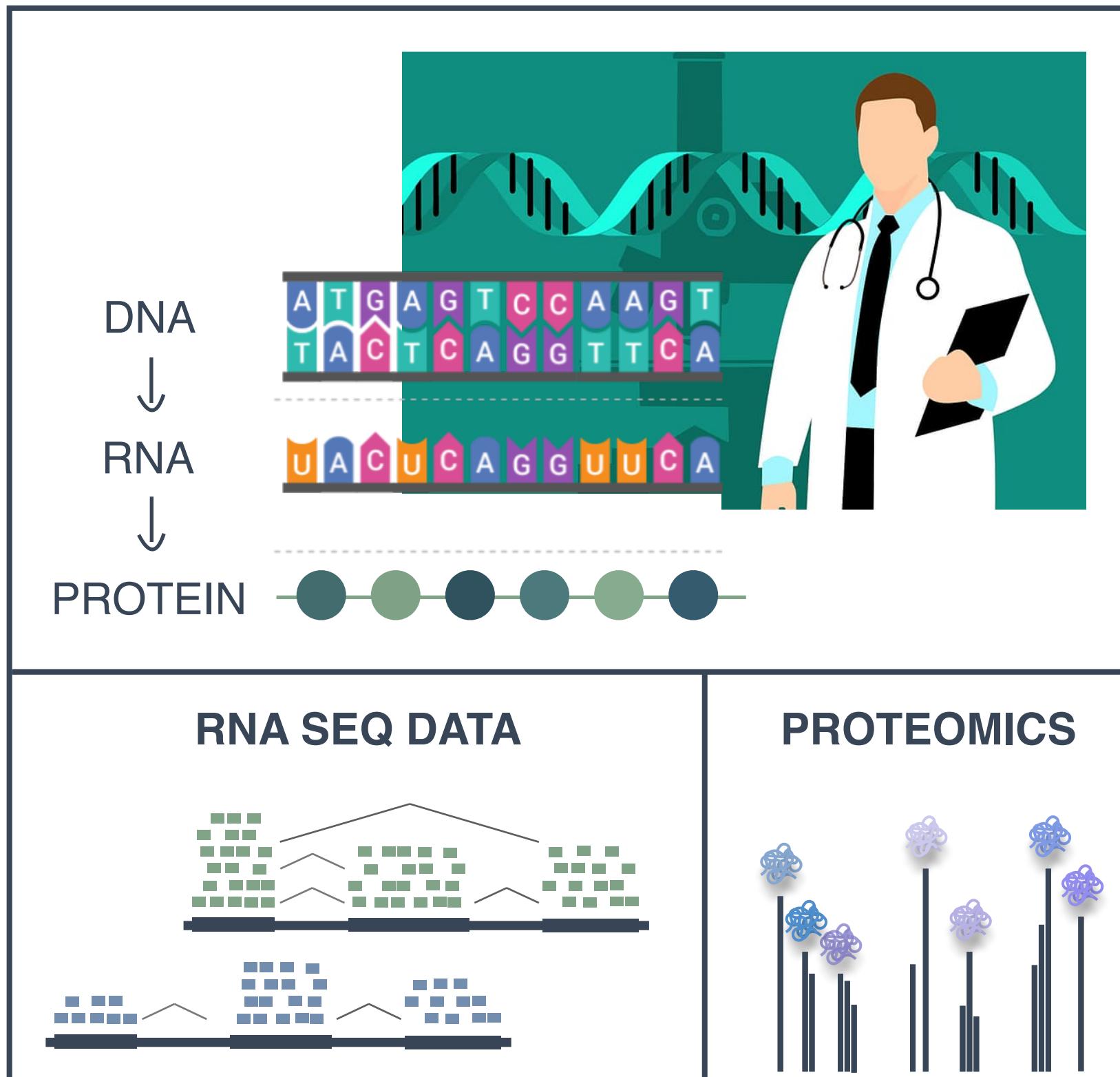


- Linear regression
- Summary Statistics
- ANOVA
- Logistic regression
- Clustering
- Correlation

— Statistics in R
EXERCISE 5

BIOINFORMATICS IN R

HIGH THROUGHPUT DATA



BIOINFORMATIC ANALYSIS

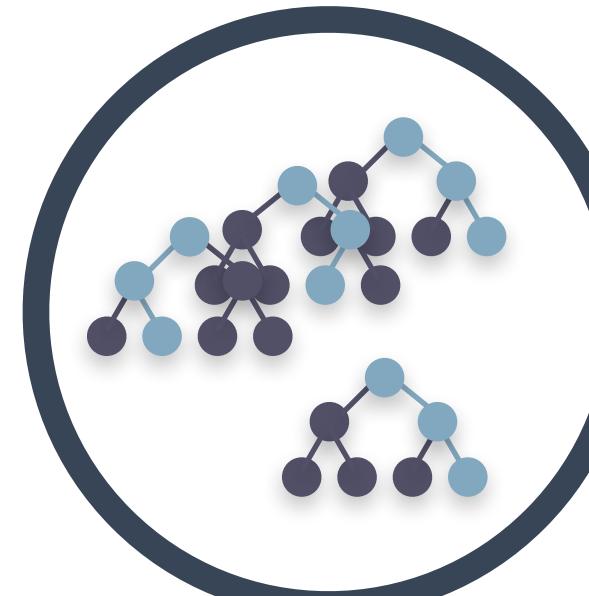
DIMENSIONALITY REDUCTION



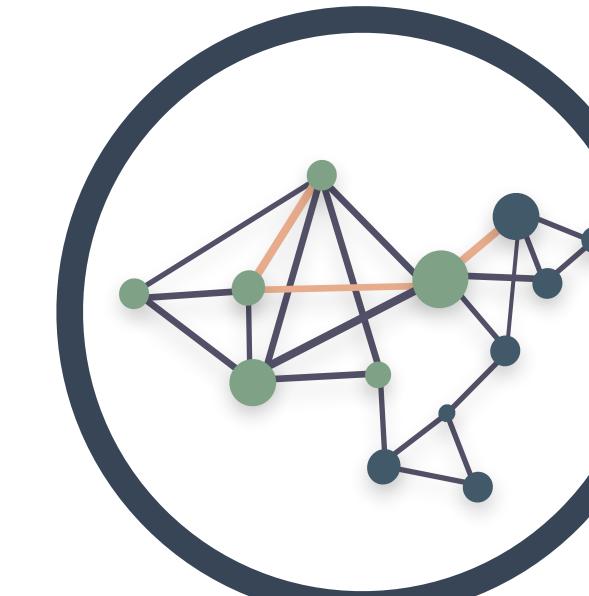
CLUSTERING



MACHINE LEARNING



NETWORK ANALYSIS



THE TOP OF THE R ICEBERG



STATISTICAL ANALYSIS

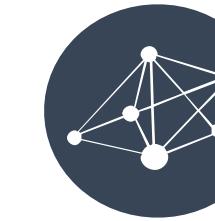
- Statistical models (linear, generalized, mixed, ...)
- Statistical tests (t-test, chisq, anova, ...)
- Survival analysis (Cox, Kaplan meier)



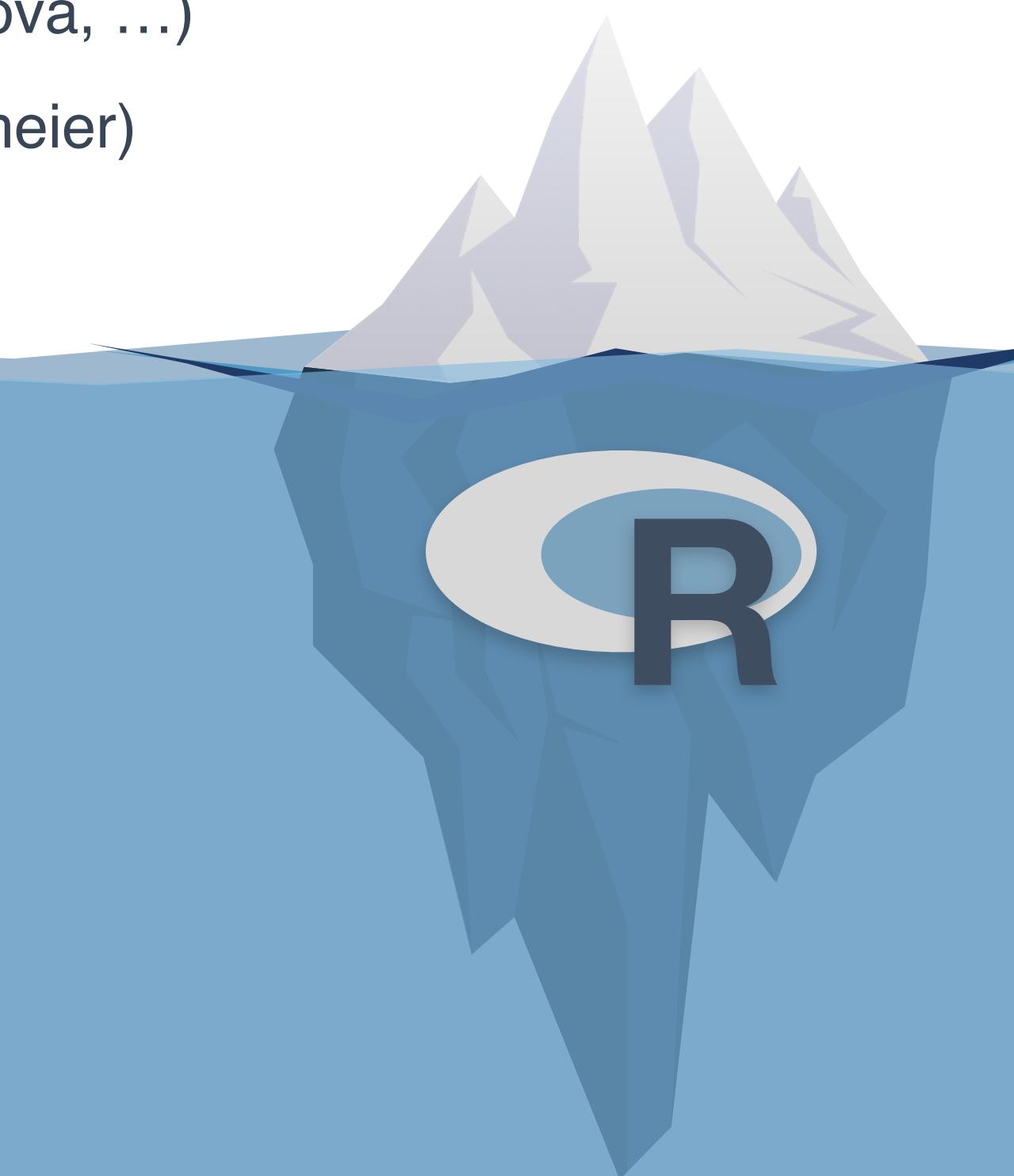
DATA MANGEMENT



EASY PLOTTING



BIOINFORMATIC ANALYSIS



— FROM EXCEL TO R

WANT MO-R-E?



The Section for Biostatistics offers a number statistics-oriented R courses:

Spring

[Basic Statistics for Health Researchers \(Danish course\)](#)

ECTS: 9,0

[Epidemiological methods in medical research](#)

ECTS: 7

[Advanced topics in health research B](#)

ECTS: 2,8

[Statistical methods in bioinformatics](#)

ECTS: 3,5

[Statistical analysis of survival data](#)

ECTS: 4,9

<https://publichealth.ku.dk/about-the-department/biostat/>

[Programming and statistical modelling in R](#)

ECTS: 1,6

[Bayesian methods in biomedical research](#)

ECTS: 2,4

[Psychometric validation of patient reported outcome measures](#)

ECTS: 2,5

[Introduction to validation of patient reported outcome measures](#)

[Causal inference I](#)

ECTS: 2,5

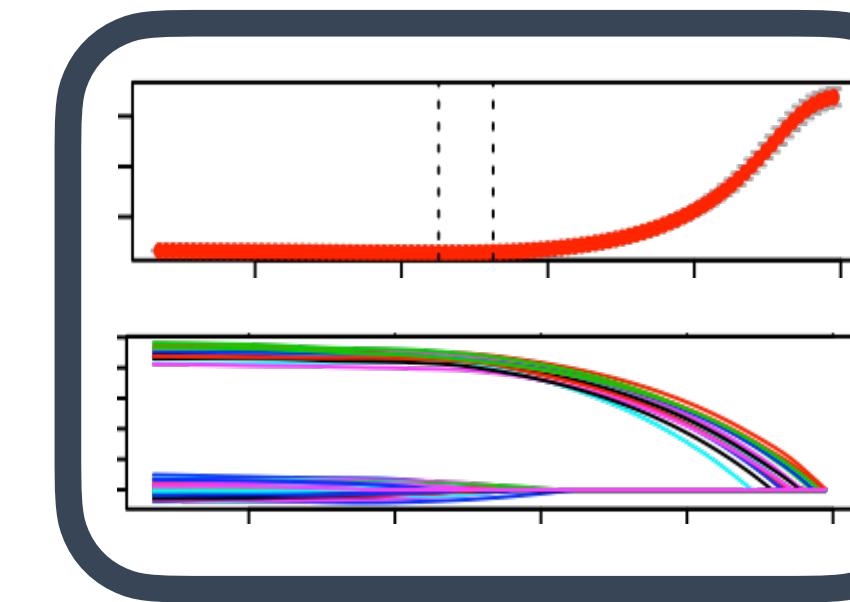
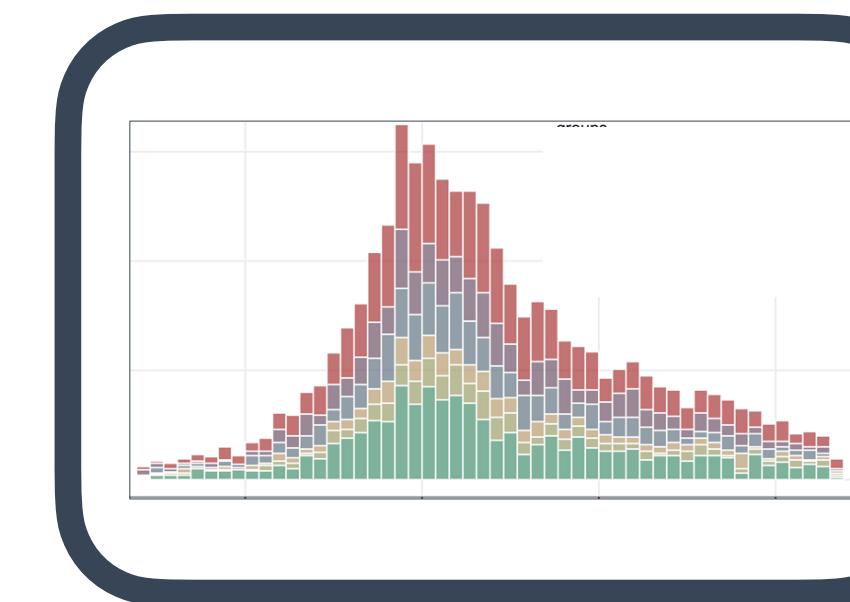
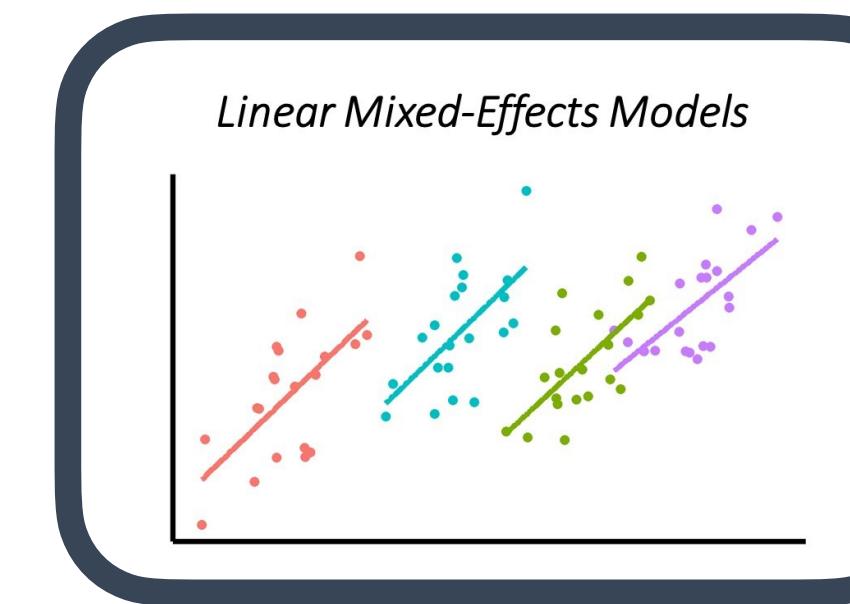
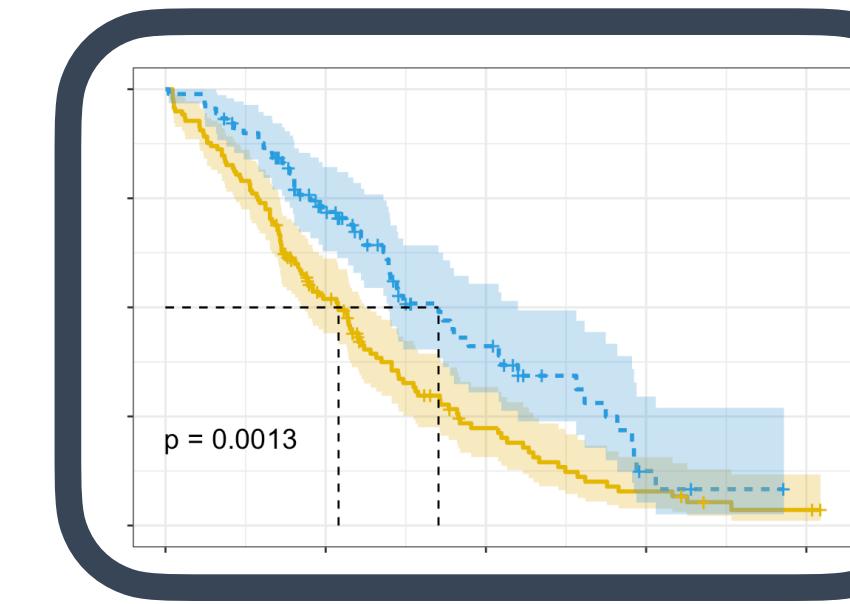
[Use of the statistical software R](#)

ECTS: 2,1

* These are screenshots. Go to the website and scroll down to ‘Teaching’



— TEASER STATISTICS in R



Survival Analysis

`survival`: <https://rviews.rstudio.com/2017/09/25/survival-analysis-with-r/>

`survminer`: <https://cran.r-project.org/web/packages/survminer/survminer.pdf>
(<https://rpkgs.datanovia.com/survminer/>)

Mixed-Effects Models

`lme4`: <https://cran.r-project.org/web/packages/lme4/vignettes/lmer.pdf>

<https://cran.microsoft.com/snapshot/2017-08-01/web/packages/sjPlot/vignettes/sjplmer.html>

`glmmTMB`: <https://cran.r-project.org/web/packages/glmmTMB/index.html>

Epidemiological Analysis

`Epi`: <https://cran.r-project.org/web/packages/Epi/index.html>

`pubh`: <https://rviews.rstudio.com/2020/03/05/covid-19-epidemiology-with-r/>

https://cran.r-project.org/web/packages/incidence/vignettes/customize_plot.html

<https://rviews.rstudio.com/2020/03/05/covid-19-epidemiology-with-r/>

Elastic-Net Regression

`glmnet`: <https://cran.r-project.org/web/packages/glmnet/glmnet.pdf>

`elasticnet`: <https://cran.r-project.org/web/packages/elasticnet/elasticnet.pdf>

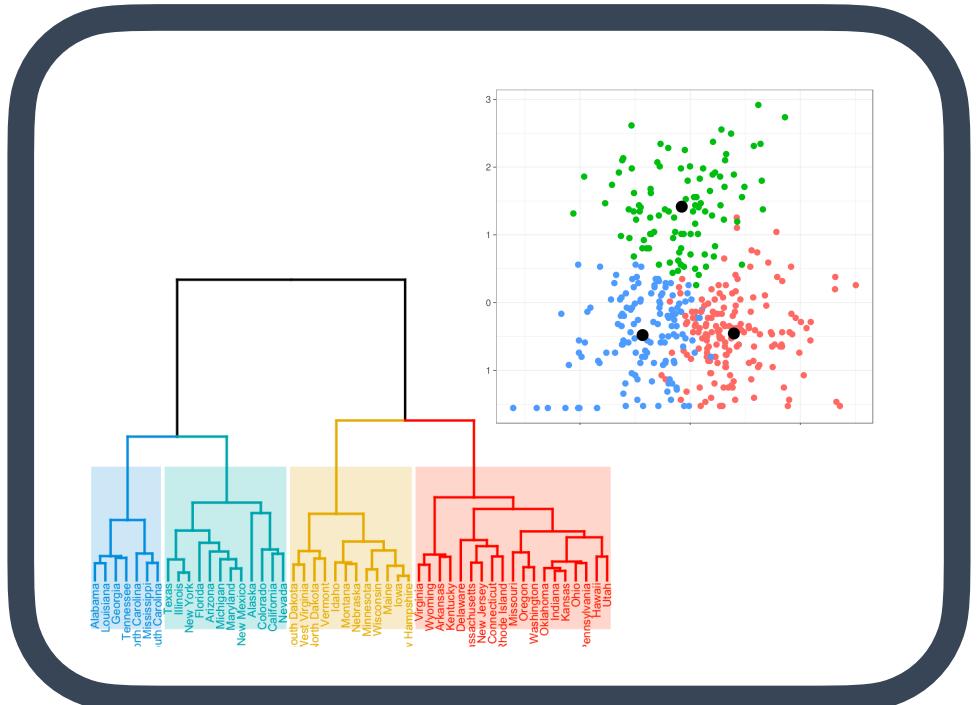
<https://www.datacamp.com/community/tutorials/tutorial-ridge-lasso-elastic-net>

TEASER

Machine Learning

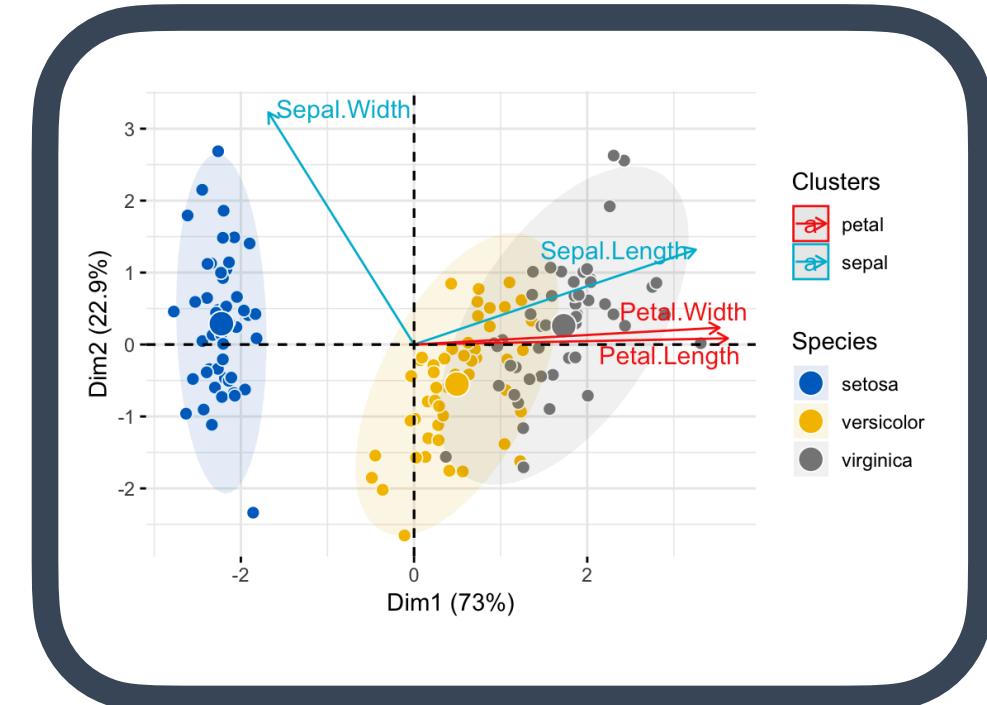
<https://lgatto.github.io/IntroMachineLearningWithR/an-introduction-to-machine-learning-with-r.html>

Clustering



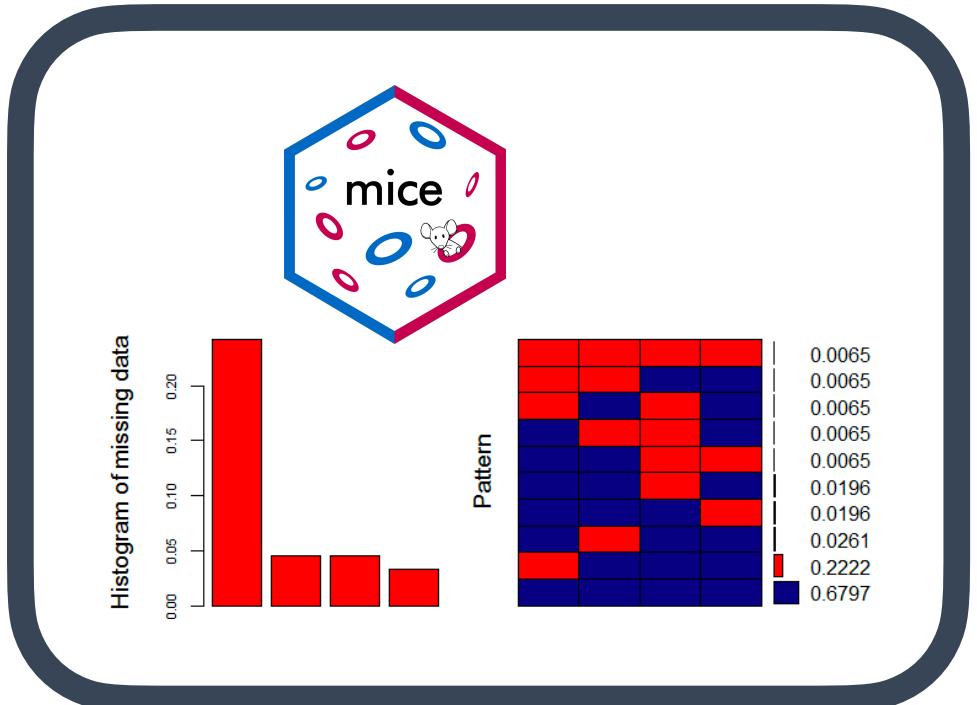
<https://statsandr.com/blog/clustering-analysis-k-means-and-hierarchical-clustering-by-hand-and-in-r/>

Feature Selection: PCA



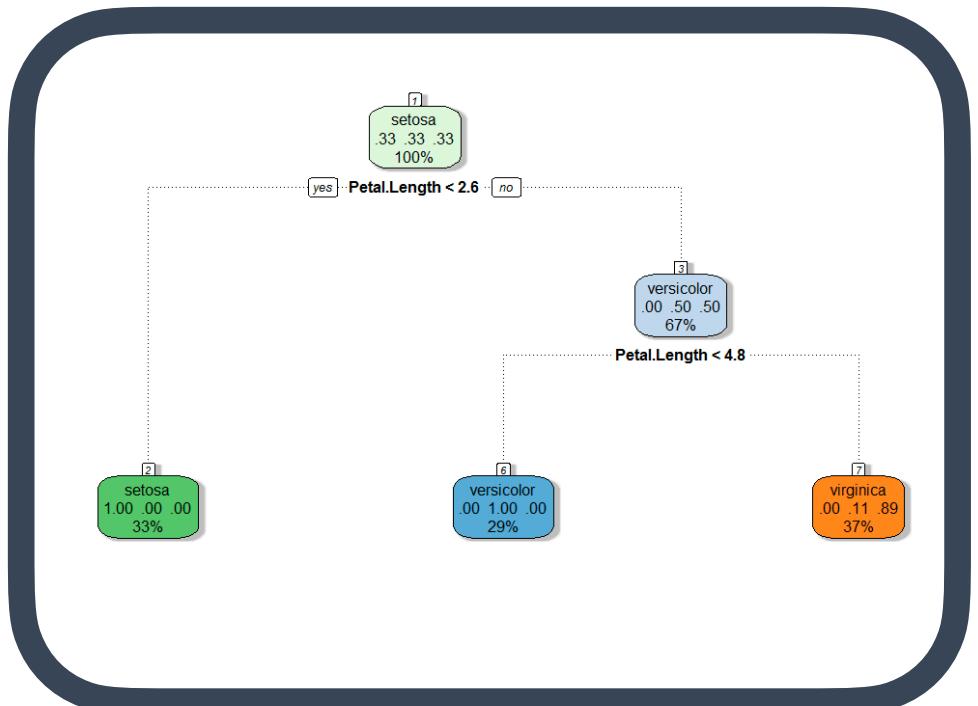
<https://bioconductor.org/packages/release/bioc/vignettes/PCATools/inst/doc/PCATools.html>

Missing Data



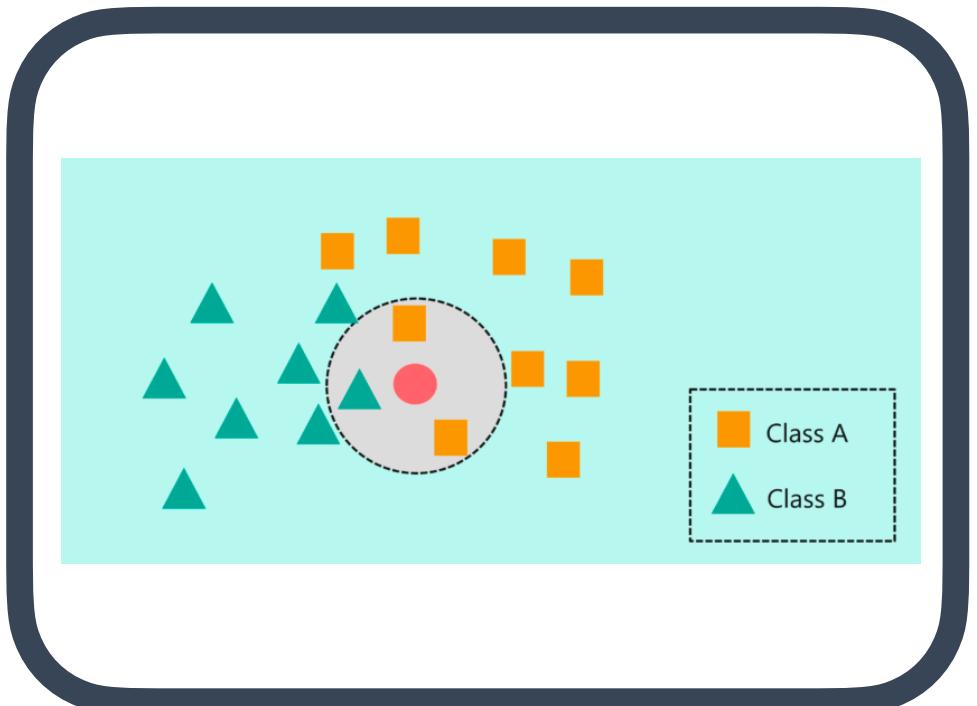
<https://amices.org/mice/>
<https://datascienceplus.com/imputing-missing-data-with-r-mice-package/>

Random Forest



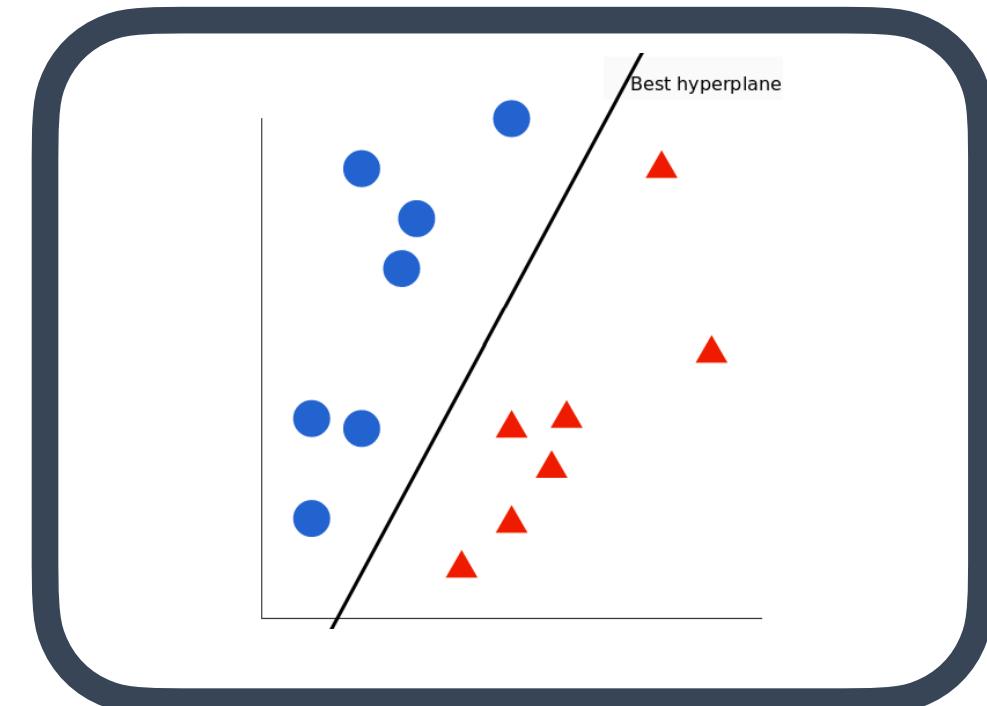
<https://www.blopig.com/blog/2017/04/a-very-basic-introduction-to-random-forests-using-r/>

kNN



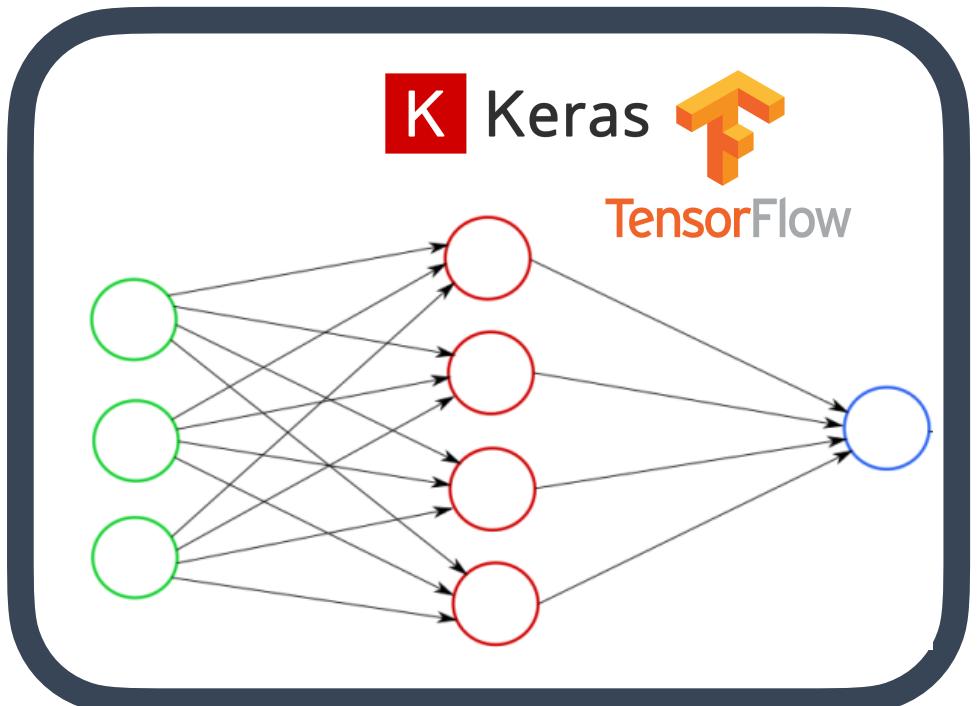
<https://www.edureka.co/blog/knn-algorithm-in-r/>

SVM



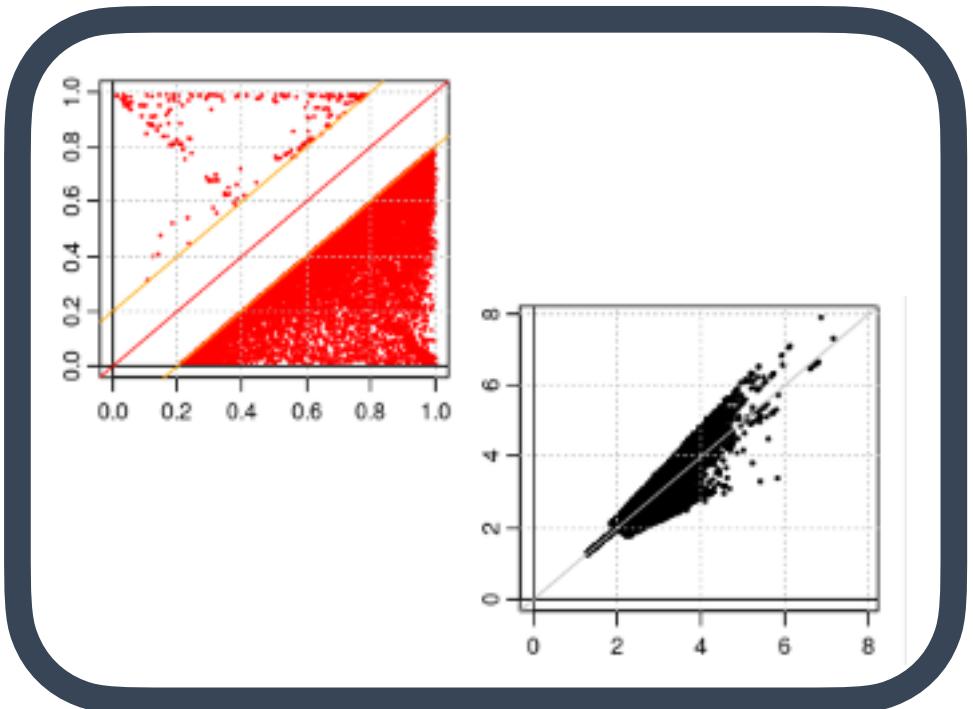
<https://cran.r-project.org/web/packages/e1071/vignettes/svmdoc.pdf>

Neural Networks



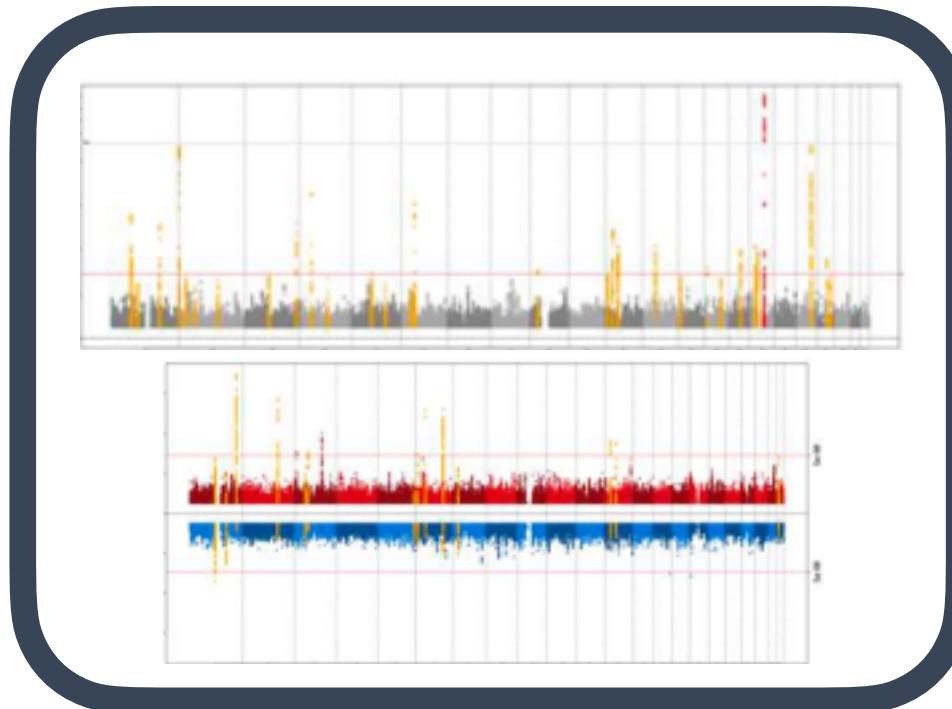
<https://keras.rstudio.com/>
<https://tensorflow.rstudio.com/>

GWAS - QC & Data Harmonization



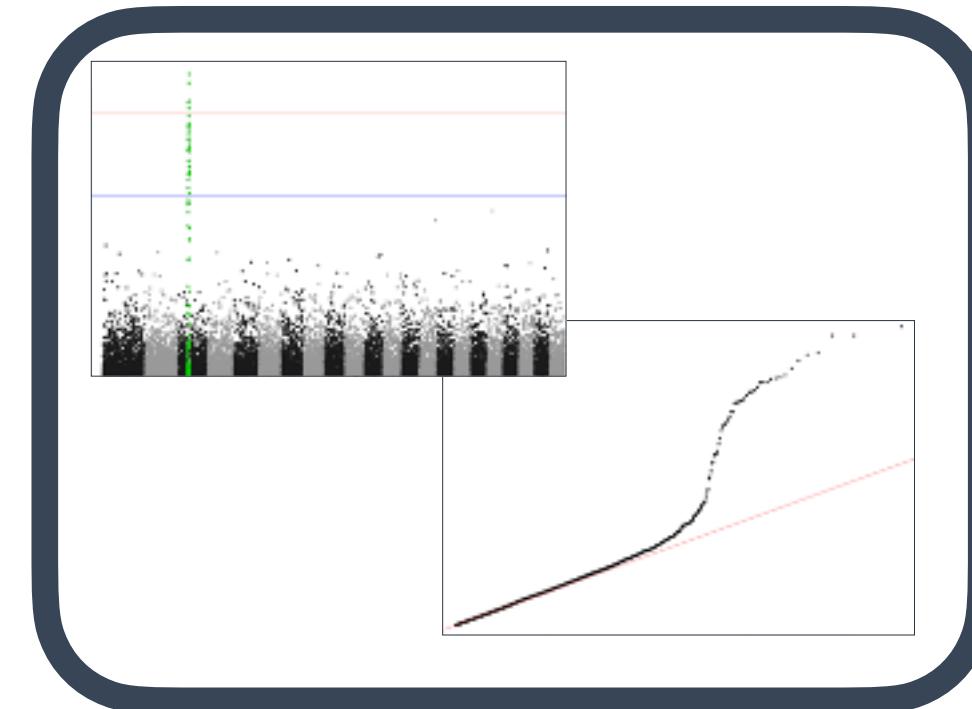
EasyQC: <https://www.uni-regensburg.de/medizin/epidemiologie-praeventivmedizin/genetische-epidemiologie/software/>

GWAS Data Management & Plots



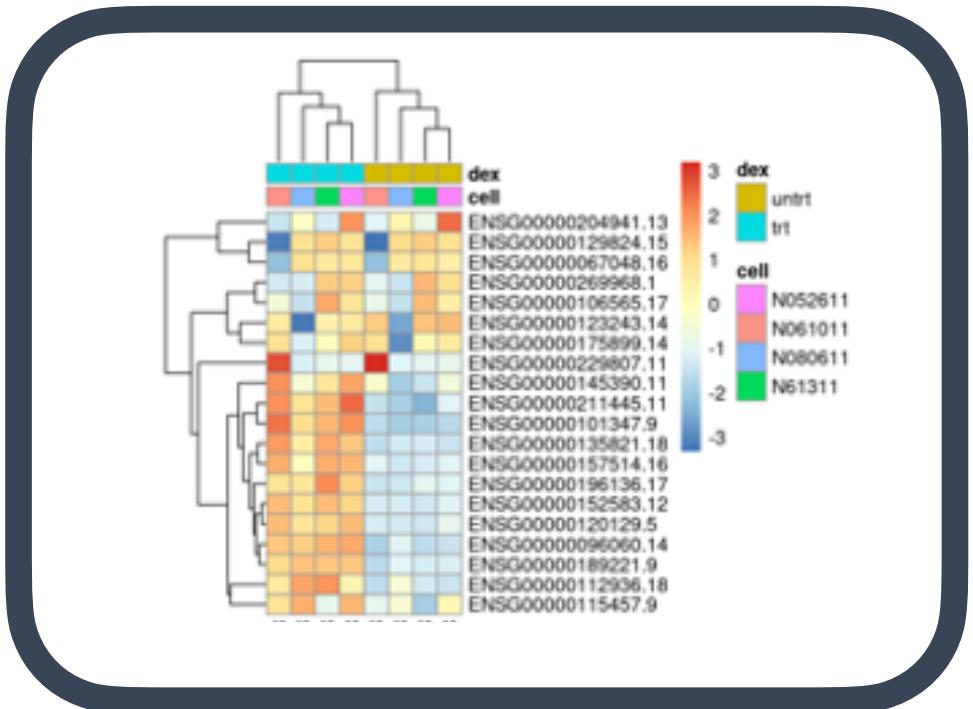
EasyStrata: <https://www.uni-regensburg.de/medizin/epidemiologie-praeventivmedizin/genetische-epidemiologie/software/>

More Plotting...



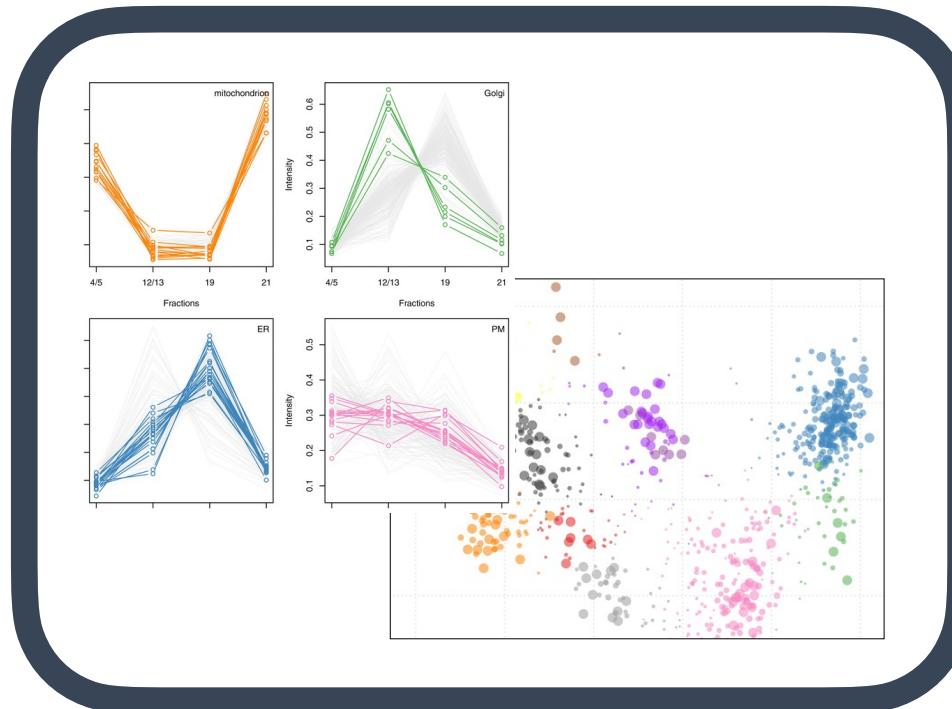
Manhattan and QQ plots: <https://cran.r-project.org/web/packages/qqman/vignettes/qqman.html>

Gene Expression Analysis



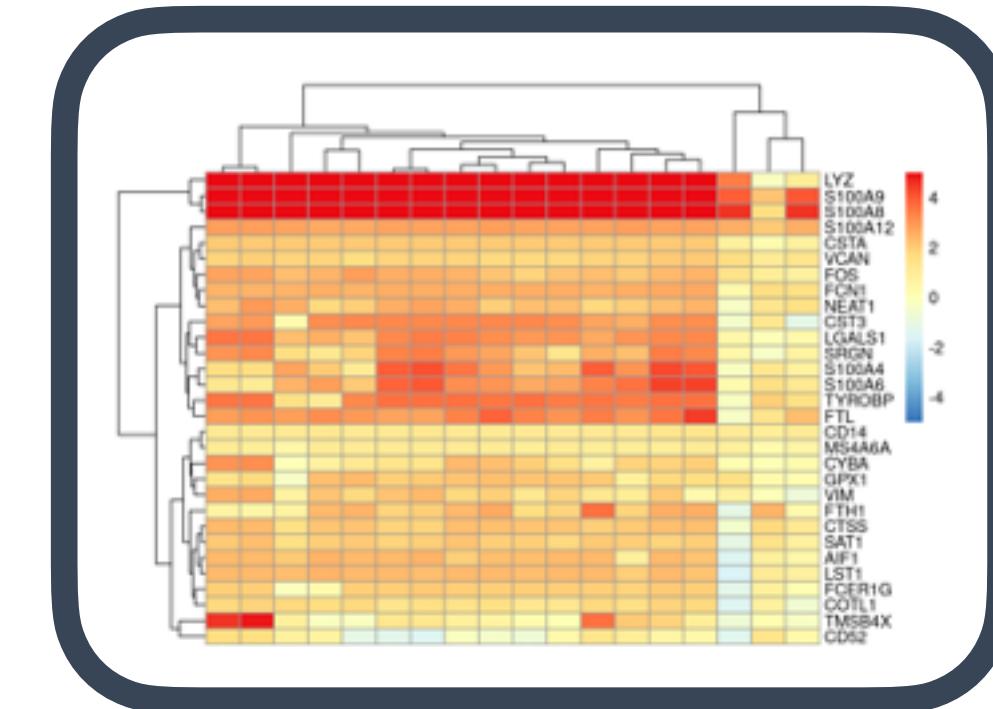
DESeq2, limma, EdgeR, etc.: http://www.bioconductor.org/packages/release/BiocViews.html#_RNASeq

Proteomics Analysis



RforProteomics: http://www.bioconductor.org/packages/release/BiocViews.html#_Proteomics

Single-Cell RNASeq



<https://cran.r-project.org/web/packages/e1071/vignettes/svmdoc.pdf>

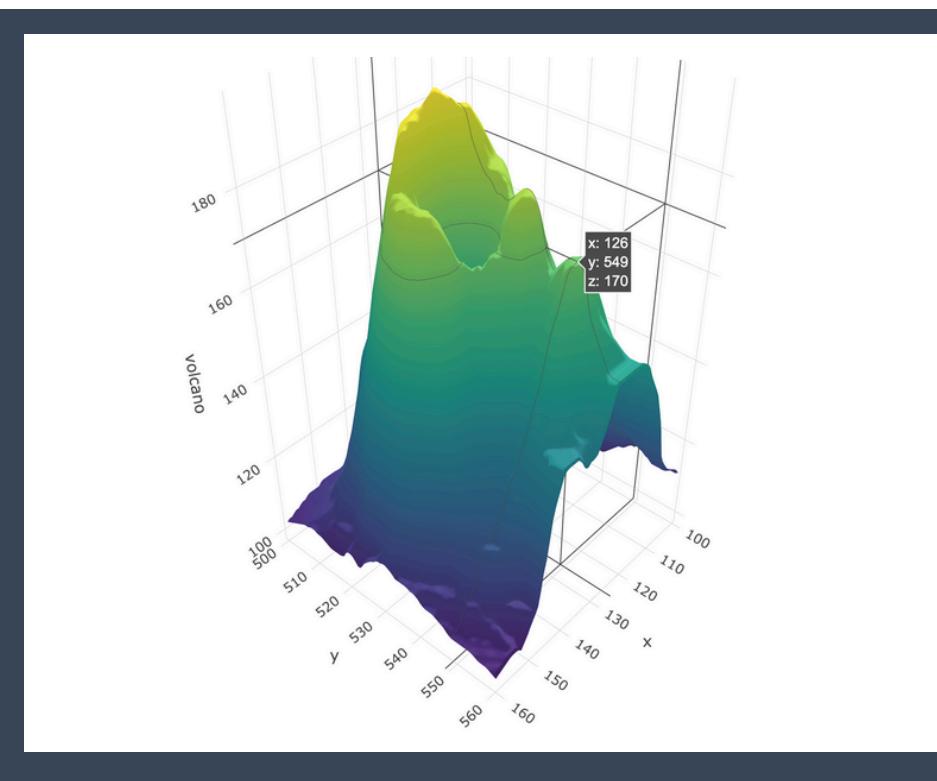
— TEASER Omics Data

<http://www.bioconductor.org/packages/release/BiocViews.html>

COOL STUFF IN R

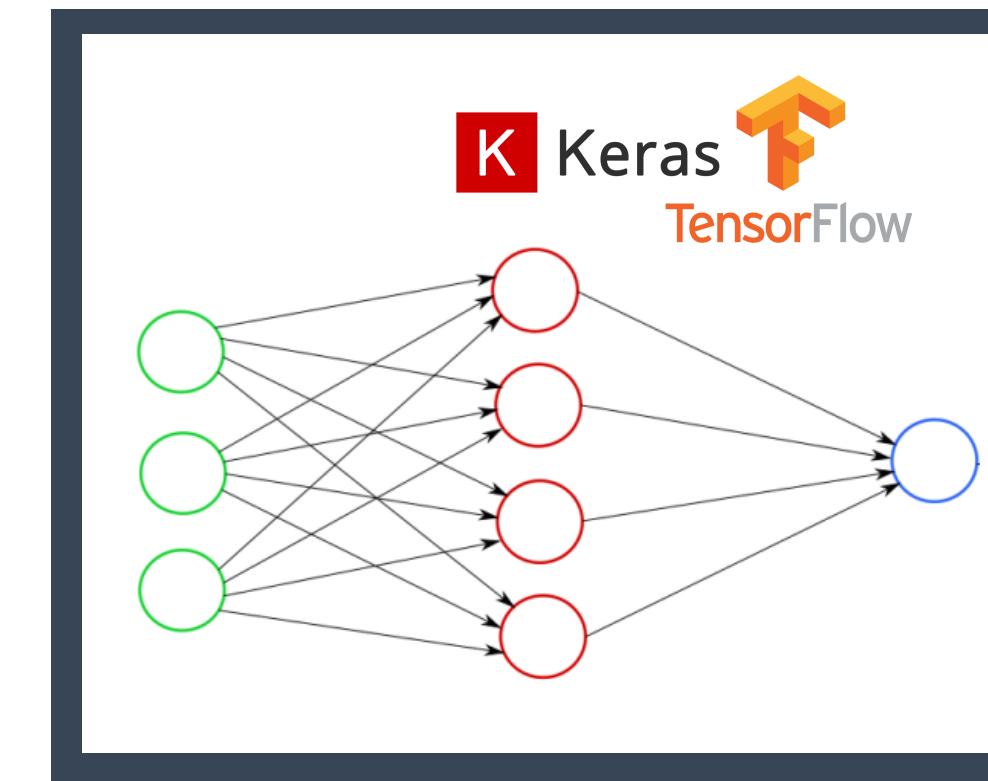
— FROM EXCEL TO R

PLOTTING IN 3D



<https://plotly-r.com/d-charts.html>

DEEP LEARNING



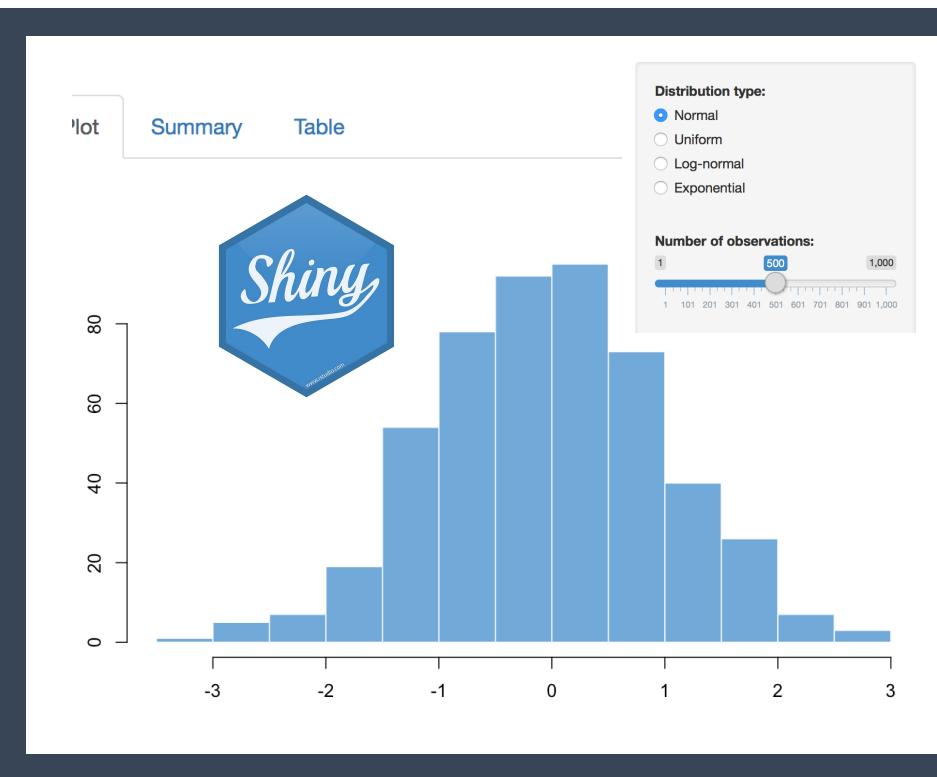
<https://keras.rstudio.com/>
<https://tensorflow.rstudio.com/>

BAYESIAN STATISTICS



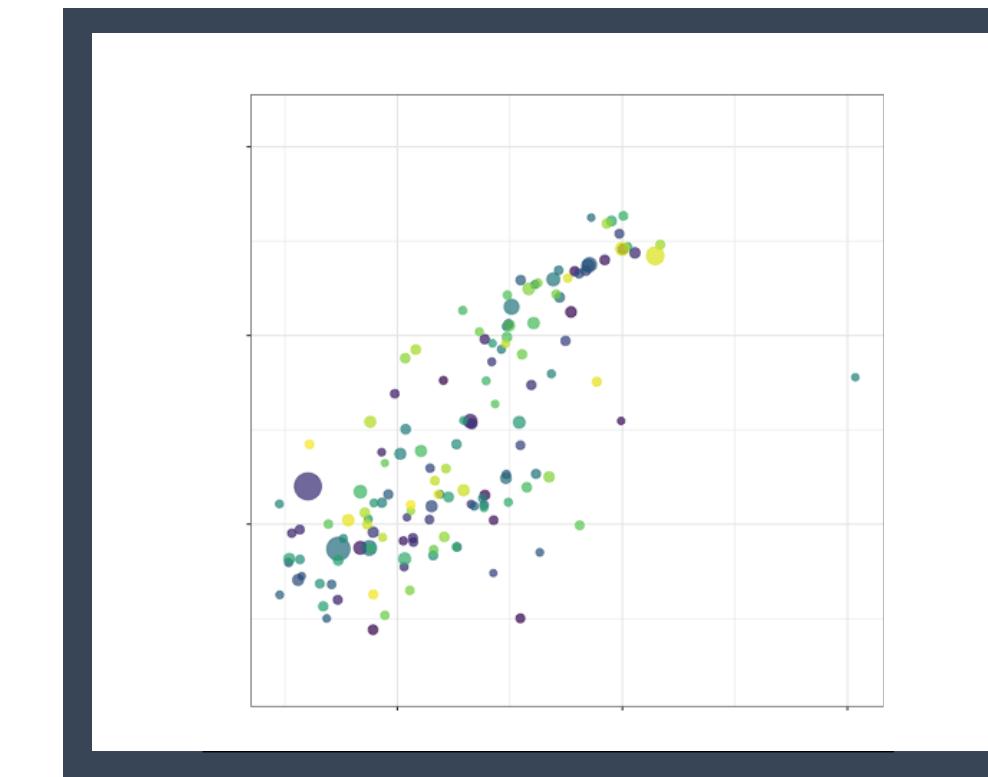
<https://mc-stan.org/users/interfaces/rstan>

WEBPAGE WITH R SHINY



<https://shiny.rstudio.com/>

INTERACTIVE PLOTS



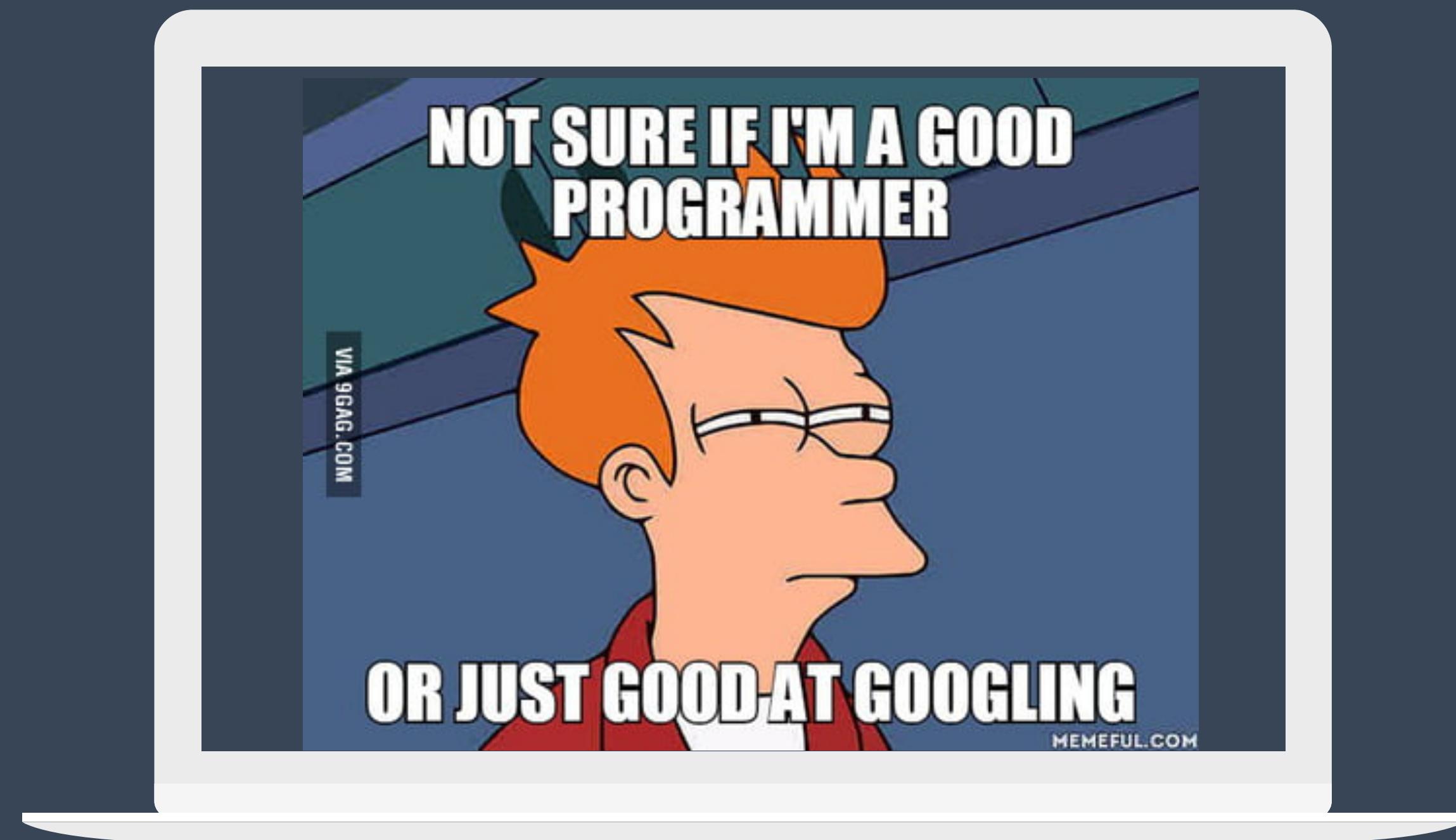
<https://gganimate.com/articles/gganimate.html>

MAIL AND MESSAGES



<https://github.com/briandconnelly/pushoverr>

THANK YOU FOR LISTENING



This keynote presentation was created by Thilde Terkelsen,
Data Scientist, Center for Health Data Science, SUND, KU.
For internal use at KU only, do not distribute commercially.