

# Reproducible research

## Module 12

*Leonardo Collado-Torres*

*June 19, 2015*

### Reproducible research

- Have you heard about it?
- Maybe you've heard about the [Coursera Reproducible Research](#) course
- How would you define it?
- Is it the same as replicability?
- Is it important?

Science moves forward then discoveries are replicated and reproduced

*[Implementing Reproducible Research](#)*

### Replication

Replication, the practice of independently implementing scientific experiments to validate specific findings, is the cornerstone of discovering scientific truth.

*[Implementing Reproducible Research](#)*

### Reproducibility

Reproducibility can be thought of as a different standard of validity from replication because it forgoes independent data collection and uses the methods and data collected by the original investigator.

*[Implementing Reproducible Research](#)*

### A bit more practical

The sharing of analytic data and computer codes uses to map those data into computational results is central to any comprehensive definition of reproducibility.

*[Implementing Reproducible Research](#)*

## Why its important?

Except for the simplest of analyses, the computer code used to analyze a dataset is the only record that permits others to fully understand what a researcher has done.

### *Implementing Reproducible Research*

## Scrapping example

- Open [Scrapping Ravens data from ESPN](#)
- Is this document reproducible?
- What information is missing?
- How could we improve it?
- What about [rMaps Mexico map](#)?

## R session information

```
sessionInfo()
```

```
## R version 3.2.0 Patched (2015-05-18 r68382)
## Platform: x86_64-apple-darwin10.8.0 (64-bit)
## Running under: OS X 10.8.5 (Mountain Lion)
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] devtools_1.8.0 knitr_1.10.5  rmarkdown_0.7
##
## loaded via a namespace (and not attached):
## [1] Rcpp_0.11.6      digest_0.6.8    git2r_0.10.1    formatR_1.2
## [5] magrittr_1.5     evaluate_0.7     stringi_0.4-1   curl_0.8
## [9] rstudioapi_0.3.1 xml2_0.1.1      tools_3.2.0     stringr_1.0.0
## [13] yaml_2.1.13      rversions_1.0.1 memoise_0.2.1   htmltools_0.2.6
```

```
library('devtools'); session_info()
```

```
## Session info -----
```

```

## setting value
## version R version 3.2.0 Patched (2015-05-18 r68382)
## system x86_64, darwin10.8.0
## ui AQUA
## language (EN)
## collate en_US.UTF-8
## tz America/New_York

## Packages -----

## package * version date source
## curl 0.8 2015-06-06 CRAN (R 3.2.0)
## devtools * 1.8.0 2015-05-09 CRAN (R 3.2.0)
## digest 0.6.8 2014-12-31 CRAN (R 3.2.0)
## evaluate 0.7 2015-04-21 CRAN (R 3.2.0)
## formatR 1.2 2015-04-21 CRAN (R 3.2.0)
## git2r 0.10.1 2015-05-07 CRAN (R 3.2.0)
## htmltools 0.2.6 2014-09-08 CRAN (R 3.2.0)
## knitr * 1.10.5 2015-05-06 CRAN (R 3.2.0)
## magrittr 1.5 2014-11-22 CRAN (R 3.2.0)
## memoise 0.2.1 2014-04-22 CRAN (R 3.2.0)
## Rcpp 0.11.6 2015-05-01 CRAN (R 3.2.0)
## rmarkdown * 0.7 2015-06-13 CRAN (R 3.2.0)
## rstudioapi 0.3.1 2015-04-07 CRAN (R 3.2.0)
## rversions 1.0.1 2015-06-06 CRAN (R 3.2.0)
## stringi 0.4-1 2014-12-14 CRAN (R 3.2.0)
## stringr 1.0.0 2015-04-30 CRAN (R 3.2.0)
## xml2 0.1.1 2015-06-02 CRAN (R 3.2.0)
## yaml 2.1.13 2014-06-12 CRAN (R 3.2.0)

```

## Reproducible documents

- Have you ever had your code in one file, your description of the results in another file?
- Ever made copy-paste mistakes?
- What if you were asked to change some models or revise the document?
- Was it easy to maintain?



- What would be a reproducible document for you?

## Reproducible docs in R

- [R Markdown](#) is the easiest
- It's based on [Markdown](#): simple human readable syntax that generates HTML docs
- Also uses [Pandoc](#) to create files in many formats
- You maintain a **single** file! It has the code, figures and description of results. It then creates a file in the format you want to share with others.

# Markdown

The image shows a side-by-side comparison of a Markdown document and its rendered HTML output in RStudio. The left pane, titled 'example.Rmd', contains the following text:

```
1 Header 1
2 -----
3 This is an R Markdown document. Markdown is a
4 | simple formatting syntax for authoring web pages.
5 Use an asterisk mark, to provide emphasis such as
6 | italics and bold.
7 Create lists with a dash:
8 - Item 1
9 - Item 2
10 - Item 3
11
12 You can write `in-line` code with a back-tick.
13
14 ```
15 Code blocks display
16 with fixed-width font
17 ```
18
19 > Blockquotes are offset
20
```

The right pane, titled 'RStudio: Preview HTML', shows the rendered output:

## Header 1

This is an R Markdown document. Markdown is a simple formatting syntax for authoring web pages.

Use an asterisk mark, to provide emphasis such as *italics* and **bold**.

Create lists with a dash:

- Item 1
- Item 2
- Item 3

You can write in-line code with a back-tick.

```
Code blocks display
with fixed-width font
```

Blockquotes are offset

## Markdown + R code

The screenshot shows the RStudio interface with two windows. The left window, titled 'chunks.Rmd', displays R code chunks. The right window, titled 'RStudio: Preview HTML', shows the rendered HTML output of the code.

**Left Window (chunks.Rmd):**

```
1 R Code Chunks
2 =====
3
4 With R Markdown, you can insert R code
5 chunks including plots:
6 ```{r qplot, fig.width=4, fig.height=3,
7 message=FALSE}
8 # quick summary and plot
9 library(ggplot2)
10 summary(cars)
11 qplot(speed, dist, data=cars) +
12   geom_smooth()
13
```

**Right Window (RStudio: Preview HTML):**

### R Code Chunks

With R Markdown, you can insert R code chunks including plots:

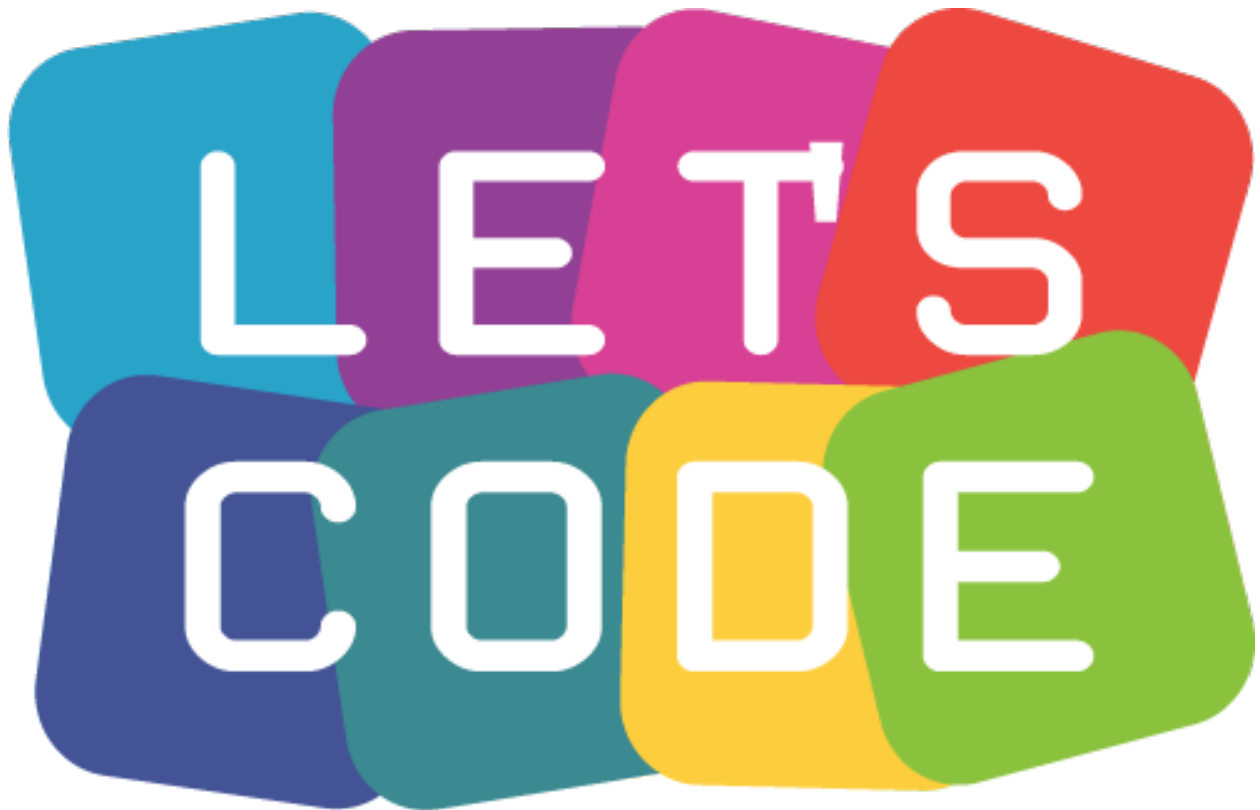
```
# quick summary and plot
library(ggplot2)
summary(cars)
```

##	speed	dist
##	Min. : 4.0	Min. : 2
##	1st Qu.:12.0	1st Qu.: 26
##	Median :15.0	Median : 36
##	Mean :15.4	Mean : 43
##	3rd Qu.:19.0	3rd Qu.: 56
##	Max. :25.0	Max. :120

```
qplot(speed, dist, data = cars) + geom_smooth()
```

## Examples

- [R Programming for Data Science](#) (can buy for \$0.00)
- [Advanced R](#), for example [Data structures](#)
- [Polyester: simulating RNA-seq datasets with differential transcript expression](#) paper which creates [this html](#).
- [Timing information report](#) which renders into [this html](#). Note that it has an interactive table.



## Misc

- [R Markdown: where to go to learn more](#)
- [Example R Markdown doc](#)
- [An example interactive document](#): beyond the scope of this course, but cool to know it exists.
- [Another intro to R Markdown presentation](#)
- [Where do I start using Bioconductor?](#)