

Doing reproducible science: from your hard-won data to a publishable manuscript without going mad

Francisco Rodriguez-Sanchez (@frod_san)

February 2017

A typical research workflow

1. Prepare data (**EXCEL**)

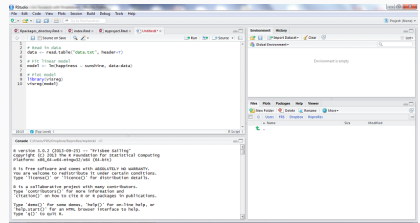
	A	B
1	happiness_index	sunshine_h
2	10.5	978.4
3	6.6	660.9
4	11.3	1093.5
5	9.6	978.9
6	10.9	1135.5
7	9.1	907.0
8	10.6	990.4
9	12.4	1172.9
10	9.6	1025.6
11	10.1	1055.0
12	10.9	1093.7
13	8.9	863.8
14	12.5	1196.6
15	10.0	995.8
16	11.0	1120.2
17	10.3	988.0
18	9.7	987.0
19	9.3	970.4
20	10.9	1076.6
21	9.0	909.8
22	7.7	733.4
23	9.0	985.2
24	10.4	1084.0
25	10.0	1066.7

data

Ready

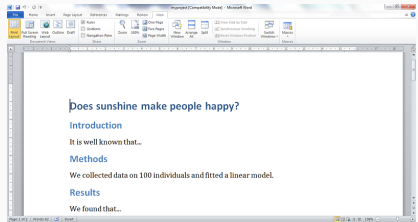
A typical research workflow

1. Prepare data (**EXCEL**)
2. Analyse data (**R**)



A typical research workflow

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2. Analyse data (**R**)
3. Write report/paper
(**WORD**)

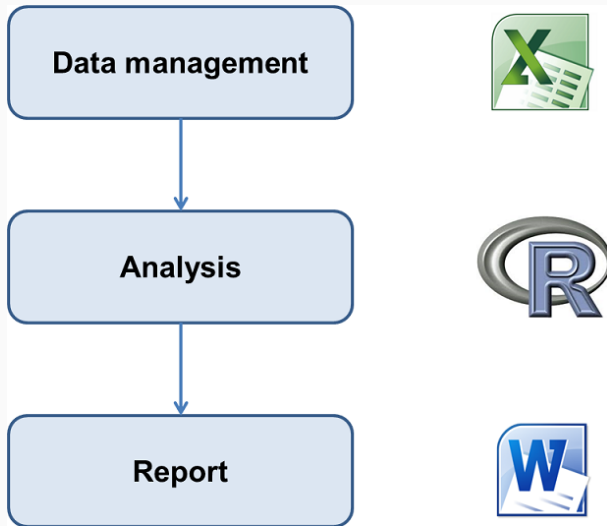


A typical research workflow

1. Prepare data (**EXCEL**)
2. Analyse data (**R**)
3. Write report/paper
(**WORD**)
4. Start the email attachments
nightmare...



This workflow is broken



Problems of a broken workflow

- How did you do this? What analysis is behind this figure? Did you account for ...?

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- What dataset was used? Which individuals were left out? Where is the clean dataset?

Problems of a broken workflow

- How did you do this? What analysis is behind this figure? Did you account for ...?
- What dataset was used? Which individuals were left out? Where is the clean dataset?
- Oops, there is an error in the data. Can you repeat the analysis? And update figures/tables in Word!



Trevor A. Branch

@TrevorABranch



Follow

My rule of thumb: every analysis you do on a dataset will have to be redone 10–15 times before publication. Plan accordingly. [#Rstats](#)

Our everyday scary movie

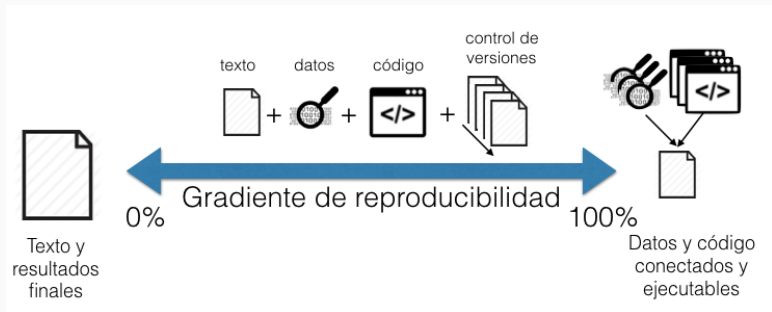
<https://youtu.be/s3JldKoA0zw>

WHAT is Reproducible Science?

A scientific article is **reproducible** if there is computer **code** that can **regenerate** all results and figures from the original data.

- Transparent
- Traceable
- Comprehensive
- Useful

Most science is not reproducible



Even **you** will struggle to reproduce **your own results** from a few weeks/months ago.

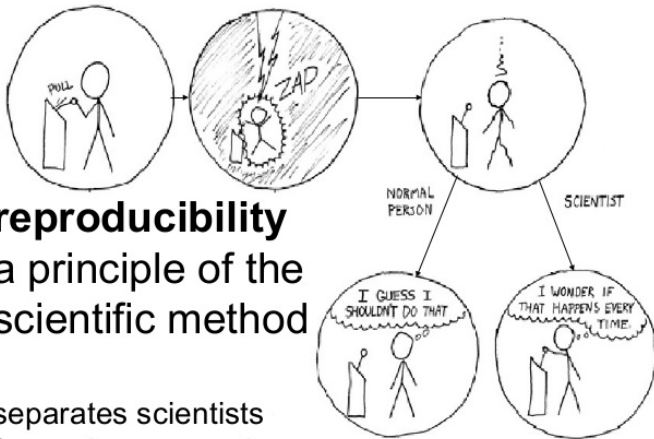
You can't reproduce if you don't understand where a number came from.

You can't reproduce what you don't remember. And trust me: you won't.

You can't reproduce what you've lost. What if you need access to a file as it existed 1, 10, 100, or 1000 days ago?

Ben Bond-Lamberty

WHY Reproducible Science?



Carole Goble

<http://www.slideshare.net/carolegoble/ismb2013-keynotecleangoble>



Noam Ross

@noamross



Follow



Gelman: "Reproducible research is even better when you're wrong" [#stancon2017](#)

- Fundamental pillar of **scientific method**

Reproducible Science: WHY

- Fundamental pillar of **scientific method**
- Much less prone to **errors**

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- **Code reuse** & sharing accelerates scientific progress
- Increasingly required by **journals**
- Higher publication **impact** (citations, future collaborations, etc)

HOW TO DO Reproducible Science?

1. File **organisation**.
2. **Data management**. Spreadsheet good practices.
3. **Code-based** data analysis. **Rmarkdown**
4. Software **dependencies**.
5. **Version control** & collaborative writing.

- All files in **same directory** (Rstudio project).

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- Figures, code, etc also have their own folder.

File organisation example

myproject

```
| - README           # general info about the project  
  
| - analysis.R       # master script that executes everything  
  
| - data-raw/        # original raw data  
  
| - data/            # clean data (produced w/ script)  
  
| - R/               # functions definitions  
  
| - doc/             # manuscript files  
  
| - figs/            # final figures
```

Data management

Editorial expression of concern

IN THE 3 June issue, *Science* published the Report “Environmentally relevant concentrations of microplastic particles influence larval fish ecology” by Oona M. Lönnstedt and Peter Eklöv (*1*). The authors have notified *Science* of the theft of the computer on which the raw data for the paper were stored. These data were not backed up on any other device nor deposited in an appropriate repository. *Science* is publishing this Editorial Expression of Concern to alert our readers to the fact that no further data can be made available, beyond those already presented in the paper and its supplement, to enable readers to understand, assess, reproduce, or extend the conclusions of the paper.

Jeremy Berg

Editor in Chief

<http://science.sciencemag.org/content/354/6317/1242.1>

Use the **cloud**: safe, persistent, easy to share

- Dropbox
- OSF
- Figshare, etc
- See all data repositories in www.re3data.org

Tidy data

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	2666	20065360
Brazil	1999	37737	17206362
Brazil	2000	80488	17404898
China	1999	212258	127205272
China	2000	213766	128053583

variables

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	2666	20065360
Brazil	1999	37737	17206362
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China	1999	212258	127205272
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observations

country	year	cases	population
Afghanistan	1999	745	19987071
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table4

<http://r4ds.had.co.nz/tidy.html>

Spreadsheet good practices

- Put **variables** in **columns** (things you are measuring: height, weight, sex)

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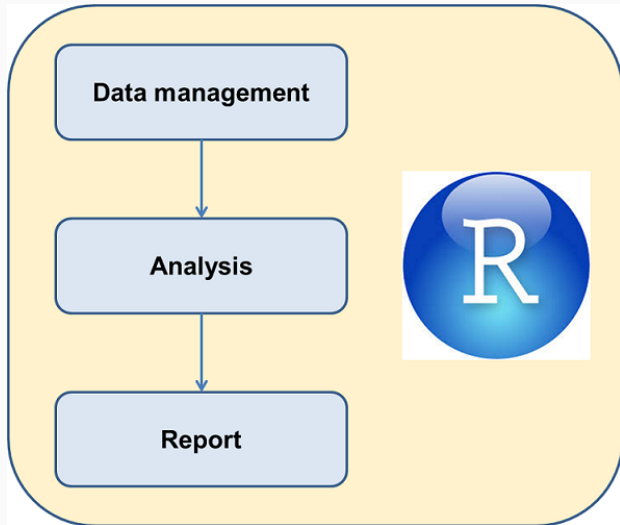
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- <http://kbroman.org/dataorg/>

Data analysis

Always use code

- Reproducible
- Reusable

Dynamic reports



Rmarkdown documents

- Fully reproducible (trace all results inc. tables and plots)
- Dynamic (regenerate with 1 click)
- Suitable for
 - documents (Word, PDF, etc)
 - presentations
 - books
 - websites
 - ...

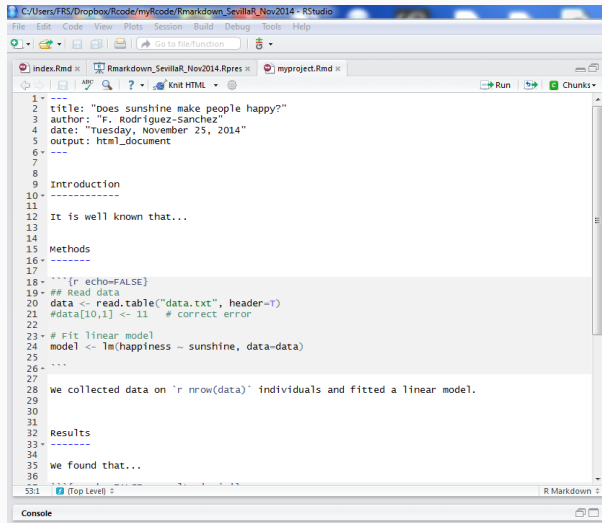


Let's see Rmarkdown in action

In Rstudio, create new Rmarkdown document and click on `Knit HTML`.

Example: Does sunshine influence happiness?

See `myproject.Rmd` (<http://bit.ly/rmdsun>)



```
1 ---
2 title: "Does sunshine make people happy?"
3 author: "F. Rodríguez-Sánchez"
4 date: "Tuesday, November 25, 2014"
5 output: html_document
6 ---
7
8
9 Introduction
10 ---
11
12 It is well known that...
13
14
15 Methods
16 ---
17
18 ```{r echo=FALSE}
19 ## Read data
20 data <- read.table("data.txt", header=T)
21 #data[10,1] <- 11 # correct error
22
23 # Fit linear model
24 model <- lm(happiness ~ sunshine, data=data)
25
26 ```
27
28 we collected data on `r nrow(data)` individuals and fitted a linear model.
29
30
31
32 Results
33 ---
34
35 we found that...
36
37
38
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53:1 (Top Level) R Markdown
```

HTML output includes text, plot and formatted table

Does sunshine make people happy?

F. Rodriguez-Sanchez

Tuesday, November 25, 2014

Introduction

It is well known that individual well-being can be influenced by climatic conditions. However, ...

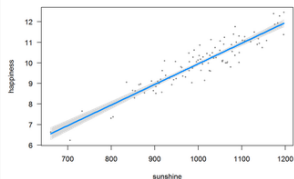
Methods

We collected data on 100 individuals and fitted a linear model.

Results

We found that...

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.0651657	0.4264970	-0.1527928	0.8788758
sunshine	0.0100228	0.0004232	23.6833264	0.0000000



Discussion

These results confirm that sunshine is good for happiness (slope = 0.0100228).

Acknowledgements

Y. Xie, J. MacFarlane, Rstudio...

Spotted error in the data? No problem!

Make changes in Rmarkdown document, click `knit` and report will **update automatically!**

Other formats: PDF, Word

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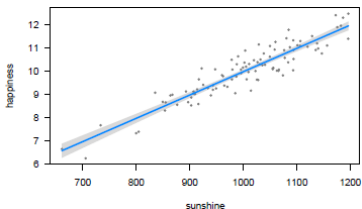
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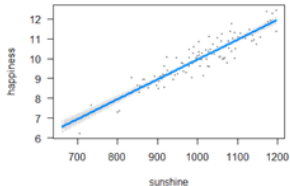
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Adding citations by DOI

rcrossref addin

Add Crossref Citations

Cancel Add Crossref Citations Done

Add a new bibliography entry through Crossref DOI

10.3390/ma8063101 Add to My Citations

Type: journal-article
Title: Photoluminescent ZnO Nanoparticles and Their Biological Applications
Author: Zheng-Yong Zhang; Huan-Ming Xiong
Time: 2015
Publisher: MDPI AG

Adding citations from BibTeX file

citr addin

<https://github.com/crsh/citr/>

- `rticles`
- `rmdTemplates`

Can write full thesis in Rmarkdown!

See `thesis.Rmd`.

See `thesis.pdf`.

Rmarkdown website

<http://rmarkdown.rstudio.com/index.html>

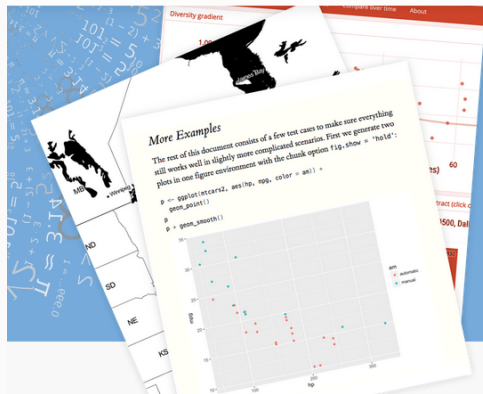
R Markdown

from  Studio

Get Started

Gallery

For



Analyze. Share. Reproduce.

Your data tells a story. Tell it with R.

Turn your analyses into high quality

reports, presentations and dashboards.

Rmarkdown reference guide



R Markdown Reference Guide

Learn more about R Markdown at rmarkdown.rstudio.com
Learn more about Interactive Docs at shiny.rstudio.com/articles

Contents:

1. Markdown Syntax
2. Knitr chunk options
3. Pandoc options

Syntax	Becomes
<p>Make a code chunk with three back ticks followed by an <code>r</code> in braces. The chunk with three back ticks:</p> <pre>```(r) paste("Hello", "World!")</pre>	<p>Make a code chunk with three back ticks followed by an <code>r</code> in braces. End the chunk with three back ticks:</p> <pre>paste("Hello", "World!") ## [1] "Hello World!"</pre>
<p>Place code inline with a single back ticks. The first back tick must be followed by an <code>r</code>, like this <code>`r paste("Hello", "World!")`</code>.</p>	<p>Place code inline with a single back ticks. The first back tick must be followed by an <code>R</code>, like this <code>Hello World!</code>.</p>
<p>Add chunk options within braces. For example, <code>echo=FALSE</code> will prevent source code from being displayed:</p> <pre>```(r eval=TRUE, echo=FALSE) paste("Hello", "World!")</pre>	<p>Add chunk options within braces. For example, <code>echo=FALSE</code> will prevent source code from being displayed:</p> <pre>## [1] "Hello World!"</pre>

Learn more about chunk options at <http://yihui.name/knitr/options>

Chunk options		
option	default value	description
Code options		
<code>child</code>	NULL	A character vector of filenames. Knitr will knit the files and place them into the main document.
<code>code</code>	NULL	Set to R code. Knitr will replace the code in the chunk with the code in the code options.
<code>engine</code>	"R"	Knitr will evaluate the chunk in the named language, e.g. <code>engine = "python"</code> . Run <code>names(knitr::knit_engines\$get())</code> to see supported languages.
<code>eval</code>	TRUE	If FALSE, knitr will not run the code in the code chunk.
<code>include</code>	TRUE	If FALSE, knitr will run the chunk but not include the chunk in the final document.
<code>port</code>	TRUE	If FALSE, knitr will not include the chunk when running <code>port()</code> to extract the source code.
Results		
<code>collapse</code>	FALSE	If TRUE, knitr will collapse all the source and output blocks created by the chunk into a single block.
<code>echo</code>	TRUE	If FALSE, knitr will not display the code in the code chunk above it's results in the final document.
<code>results</code>	"markup"	If "hide", knitr will not display the code's results in the final document. If "hold", knitr will delay displaying all output pieces until the end of the chunk. If "skip", knitr will pass through results without reformatting them (useful if results return raw HTML, etc.)
<code>error</code>	TRUE	If FALSE, knitr will not display any error messages generated by the code.
<code>message</code>	TRUE	If FALSE, knitr will not display any messages generated by the code.
<code>warning</code>	TRUE	If FALSE, knitr will not display any warning messages generated by the code.
Code formatting		
<code>background</code>	"FFFFFF"	A background color for chunks in LaTeX output.
<code>comment</code>	"#"	A character string. Knitr will append the string to the start of each line of results in the final document.
<code>highlight</code>	TRUE	If TRUE, knitr will highlight the source code in the final output.
<code>prompt</code>	FALSE	If TRUE, knitr will add <code>></code> to the start of each line of code displayed in the final document.
<code>size</code>	"normalsize"	Fontsize for LaTeX output.
<code>strip.white</code>	TRUE	If TRUE, knitr will remove white spaces that appear at the beginning or end of a code chunk.
<code>tidy</code>	FALSE	If TRUE, knitr will tidy code chunks for display with the <code>tidy_source()</code> function in the <code>formatR</code> package.

8 Studio

Updated 10/30/2014

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Managing software dependencies

Managing package dependencies in R

- **sessionInfo** (or `session_info`)
- `switchr`
- `rctrack`
- **checkpoint**
- **packrat**
- `docker`

Version control

"FINAL".doc



FINAL.doc!



FINAL_rev.2.doc



FINAL_rev.6.COMMENTS.doc



FINAL_rev.8.comments5.
CORRECTIONS.doc



FINAL_rev.18.comments7.
corrections9.MORE.30.doc



FINAL_rev.22.comments49.
corrections.10. #@\$%WHYDID
ICOMETOGRADSCHOOL?????.doc

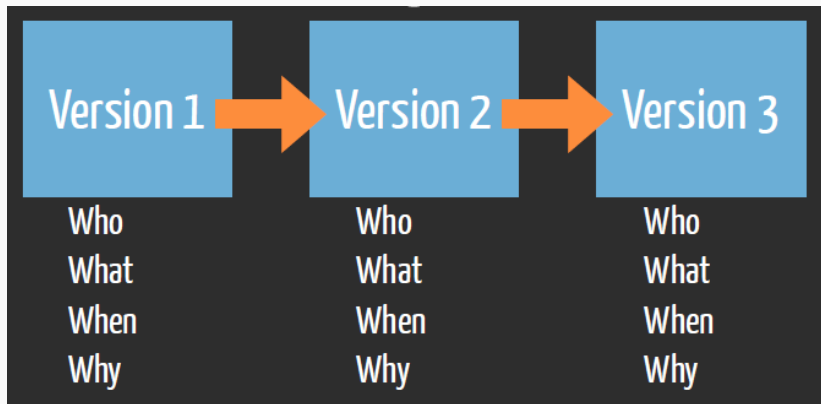
Dropbox keeps record of deleted/edited files for 30 days

Open Science Framework

Automatic version control, no time limit.

The screenshot displays the Open Science Framework (OSF) interface. The top navigation bar includes links for Dashboard, My Projects, Browse, and a search icon. The user profile 'Sara Bowman' is visible. The main content area shows a file named 'manuscript green chemistry.docx'. Below the file name are buttons for Delete, Check out, Download, View, and Revisions. The Revisions button is highlighted with a pink box, and a pink arrow points to it. Below the buttons is a table titled 'Revisions' with columns: Version ID, Date, User, Download, MD5, and SHA2. The Version ID column is highlighted with a pink box. The table lists five revisions, all by Sara Bowman, dated 2016-03-01.

Version ID	Date	User	Download	MD5	SHA2
5	2016-03-01 04:51 PM	Sara Bowman	3	605360a9d897969845e	0a15b7a38d21268e87;
4	2016-03-01 04:51 PM	Sara Bowman	2	d36862941d1f3a9834a	0b26a8c8d5aaa9a26d2
3	2016-03-01 04:50 PM	Sara Bowman	1	4f9731f49aea5b8eafa9	1c86e4964c495201460
2	2016-03-01 04:50 PM	Sara Bowman	1	bc165cff2a8ad6b3a8bc	401cdd53dbcb3c54a45
1	2016-03-01 03:32 PM	Sara Bowman	4	96f5aa2525e176ec2e9;	59ec22c26e9510bac3



R. Fitzjohn

(<https://github.com/richfitz/reproducibility-2014>)



Ecosistemas 25(2): 83-92 [Mayo-Agosto 2016]
Doi: 10.7818/ECOS.2016.25-2.11

Artículo publicado en Open Access bajo los términos
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REVISIONES

ecosistemas

REVISTA CIENTÍFICA DE ECOLOGÍA Y MEDIO AMBIENTE

ISSN 1697-2473 / Open access
disponible en www.revistaecosistemas.net

Ciencia reproducible: qué, por qué, cómo

F. Rodríguez-Sánchez^{1,*}, A.J. Pérez-Luque^{2,**}, I. Bartomeus^{1,**}, S. Varela^{3,**}

[http://www.revistaecosistemas.net/index.php/ecosistemas/article/
viewFile/1178/973](http://www.revistaecosistemas.net/index.php/ecosistemas/article/viewFile/1178/973)

Happy writing!



Slides and source code available at

<https://github.com/Pakillo/ReproducibleScience>