R for Reproducibility

Daniela Palleschi

2024-02-29

Table of contents

Pre	eface	4
	Aims of the book	4 4 5
I	Conceptualisation	6
1	Open Science Practices	7
2	Reproducibility	8
3	Documentation	9
		4.0
II	Basic workflow	10
4	RProjects	11
5	Folder structure	12
6	Writing reproducible code	13
7	Code review	14
Ш	Research workflow	15
8	Pre-registration	16
9	Version control	17
10	Storing results	18
IV	Writing	19
11	Publishing your analyses	20

12	Writing it up	21
13	Collaboration	22
14	Summary	23
V	Exercises	24
15	Exercises	25
Appendices		27
Α	Glossary	27
В	Project setup B.1 CRAN packages	28 28 28

Preface

Under construction

This web-book will serve as a full-length version of the materials for the course 'Open Science Practices: Implementing a Reproducible Workflow in R'. Until it is up-andrunning, the slides can be viewed at the following link: https://daniela-palleschi.github. io/r4repro SoSe2024/

This web-book is the companion to the course 'Open Science Practices: Implementing a Reproducible Workflow in R' (and other iterations) given by Daniela Palleschi at the Humboldt-Universität zu Berlin. The tools discussed in this book are specific to the R environment, but the concepts are universal and programming language agnostic.

Aims of the book

- introduce students to basic concepts of reprodubility and replication in the scope of language research
- teach students how to implement a self-contained project-oriented analysis workflow
- help students develop good habits for project/data management
- teach students how to write up analyses for a paper or thesis using dynamic reports
- introduce students to concepts like version control and containerization

What you will not learn in this book

- how to analyse data
- how to fit models
- how to plot data
- how to design an experiment

About this book

This book was created using a Quarto Book RProject. The glossary for each chapter was created using the glossary package (DeBruine, 2023). The source code for each chapter is viewable by clicking the <> Code button at the top right of the page. The source code for the whole book is also available via GitHub, and is fully reproducible.

Part I Conceptualisation

1 Open Science Practices



▲ Under construction

Coming soon! In the meantime, you can view the slides for this topic here (Open Science) and here (Replciation Crisis).

2 Reproducibility



▲ Under construction

Coming soon! In the meantime, you can view the slides for this topic here (Reproducibility).

3 Documentation



▲ Under construction

Part II Basic workflow

4 RProjects

An ode to self-contained projects



⚠ Under construction

Coming soon!In the meantime, you can view the slides for this topic here (Rprojects).

5 Folder structure



⚠ Under construction

6 Writing reproducible code



△ Under construction

Coming soon!In the meantime, you can view the slides for this topic here (Writing reproducible code) and here (Data wrangling).

7 Code review



⚠ Under construction

Part III Research workflow

8 Pre-registration



⚠ Under construction

9 Version control



▲ Under construction

10 Storing results



▲ Under construction

Part IV Writing

11 Publishing your analyses



⚠ Under construction

12 Writing it up



▲ Under construction

13 Collaboration



▲ Under construction

14 Summary



▲ Under construction

Part V Exercises

15 Exercises



⚠ Under construction

References

DeBruine, L. (2023). Glossary: Glossaries for markdown and quarto documents. https://github.com/debruine/glossary

A Glossary

```
library(tidyverse)
library(glossary)
glossary_path(here::here("glossary/glossary.yml"))
# glossary_persistent(TRUE)

x <-
    readr::read_delim(here::here("glossary/glossary.yml"), col_names = F, delim = "|") |>
    mutate(col = rep(c("term", "definition"), times = 4)) |>
    mutate(X1 = str_replace(X1, ': ', '')) |>
    mutate(X1 = str_trim(X1)) |>
    mutate(col = as.factor(col)) |>
    select(-X2)

glossary_table() |> as_tibble()
```

A tibble: 0 x 0

B Project setup

B.1 CRAN packages

You don't necessarily need to keep track of these, as usually RStudio will automatically inform you of missing packages used in an Rmd or qmd script (e.g., after updating R). But still useful to keep track

```
install.packages("remotes")
install.packages("devtools")
```

B.2 Developer packages

B.2.1 Glossary

Set-up glossary using the glossary package (glossary?). For now I'll use an as-yet-unreleased version, which includes terms that share first words with other terms (e.g., replication and replication crisis).

```
# install.packages("glossary")
remotes::install_github("https://github.com/debruine/glossary")
library(glossary)
```

Add a glossary term.

Set my preferred glossary theme.

glossary_path("glossary/glossary.yml")

```
# append default styles to an external CSS file
write(glossary_style(), "glossary/glossary.css", append = TRUE)
```

B.3 Workflow

B.3.1 renv

Initialise a lockfile and take a snapshot.

Listing B.1 in the Console

```
# initialise
renv::init()

# take snapshot
renv::snapshot()
```

If it's been a while since you're updated your packages, you can update them all with:

```
renv::hydrate()

# or
renv::update()
```

If you've recently updated R, you'll need to re-install your packages. You can simply restore your lockfile.

```
# restore to your most recent package versions
renv::restore()
```

B.3.2 rbbt

Install rbbt. N.B., if you're not in a remote RProject and are using renv, this might return an error. To mitigate this, try running renv::deactivate, and renv::activate() to activate it again.

devtools::install_github("paleolimbot/rbbt")