Computational notebooks for more openness, reproducibility, and productivity in research

A minimal setup to get you started

Dr. Ludmilla Figueiredo
Data & Code Curator (iBID, iDiv)
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Agenda

- Round of introductions
- Motivation
- The kit
- (Break)
- Demo/Try out time
- Q&A

Round of introductions

- Who are you
- What are you working on
- How much data and code do you work with
- Which tools do you use: programming languages, editors, version control

Round of introductions



Dr. Ludmilla FigueiredoiDiv Data & Code Curator



Prof. Juliano Sarmento Cabral, University of Birmingham *The road to integrate climate change projections with regional land-use-biodiversity models*Thursday, 7th September, Hall 1 CD, S 30a

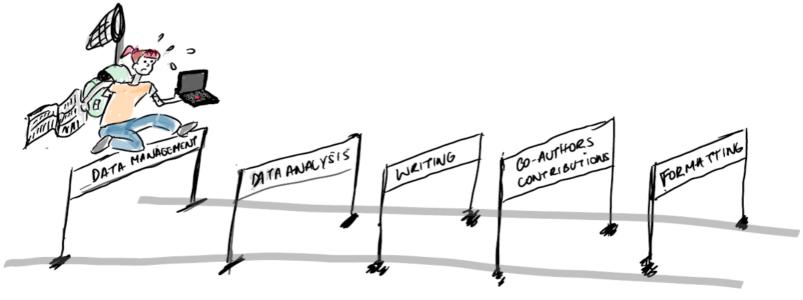




Round of introductions

- Who are you
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Facilitate traceability and reproducibility









Open science







Open science



- = Open data
- = Open access
- = Open source
- = Pre-registration
- = Open peer reviews
- = Open standards
- = Open protocols





Open science



Requirements for FAIR/O:

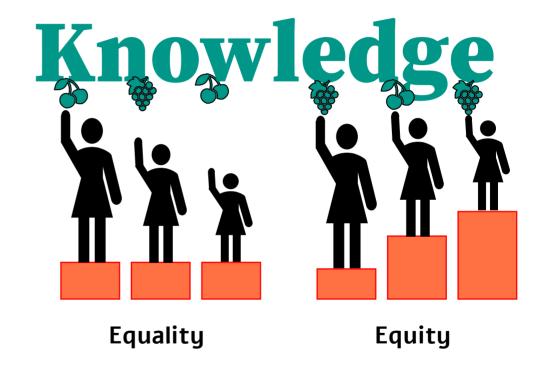
- . Journals
- Institutions
- Funding agencies:
 - . European Research Counsil
 - Horizon Europe
 - . DFG





Knowledge equity

- Done is better than perfect
- Ease of use despite computational background







Facilitate traceability and reproducibility

© Knowledge equity



HOW STANDARDS PROLIFERATE: (SEE: A/C CHARGERS, CHARACTER ENCODINGS, INSTANT MESSAGING, ETC.)

SITUATION: THERE ARE 14 COMPETING STANDARDS.



SOON:

SITUATION:

THERE ARE

15 COMPETING

STANDARDS.

https://xkcd.com/927







PLOS COMPUTATIONAL BIOLOGY

EDUCATION

A simple kit to use computational notebooks for more openness, reproducibility, and productivity in research

Ludmilla Figueiredo 1 ra¤b *, Cédric Scherer 2, Juliano Sarmento Cabral 1,3

1 Ecosystem Modeling, Center for Computational and Theoretical Biology (CCTB), University of Würzburg, Würzburg, Germany, 2 Department of Ecological Dynamics, Leibniz Institute for Zoo and Wildlife Research (IZW), Berlin, Germany, 3 Biodiversity Modelling and Environmental Change, School of Biosciences, College of Life and Environmental Sciences, University of Birmingham, Birmingham, United Kingdom

https://doi.org/10.1371/journal.pcbi.1010356

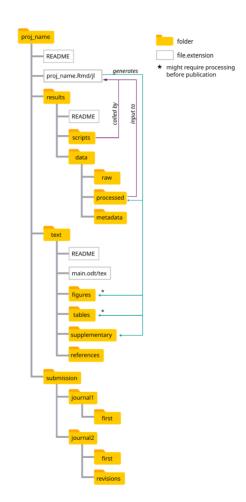


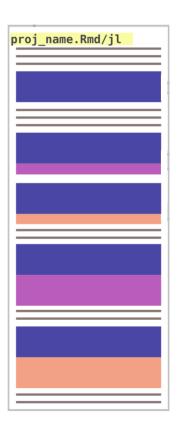


Contents

Direct download

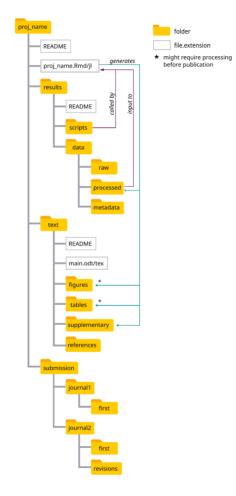
- →set_kit.R
- →set_kit.jl







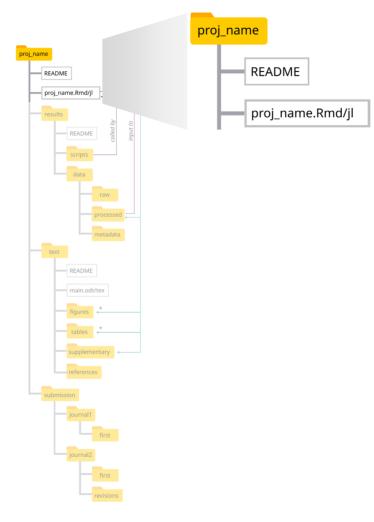




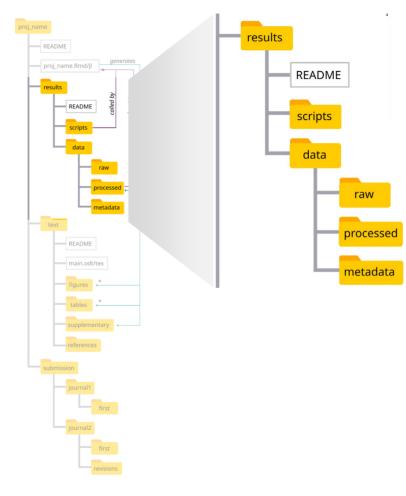






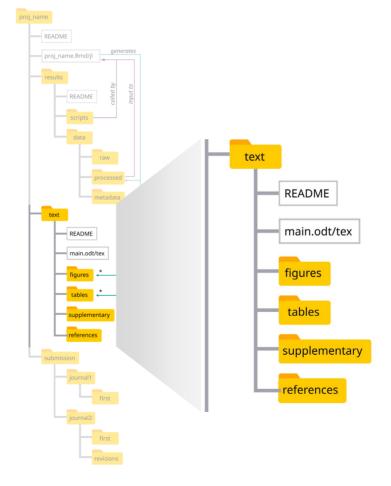








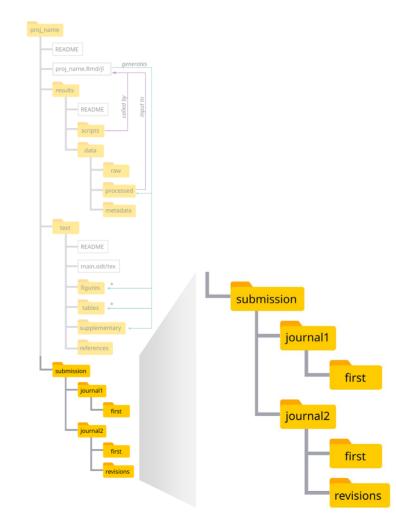




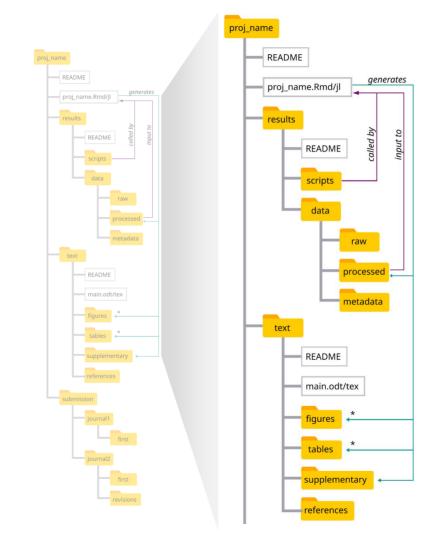




Q&A



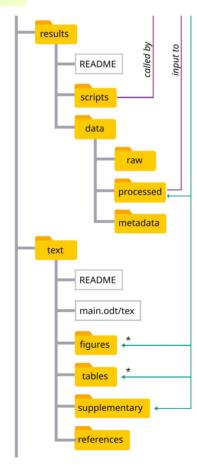




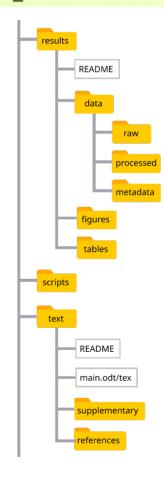


New file structure

main branch:



file_struct2 branch:





Notebook is adapted to handle both

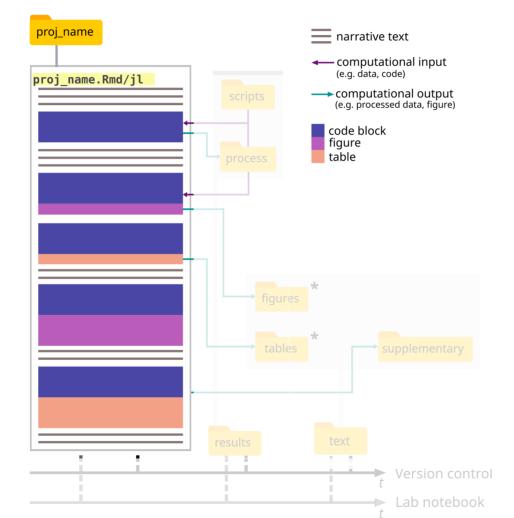
main branch:

```
data_dir <- file.path("results", "data", "processed")
scripts_dir <- file.path("results", "scripts")
suppl_dir <- file.path("results", "supplementary")
figs_dir <- file.path("text", "figures")
tabs_dir <- file.path("text", "tables")</pre>
```

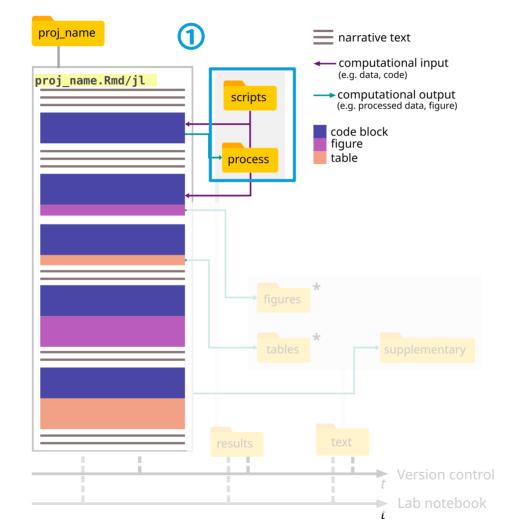
file_struct2 branch:

```
data_dir <- file.path("results", "data", "processed")
scripts_dir <- file.path("scripts")
figs_dir <- file.path("results", "figures")
tabs_dir <- file.path("results", "tables")
suppl_dir <- file.path("text", "supplementary")</pre>
```

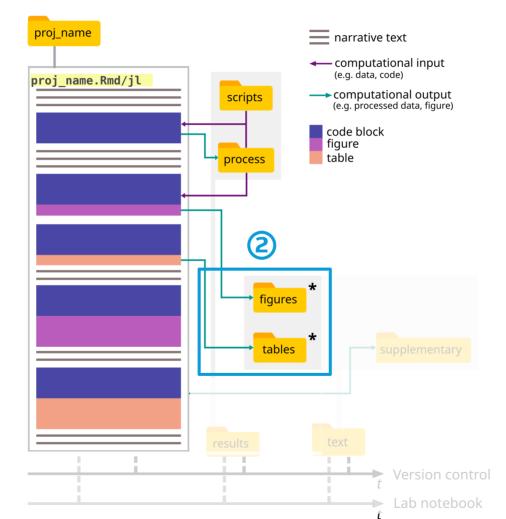




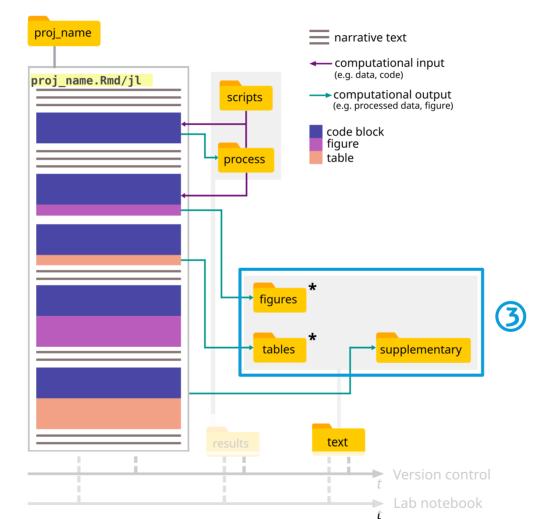




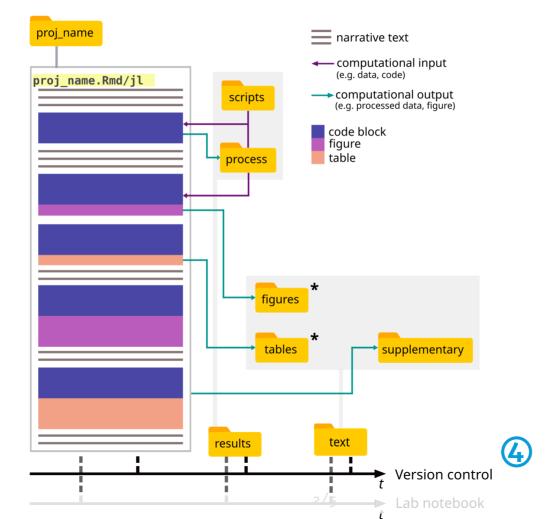




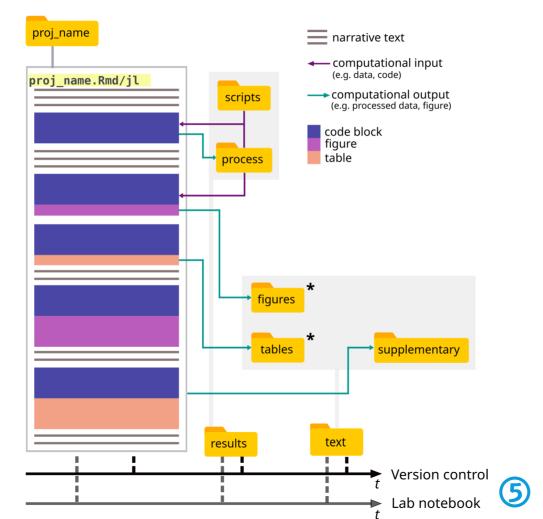














toc: true pdf_document: default 10 <! --11 12 This is your notebook. When filling it, you should obey the 13 Text written directly in the file like this one you are rea this file (click the "Knit" button in the upper-left part 15 <!-- Text written like this will be not be included in html 17 This line includes 'inline code'. Inline code is only forma 18 19 Blocks of code (also called 'chunks') are marked like this: 21 print("This is an example of a code block") 23 24 To execute the code in the chunk and have it's results appe 26 To insert add a chunk, click the 'Insert' button on the too 28 It is also possible to include hyperlinks: [link to further 30 Very important, check how to adjust how much of your code a 31 32 36 # Create objects to provide easy access to the folders rela-# The code in this chunk is not relevant for the reader, and 38 # This is also useful if you are not using the folder struc # Feel free to edit this path as you wish. 41 data_dir <- file.path("results", "processed") ## DO NOT pla scripts_dir <- file.path("results", "scripts") suppl_dir <- file.path("results", "supplementary") semiprods_dir <- file.path("results", "semi_products")</pre> 45 figures_dir <- file.path("text", "figures")
46 tables_dir <- file.path("text", "tables")

RStudio Source Editor

output:

□□ | □ | □ | △□ Q | 6 Knit • □ •

code folding: hide

2 title: "Your title here"

html_document:

test.Rmd ×

48 49 - # Read data

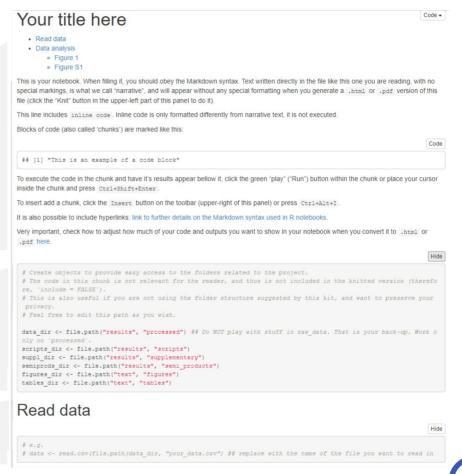
50 + ```{r}

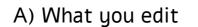
51 # e.g.

Examples of Markdown svntax

Short paths to relevant directories

Suggestion of use





52 # data <- read.csv(file.path(data_dir, "your_data.csv") ##



Live Demo





Try it out!

Kits:

https://github.com/FellowsFreiesWissen/computational_notebooks

http://tinyurl.com/notebooks2023



Data:

https://idata.idiv.de/ddm/Data/ShowData/3519?version=o

http://tinyurl.com/dataECEM2023







Next steps

- Electronic lab notebooks
- > Writing manuscripts in Rmarkdown or Quarto
- R packages to build workflow:
 - workflowr (Blischak et al. 2019)
 - > targets (Landau 2021)
- > Organize your work as an R package or research compendium
- > renv, Binder, Docker image/containers
- > Emacs' Org mode: fully customizable digital lab notebook





Thank You!

Prof. Juliano Sarmento Cabral & Dr. Cedrid Scherer Center for Computational and Theoretical Biology, Uni Wuerzburg

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