# A desperate guide to data analysis workflow improvements

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"Reproducible Data Analysis" Data Science @ BOKU initiative

> 2024-02-29 **BOKU** Wien



THIS IS YOUR MACHINE LEARNING SYSTEM?

YUP! YOU POUR THE DATA INTO THIS BIG PILE OF LINEAR ALGEBRA, THEN COLLECT THE ANSWERS ON THE OTHER SIDE.

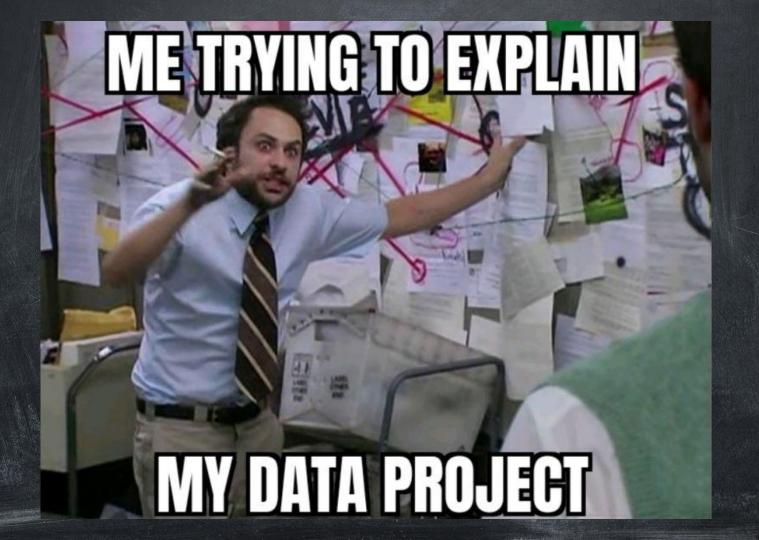
WHAT IF THE ANSWERS ARE WRONG?

JUST STIR THE PILE UNTIL THEY START LOOKING RIGHT.









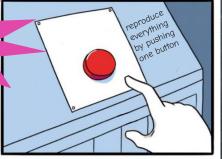
#### Our goal: reproducibility!

...but in practice:

nobody re-runs it or it is broken

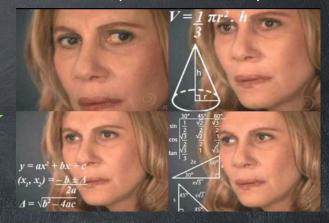
irreproducibility
by hardware
demands

irreproducibility
by obscurity





comprehensibility



traceability



some users might be interested mostly in results

#### Make things more usable - for yourself and others!

Reproducibility in practice: How easily can somebody use the thing later?

Think about your target audience to set priorities!

Do they want to...

- re-run all your scripts?
- run the code as piece of software?
- use the code and adapt it?
- use the computation results only?
- understand what and how we did it?





# 7 Steps towards more use-able data analysis projects

Gamification: count every  $\checkmark$ , if you follow the guideline already!





#### (1) Publish your result data sets 🗸

Wicherts, J. M., Bakker, M., & Molenaar, D. (2011). Willingness to Share Research Data Is Related to the Strength of the Evidence and the Quality of Reporting of Statistical Results. In R. E. Tractenberg (Ed.), PLoS ONE (Vol. 6, Issue 11, p. e26828). Public Library of Science (PLoS). https://doi.org/10.1371/journal.pone.0026828

Willingness to Share Research Data

is related to

Strength of the Evidence

(1) Publish your result data sets: in a fair way!

Findable Accessible Interoperable Reusable

O Physical Reusable

https://www.go-fair.org/fair-principles/https://doi.org/10.1038/sdata.2016.18

by SangyaPundir under CC-BY-SA 4.0

#### (1) Publish your result data sets: how?

- ✓ Use an open license if possible! (Allows others to publish your results as input data!)
- Label and document the fields of your result data set properly
- ✓ A CSV or NetCDF file on Zenodo is already pretty good
- X tables in the PDF of the supplementary material
- non-open or non-standard file formats like \*.gdx (GAMS)
- X units for quantities not documented

More data providers: Dataverse, FigShare, AUSSDA, Dryad, Mendeley Data, DataHub, DANS, EUDat, ...

#### (2) Publish your code 🗸

- use an open license, e.g. MIT license or CC-BY ✓ (unless you need GPL libraries)
- For maintained projects: Use GIT / Github
- Include a README file
  - What does it do? (summary & link to article)
  - Requirements: hardware & software
  - How to run / use the thing

## (3) Use reproducible virtual environments /

Virtual environments (e.g. Conda) allow you to:

- Multiple versions of a library on one machine
- Pretty portable (Windows, Linux, Mac OS)
- Easier to use than Docker

...and to export all used libraries & precise versions!

Alternatives to conda: renv, pipenv, ...

Export all used dependencies using conda: 🗸

conda env export --no-builds | grep -v "^prefix: " > env.yml

Use micromamba - conda is too slow! 🗸

https://mamba.readthedocs.io/en/latest/installation/micromamba-installation.html

#### env.yml:

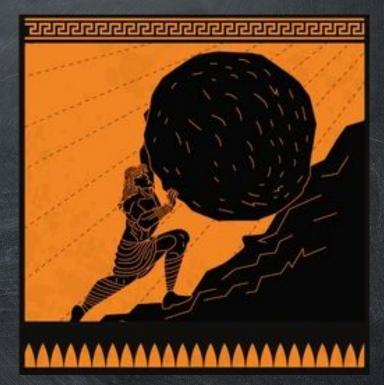
```
name: my project
 - conda-forge
  defaults
  ca-certificates=2023.11.17
  liblapack=3.9.0
  libzlib=1.2.13
  ncurses=6.4
  - numpy=1.26.3
  openssl=3.2.1
  pip=23.3.2
  python=3.10.13
  wheel=0.42.0
  -xz=5.2.6
  zstd=1.5.5
    - cplex==22.1.1.0
```

# (4) Automate everything Manual steps are...

- error prone
- time consuming
- not documented

Every command or click should be stored in a way, such that it can be executed again!

Be realistic! You will do things over and over again...



But you can automate all steps!



#### (4) Automate everything

Goal: a single executable script 🗸

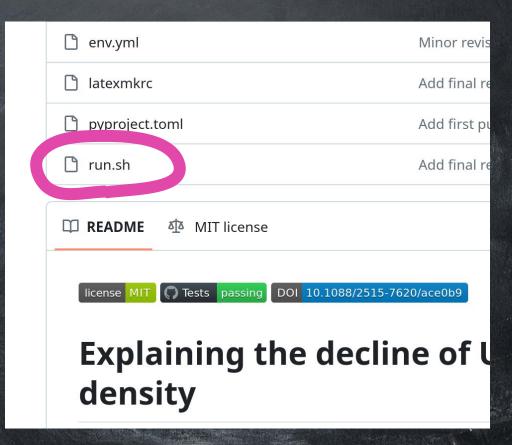
- download of input data
- preprocessing
- computation
- write results to files
- create figures

article

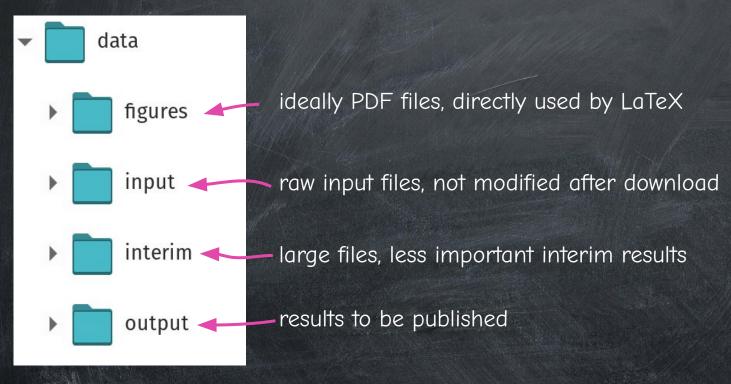
IAT<sub>E</sub>X

how to & links in bonus slides

**+** quarto



## (5) Use a good folder structure 🗸



#### (5) Use a good folder structure

your project

Follow common conventions!

- no spaces in file names
- file names in lower case or ALL\_CAPS
- avoid CamelCase for files and folders

/ auto-create from templates



https://github.com/cookiecutter/cookiecutter

#### Literature:

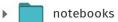
[1] G. Wilson, J. Bryan, K. Cranston, J. Kitzes, L. Nederbragt, and T. K. Teal, "Good enough practices in scientific computing," PLOS Computational Biology, vol. 13, no. 6, p. e1005510, Jun. 2017, doi: 10.1371/journal.pcbi.1005510.

[2] W. S. Noble, "A Quick Guide to Organizing Computational Biology Projects," PLOS Computational Biology, vol. 5, no. 7, p. e1000424, Jul. 2009, doi: 10.1371/journal.pcbi.1000424.

[3] "Cookiecutter Data Science." <a href="https://drivendata.github.io/cookiecutter-data-science/">https://drivendata.github.io/cookiecutter-data-science/</a>

















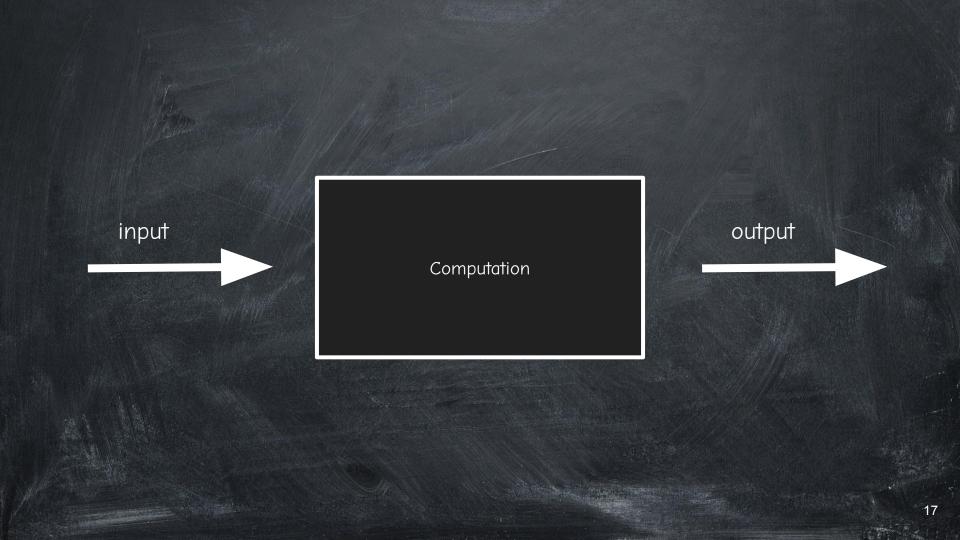


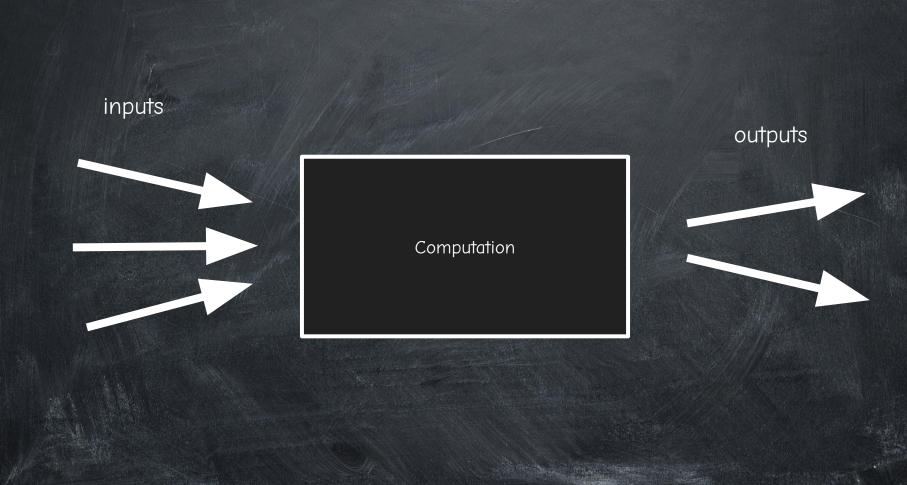


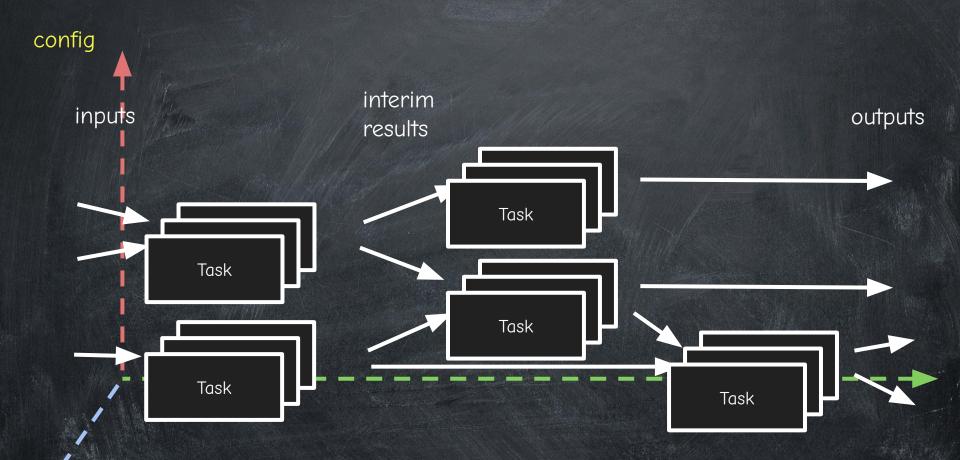




Organize your computation pipeline







parameters

## (6) Use a pipeline tool 🗸

- Caching / incremental builds
  - $\rightarrow$  skip tasks if nothing changed (inputs, code, config, parameters)
- Parametrizing tasks
  - → specify (multi dimensional) parameter spaces via ranges or lists
- Run tasks in parallel
  - $\rightarrow$  on different CPUs or on different nodes on a cluster (e.g. VSC)

Rabix KNIME Analytics Platform Fission Workflows Polynote

Dask

Sushi<sub>stpipe</sub>
OpenMOLE DSL

Reflow

Mario Remake

Pydra Common Workflow Language

Vet Another Workflow Language

Workflow Patterns Initiative

Pydra Common Workflow Language

Redun

Pyperator Zennelin

Redun pyperator zeppelin Flyte Arvados Factual's Drake SciLuigi Joblib Swif Just Snakemake Pinball Spyppol Cadence ecflow Zenaton Nestral Workflow Description Language Substation Substation Orbital Scotland Pipengine Toil Source Action Chain Martian Source Substation Su Ospiff MetaFlow Dataform Bpipe Couler Nessie Stolos Sure DVC Steppy Vali ZenML Yapp StreamPipes Sure DVC Steppy Vali ZenML Yapp StreamPipes Sure DVC Polyaxon OpenMOLE Ray Polyaxon OpenMOLE Ray Prodmodel Popper Maestro Temporal Piper StreamFlow Pachvderm Scoop Make DataLad Dagr GC3pie Flowr Make DataLad Dagr GC3pie Flowr Make Cradle Dray Butler Scoop Rapping Condition Dray Rapp Note Data Popper Prodmodel Pachvderm Scoop Make Cradle Dray Prodmodel Dray Rapp Note DataLad Dagr GC3pie Flowr Make Cradle omegalmi DataOps Platform Rain Fireworks Reana CDS https://github.com/pditommaso/awesome-pipeline aso/awesome-pipeline s Prefect Gradle CULICITOITI

#### (7) Keep old versions of your data 🗸

Re-running computations might be difficult or take long time... Keep old versions!

It would be nice to...

- know how artefacts (figures, output files, ...) were generated
   → record Git hash / code, all inputs, parameters, conda environment, config, ...
- delete large files later when disk space is an issue
- compare between different versions of data

Git alone is not enough to fulfill these requirements!

Note that Git offers more features:

- Sync to other machines / collaboration with others
- Publish the repository but data is not always publishable due to licence issues

Tools for large data: DVC, GIT-LFS, git-annex, DataLad, ...

# My poor-man data versioning run.sh script:

- runs the computation
- creates a Git tag
- copies outputs to an archive folder

#### Open issues:

- How to sync data between machines? Use Unison?
- How to collaborate with others using the same data?
- How to sync figures to Overleaf without Git?

https://github.com/inwe-boku/windpower-decomposition-usa/blob/main/run.sh



#### Cookies for all of you!





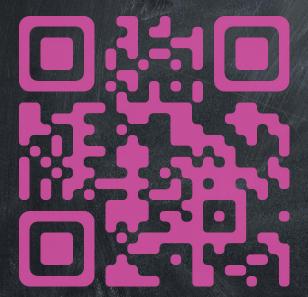
Everything we have seen today in a cookiecutter template:

- folder structure
- code snippet for including results in LaTeX
- Script for building a zip file for arXiv.org
- run.sh script
- Snakemake file
- README.md, .gitignore, LICENSE, ...
- ...

→ https://github.com/inwe-boku/cookiecutter-data-research

#### Reproduce all the things!

Download slides:



https://bit.ly/desperate-guide

peter.regner@boku.ac.at

https://github.com/inwe-boku/

https://refuel.world/



We gratefully acknowledge support from the European Research Council ("reFUEL" ERC-2017-STG 758149).

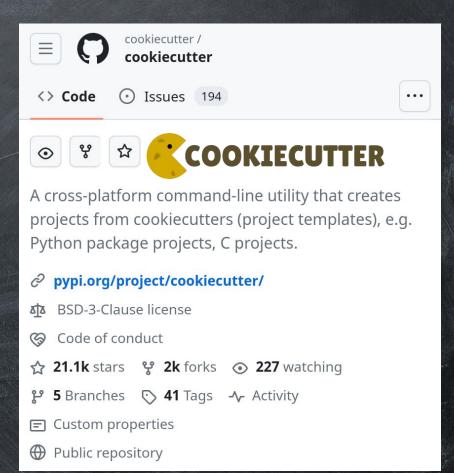


#### Automate project creation <

Create a new folder from a template:

- folder structure
- LICENSE file
- gitignore file
- code snippets
- ..

https://github.com/cookiecutter/cookiecutter



#### Include figures in LaTex documents 🗸

MS Word: use linked images

Latex:

```
\begin{figure}
                                    folder for all figures
    \centering
    \includegraphics{data/figures/capacity_factors.pdf}
    \label{fig:capacity_factors}
\end{figure}
                         no figure number
```

Overleaf: use GIT, Github or dropbox to sync figures - premium only...:-/

https://www.overleaf.com/learn/how-to/Dropbox\_Synchronization https://www.overleaf.com/learn/how-to/Git\_Integration\_and\_GitHub\_Synchronization

#### Include results in LaTex documents <

Turns out to be surprisingly difficult! 😱

```
result_values = {}
meaning_of_life = 42
result_values['meaning_of_life'] = f'{meaning_of_life:d}'
gravity_ms2 = 9.80665
result_values['gravity'] = f'{gravity_ms2:.2f}', 'm/s^2'
write_result_values(result_values)
```

script writes results to a file with custom LaTeX commands

\newcommand{\meaning\_of\_life}{42}
\newcommand{\gravity}{\qty{9.81}{m/s^2}}

auto-generated LaTeX commands

More details and other Solutions like Knitr: <a href="https://tex.stackexchange.com/a/711627/8964">https://tex.stackexchange.com/a/711627/8964</a>

#### Do more good things

- Write good code
  - Don't do no magic numbers
  - No absolute file paths /
  - No unnecessary Abbrev.
  - Follow code conventions (e.g. pep8)
- Write unit / functional tests /
   Automatically check if things are behaving as expected!
- Use a code linter (e.g. as GIT hook) 

  Tools to automatically check code for errors and style violations, for Python: flake8, black (auto-formatter),
- Use Continuous integration 
   Run tests, code linter or computations automatically, e.g. via Github Actions

#### Tracking additional computation information

```
@task
def my_fancy_func(some_param, inputs, outputs):
    # do something here
...
```

The @task decorator creates a file with metaparameters for each output:

```
function: concat_solution_chunks
git_commit: 59f9737-dirty
hostname: nora
input_files:
- data/interim/network_solution/network_solution_740_560.nc
[...]
- data/interim/network_solution/network_solution_795_615.nc
input_params: {}
output_files:
- data/output/network_solution/network_solution.nc
runtime: 7.013682842254639
start_time: '2024-01-10T12:57:21.699562+01:00'
```

#### Link to source:

https://github.com/inwe-boku/cookiecutter-data-research/blob/main/%7B%7B%2Ocookiecutter.repo\_name%20%7D%7D/src/task.py

#### Data Analysis

## Software Development

little maintenance

single use projects

large data

long run-time

few users

plots

few developers

interactive programming

literate P'

maintained for long time

Continuous

Integration

programming

GIT / Github

unit tests

code review

build system

packages

deployment

large user

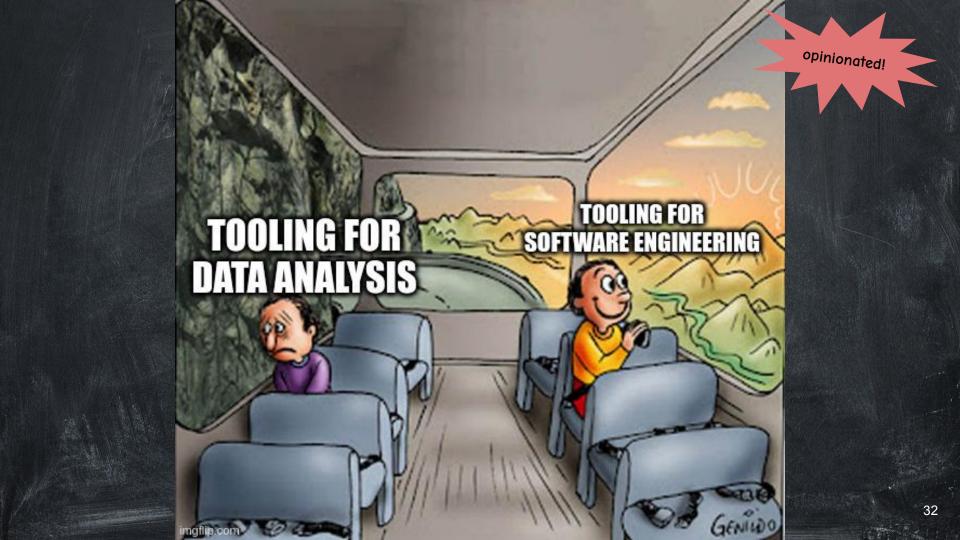
GUI design

base

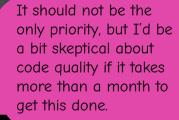
many

developers

debugger



#### Reproducibility is a hot topic...





Texting with a colleague at another uni who is mired in the AEA replication process. Wasting > month getting all in order so that some RA can just push one button and get all the data merges right so that the results are perfect to the 3rd decimal place.

6:05 PM · Feb 17, 2024 · **429K** Views



1/5

**1** 58



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https://twitter.com/toniwhited/status/1758900448596291953