# Computational Reproducibility (Research Reproducibility in Theory and Practice, Day 3, FSCI2021)

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Slides and examples are in <a href="https://bit.ly/CompRepro">https://bit.ly/CompRepro</a>



NCSA | National Center for Supercomputing Applications



#### **Exercise**

- Take a look at this dataset: <a href="https://osf.io/z274d/">https://osf.io/z274d/</a>
- Download it via: <a href="https://osf.io/z274d/download">https://osf.io/z274d/download</a>
- Contains demographic data: tab-separated with a header row
- Using this data, create a graph that shows life expectancy in Canada between 1980 and 2000
- Write down how you did it, and give it to someone else, then ask them to reproduce it

```
year poplifeexp gdppercap country continent
1952 8425333 28.801 779.4453145 afghanistan asia
1957 9240934 30.332 820.8530296 afghanistan asia
...
```



#### Goals

- Reproducibility
- Of what?
  - Papers, results, figures
- By whom?
  - Future you, someone else knowledgeable in your field, anyone else
- When?
  - Tomorrow, six months, 5 years, 50 years
- How much?
  - Close enough (you decide what this means), not necessarily all the bits
  - Plausible vs. practical



### **Defining R\* - terms**

- Reproducibility, Replicability, Repeatability, etc.
- Confusing terms see "Replicability vs. reproducibility or is it the other way around?" (blog) and
   "Reproducibility vs. Replicability: A Brief History of a Confused Terminology" (paper) for some discussion
- Maybe these are getting to be more standardized? But still, define what you mean!

Goodman	Claerbout	ACM (2020+)	AMC (2020-)
		Repeatability	Repeatability
Methods Reproducibility	Reproducibility	Reproducibility	Replicability
Results Reproducibility	Replicability	Replicability	Reproducibility
Inferential Reproducibility			

Goodman, S. N., Fanelli, D., and Ioannidis, J. P. A. (2016). <u>What does research reproducibility mean?</u> *Sci. Transl. Med.*8:341ps12. Claerbout, J. F., and Karrenbach, M. (1992). <u>Electronic documents give reproducible research a new meaning</u>. *SEG Expanded Abstracts* 11, 601–604. Association of Computing Machinery (ACM) (2020). <u>Artifact Review and Badging (Version 1.1)</u>.



#### **Context:** data science

- Organize and analyze large (or small) data sets to learn from them
  - Steps: capture/acquire, organize, process, analyze, communicate
- Examples
  - How fast are stars moving away from us, and how does this vary with their distance?
  - Which credit card transactions are fraudulent?
  - What does this German document say in English? What does this recording of someone speaking Spanish say?
  - Which patient scans contain tumors?
  - Who's going to win the election?
  - If a patient has these symptoms, what disease do they have?
  - What treatment is best for this particular patient?
- Relevant: statistics, preregistration (declare your hypothesis before doing your analysis), random studies, false positives/negatives, sample size, confidence, power
- Typical outputs: data, tools and methods (algorithms, models, software), conclusions (understanding data)



#### Context: computational science

- Modeling or simulating a (physical) process in a predictive way, often using one or more equations
- Examples, simulation or analysis of:
  - Atmospheric or oceanic circulation, coupled together with other physical processes into a climate simulation
  - The interactions of atoms in one or more molecules (drug design)
  - The atoms and forces in a material (material design)
  - Engineering analysis of the stress or deformation of a structure under some load (mechanical engineering)
  - Electrical signals in a circuit board or a set of synapses (electrical engineering or neuroscience)
  - Microwaves focused on a breast tumor (patient-specific medicine)
- Often called computational science & engineering (CSE)
- Relevant: mathematics, error bounds
- Typical outputs: algorithm, method, software, conclusions (understanding processes)



#### Computational reproducibility principles

- 1. Provide structure
- 2. Control the source & changes
- 3. Use notebooks to explain and document
- 4. Automate steps
- 5. Automate everything
- 6. Capture the environment
- 7. Provide a license & make citable



### First thing – get a terminal

- On a Mac
  - Click the Launchpad icon in the Dock, type Terminal in the search field, then click Terminal.
  - In the Finder , open the /Applications/Utilities folder, then double-click Terminal.
- On Windows
  - Open your computer's Start menu. Click the Windows ☐ icon on the bottom-left corner of your desktop or press the ⊞ Win key on your keyboard
  - Type cmd or Command Prompt. After opening the Start menu, type this on your keyboard to search the menu items. Command Prompt will show up as the top result.
  - Click the Command Prompt app on the menu. This will open the Command Prompt terminal in a new window.
- Using Binder
  - Go to <a href="https://github.com/danielskatz/repro-fdtd1d">https://github.com/danielskatz/repro-fdtd1d</a>, click on <a href="https://github.com/danielskatz/repro-fdtd1d">https://github.com/danielskatz/repro-fdtd1d</a>, click on</a>
  - Once binder starts the repo, use "New" -> "Terminal" to get a terminal



#### Principle 1 – Provide structure

- Use directories for different things, all inside a project directory, with a top-level readme and license
  - E.g., data, docs, models, notebooks, references, reports, src (for Python data science, <u>Cookiecutter Data Science</u> is an example)
- Use relative paths, so that you can move and share
  - (../data/file.dat)
- Use names that have meaning (and avoid using "final")
  - 00-dsk-data\_acquisition.py

```
My_project
```

- --data
- --docs
- --notebooks
- --references
- --reports
- --src



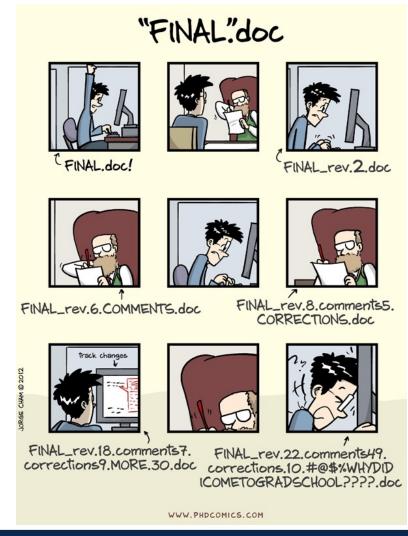
## Principle 2 – Control the source & changes

- 1. For data, store the original (raw) data archivally somewhere and build other versions elsewhere using scripts (including accessing the data from the archive)
  - Note: GitHub is not archival, and isn't good for large datasets
    - Two previous versions of this class used data in <a href="https://raw.githubusercontent.com/csoderberg/test\_study/master/gapminder\_copy.txt">https://raw.githubusercontent.com/csoderberg/test\_study/master/gapminder\_copy.txt</a> but this no longer exists
  - However, it is still in OSF: <a href="https://osf.io/z274d/">https://osf.io/z274d/</a>
  - Get it via
     wget https://osf.io/z274d/download -O gapminder\_copy.txt
     (You may have to install wget Google it)

```
My_project
  --data
         --raw
        |--derived
|--results
  --docs
  --notebooks
  --references
  --reports
  --src
```

## Principle 2 – Control the source & changes

- For software, use a version control system to save versions and changes, and explain the reason for the changes
  - Git is the standard these days
  - Basics
    - Software is stored somewhere (e.g., GitHub, GitLab), either privately or publicly
    - New versions can be added
      - Author, changes, message about change stored
    - Multiple people can make changes in different parts of a project or even a file, and these can be merged together, mostly automatically
  - See <u>Software Carpentry's "Version Control with Git"</u>
  - Version numbers
    - Consider releases, use semantic versioning
      - A release is a tagged version
      - major.minor.patch (API-breaking.API-maintaining.bug-fixes)



### Principle 2 – Control the source & changes

- 3. For published documents, people, etc. find a permanent identifier (PID, e.g., DOI, PubMed ID, ORCID) and use it to find the details (e.g., for references)
  - Get data from ORCID for a person (in Python):

Get metadata about a paper from a DOI (in bash):

```
curl https://api.crossref.org/works/10.1145/3307681.3325400/transform/application/vnd.crossref.unixsd+xml
```

Get bibtex for a paper from a DOI (in bash):

```
curl -LH "Accept: application/x-bibtex" https://doi.org/10.1145/3307681.3325400
curl https://data.crosscite.org/application/x-bibtex/10.1145/3307681.3325400
```



#### Principle 3 – Use notebooks to explain & document

- Notebooks are great for showing what code does
- And teaching people how to use it
- Intersperse cells with text, equations, runnable code, outputs, images
  - Demo: go to https://github.com/danielskatz/repro-fdtd1d, click on launch binder
  - Once binder starts the repo, click on Notebook Demonstration.ipynb
- This uses binder (mybinder.org) you can too
- Turn a Git repo into a collection of interactive notebooks, making your code immediately reproducible by anyone, anywhere
  - Use requirements.txt to tell binder what dependencies to install in the environment
  - Also take a look at binderhub and jupyterhub if you want to run your own instance
- But don't write code to do the same task in multiple notebooks
  - Pull it out (refactor it) into a (reusable) package, then import that package in the notebooks

#### **Jupyter Notebook Example**

Credit: This is slightly modified from examples used in the FSCI 2 (<a href="https://osf.io/sbnz7/">https://osf.io/sbnz7/</a>), which was created by Courtney Soderberg

#### **Setting up the notebook**

#### Lets get started

The notebook is built up from separate editable areas, or cells.

A new notebook contains a single code cell.

Add a line of code and execute it by:

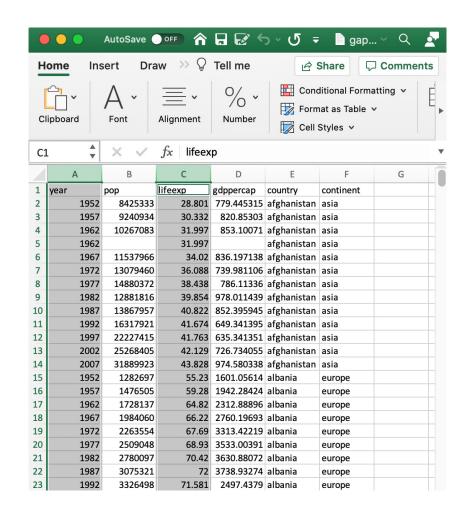
- · clicking the run button, or
- click in the cell, and press shift-return

```
In [1]: print('hello world')
```

hello world

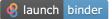


- 1. Anything you do by hand is subject to irreproducible errors
  - GUIs can be intuitive, but they don't support scalability or reproducibility well
    - Imagine having to extracts a column of data from 1000 Excel files
  - Goal: capture what you do in some way so that you can repeat it in one step, such as
    - Capture a set of commands in a single script
    - Use <u>pyexcel</u> to read and write data to/from Excel files
    - Script GUI actions see <u>AutoHotkey\_L</u> for an example of how this can be done for Windows programs



- 2. Scripts for simple things (steps)
  - Shell scripts
    - See Software Carpentry lesson "The Unix Shell"
      - "The Unix shell has been around longer than most of its users have been alive. It has survived so long because it's a power tool that allows people to do complex things with just a few keystrokes. More importantly, it helps them combine existing programs in new ways and automate repetitive tasks so they aren't typing the same things over and over again."
    - At the simplest, the shell is the process you interact with when you type in a terminal window
    - Multiple commands can be placed in a script and rerun
    - And the shell supports variables and control flow (e.g. if-then, loops)

Go to <a href="https://github.com/danielskatz/repro-fdtd1d">https://github.com/danielskatz/repro-fdtd1d</a>, click on



Once binder starts the repo, use "New" -> "Terminal" to get a terminal

```
mkdir raw mkdir proc Make directories
```

Get raw input files (into raw directory)

```
wget https://raw.githubusercontent.com/danielskatz/parsl-example/master/data/0001.jpg -0 raw/0001.jpg wget https://raw.githubusercontent.com/danielskatz/parsl-example/master/data/0002.jpg -0 raw/0002.jpg wget https://raw.githubusercontent.com/danielskatz/parsl-example/master/data/0003.jpg -0 raw/0003.jpg wget https://raw.githubusercontent.com/danielskatz/parsl-example/master/data/0004.jpg -0 raw/0004.jpg python3 bin/sharpen_image.py raw/0001.jpg proc/0001_sharp.jpg
```

```
python3 bin/sharpen_image.py raw/0001.jpg proc/0001_sharp.jpg
python3 bin/sharpen_image.py raw/0003.jpg proc/0003_sharp.jpg
python3 bin/sharpen image.py raw/0004.jpg proc/0004 sharp.jpg
```

Process raw input files

python3 bin/local\_build\_mosaic.py 2 proc/mosaic.jpg proc/0001\_sharp.jpg proc/0002\_sharp.jpg proc/0003\_sharp.jpg
proc/0004\_sharp.jpg

Further process processed files



- Go to <a href="https://github.com/danielskatz/repro-fdtd1d">https://github.com/danielskatz/repro-fdtd1d</a>, click on <a href="https://github.com/danielskatz/repro-fdtd1d">https://github.com/danielskatz/repro-fdtd1d</a>, click on
- Once binder starts the repo, use "New" -> "Terminal" to get a terminal

Automate by:

sh script/build\_mosaic.sh

Contains all the commands from the previous slide

- 3. Can use notebooks like scripts/programs with tools such as
  - <u>nbclient</u>, a very lightweight python API for executing notebooks
  - Papermill, a tool for parameterizing and executing Jupyter Notebooks
  - <u>Jupytext</u>, a converter between notebooks and code and vice versa
- 4. An interesting-looking new project
  - <u>nbmake-action</u> A Notebook-First Continuous Integration Framework
    - A GitHub Action for testing notebooks, runs them from top-to-bottom
    - Intended to raise the quality of scientific material through better automation
    - For scientists/developers who have written docs in notebooks and want to CI test them after every commit

#### 4. Make randomness repeatable

- Many simulations and data analysis involve random seeds, used to start generating a series of "random" numbers
  - Capture these seeds as part of your step so that you can repeat the same "randomness"
  - And get the "same" results
- But be aware of tradeoffs
  - Example
    - When adding a list of floating point numbers, order can matter due to numerical roundoff
    - When using parallel computing, order can change with the same or different numbers of processes
    - Can force order at the cost of performance (extra sync/lock/messages)
    - Better to know what accuracy counts
    - Or to have a debug mode and a production mode





## Principle 5 – Automate everything

- Make or something make-like to handle multiple steps (dependencies) and not redo what isn't needed
  - In the previous example, what happens if we just change the final program?
  - We don't really want to have to rerun the whole script
  - Learn about make (GNU make, gmake) from <u>Software Carpentry's lesson</u>
  - Short version
    - A program that defines rules for how to make one thing from others (dependencies)
    - Can use variables to make rules general
    - Make knows how to only make a thing when its dependencies have changes
- Other options: workflow (management) systems & languages, e.g., in bioinformatics, snakemake, cwl, wdl, nextflow, ... (there's <u>a CWL wiki page</u> with 298 examples)
- Consider continuous integration to automatically rebuild/test when things change
  - Integrate with GitHub Actions, CircleCl, Travis Cl, etc.



**GNU Make** 





## Principle 5 – Automate everything

```
.PHONY: clean all
all: proc/mosaic.jpg
                                                                             This is in script/Makefile-explicit
clean:
        -rm -rf raw proc
raw:
                                                                     To run it, from the terminal in binderhub, use:
       mkdir raw
                                                                             make -f script/Makefile-explicit
proc:
       mkdir proc
raw/0001.jpg: | raw
       wget https://raw.githubusercontent.com/danielskatz/parsl-example/master/data/0001.jpg -O raw/0001.jpg
[...]
raw/0004.jpg: | raw
       wget https://raw.githubusercontent.com/danielskatz/parsl-example/master/data/0004.jpg -O raw/0004.jpg
proc/0001 sharp.jpg: raw/0001.jpg bin/sharpen image.py | proc
        python3 bin/sharpen_image.py raw/0001.jpg proc/0001_sharp.jpg
[...]
proc/0004 sharp.jpg: raw/0004.jpg bin/sharpen image.py | proc
       pvthon3 bin/sharpen image.py raw/0004.jpg proc/0004_sharp.jpg
proc/mosaic.jpg: bin/local_build_mosaic.py_proc/0001_sharp.jpg_proc/0002_sharp.jpg_proc/0003_sharp.jpg_proc/0004_sharp.jpg
       python3 bin/local build mosaic.py 2 proc/mosaic.jpg proc/0001 sharp.jpg proc/0002 sharp.jpg proc/0003 sharp.jpg proc/0004 sharp.jpg
```



### Principle 5 – Automate everything

```
LANGUAGE=python3
FILE NOS=0001 0002 0003 0004
RAW_FILES=$(FILE_NOS:%=raw/%.jpg)
PROC_FILES=$(FILE_NOS:%=proc/%_sharp.jpg)
SHARPEN = bin/sharpen image.py
MOSAIC = bin/local build mosaic.py
RAW SOURCE DIR=https://raw.githubusercontent.com/danielskatz/parsl-example/master/data
.PHONY: clean all
all: proc/mosaic.jpg
clean:
      -rm -rf raw proc
raw:
      mkdir raw
proc:
      mkdir proc
$(RAW FILES): | raw
      wget $(@:raw/%.jpg=$(RAW_SOURCE_DIR)/%.jpg) -O $@
proc/%_sharp.jpg: raw/%.jpg $(SHARPEN) | proc
      $(LANGUAGE) $(SHARPEN) $(@:proc/%_sharp.jpg=raw/%.jpg) $@
proc/mosaic.jpg: $(MOSAIC) $(PROC FILES)
      $(LANGUAGE) $(MOSAIC) 2 $@ $(PROC FILES)
```

And make the automation as general as possible

This is in script/Makefile

To run it, from the terminal in binderhub, use: make -f script/Makefile



#### **Principle 6 – Capture the environment**

- Containers
  - In Python, use virtualenv (and `pip freeze > requirements.txt` to capture current state) or docker
  - In R, use add\_dependencies\_to\_description() or use <u>renv package</u> or <u>rocker</u>
- VMs (heavier weight than containers, includes OS)
- Reproducible builds a set of software development practices that create an independently-verifiable path from source to binary code
  - Reliant on package identification and management, e.g., <u>Guix</u>, <u>PyPI</u>, <u>CRAN</u>, ...
- Lots of tools and systems see "<u>Publishing computational research a review of infrastructures for reproducible and transparent scholarly communication</u>" for a 2020 survey of 11

- Copyright defines ownership, license gives permission to do something
- But facts aren't copyrightable, while works of authorship are (at least in the US)
  - A particular arrangement of facts might be eligible for copyright protection if that arrangement demonstrates sufficient creativity, but not if the arrangement is something uncreative like chronological or alphabetical order
  - Even with creative arrangement, underlying facts cannot be copyrighted; it's perfectly legal for someone else to pull them out, rearrange them, and use them in something new
  - See "Who 'owns' your data?"
- If you are employed, your employer may own the copyright to things you create at work, and maybe even outside
  - Common in the US and in universities, but students own work they develop, even in their own coursework
- Use a common license, don't create your own
  - Common licenses are understood, uncommon one will prevent people from using your work just because they may not understand the license



- Creative Common licenses for text and data
  - CC0 waive copyright, dedicate to the public domain (not really a license)
  - CC BY (Attribution): material is free to use and adapt, but credit must be given
  - CC BY-SA (Attribution-ShareAlike): free to use and adapt, but credit must be given and adapted material must also be
    distributed with this same license
  - CC BY-ND (Attribution-NoDerivs): free to use, but credit must be given and can't be adapted
  - CC BY-NC (Attribution-NonCommercial): free to use and adapt but credit must be given and can't be used commercially
  - CC BY-NC-SA (Attribution-NonCommercial-ShareAlike): free to use and adapt, but credit must be given, can't be used commercially, and adapted material must also be distributed with this same license
  - CC BY-NC-ND (Attribution-NonCommercial-NoDerivs): free to use, but credit must be given, can't be used commercially, and can't be adapted
- Creative Commons provides a guide/decision tree
- Be aware someone might argue that the data are facts and not subject to copyright, so the license doesn't hold
- Scholarly norms and principles of attribution/credit/provenance/authority might hold more sway
- (for more, see "CC BY and data: Not always a good fit")



- Open Source Initiative licenses for software
  - Don't use a CC license for software
  - At high level, two types of licenses
  - Permissive: MIT, Apache, BSD, ...
  - Copyleft ("viral"): GPL, LGPL
- Use <u>choosealicense.com</u> to pick one
- Pick a very common one if possible
- How to apply (MIT):
  - Create a text file (typically named LICENSE or LICENSE.txt) in the root of your source code and copy
    the text of the license into the file. Replace [year] with the current year and [fullname] with the name
    (or names) of the copyright holders.

- Citeable isn't required for reproducibility, but it's a good idea if you want credit
- Make your data citable
  - Deposit it in an archival repository (e.g., Zenodo, OSF, see <u>re3data.org</u> for more) along with metadata,
     receive a DOI, advertise the DOI and metadata (suggested citation)
- Make your software citable
  - Less well-defined practice
  - GitHub is not an archival repository
  - Can follow data practice (can link GitHub repo to Zenodo to automatically deposit new releases guides.github.com/activities/citable-code)
  - Record metadata in the repository (using CodeMeta or citation.cff), some repositories will pick up
  - Also can use Software Heritage ("archive.org for software") to cite archive of GitHub software
  - See <u>cite.research-software.org</u> for more



#### Exercise(s)

- Try out one of the project structure tools, or look at them and try to organize a project you have similarly
  - Python: Cookiecutter Data Science
  - R: ProjectTemplate
- Redo the exercise from the beginning in a more reproducible manner
- Automate a paper you have written
  - Or try to do this for a paper someone else has written (start by finding the data and code, see how far you can get)

### Final thoughts

- "I was inspired more than 15 years ago by John Claerbout [...] He pointed out to me, in a way paraphrased in Buckheit and Donoho (1995): 'an article about computational result is advertising, not scholarship. The actual scholarship is the full software environment, code and data, that produced the result.'" David Donoho (in <a href="https://doi.org/10.1093/biostatistics/kxq028">https://doi.org/10.1093/biostatistics/kxq028</a>)
- "You shouldn't try to do these things all at once; start with one, or part of one. Then in your next project, do that plus another thing." Karl Broman (in <a href="https://kbroman.org/steps2rr/">https://kbroman.org/steps2rr/</a>)
- It's no secret that good analyses are often the result of very scattershot and serendipitous explorations. [...] That being said, once started it is not a process that lends itself to thinking carefully about the structure of your code or project layout, so it's best to start with a clean, logical structure and stick to it throughout. (in <a href="https://drivendata.github.io/cookiecutter-data-science/">https://drivendata.github.io/cookiecutter-data-science/</a>)



#### Resources (1)

- Organizing projects:
  - Python: Cookiecutter Data Science <a href="https://drivendata.github.io/cookiecutter-data-science">https://drivendata.github.io/cookiecutter-data-science</a>
  - R: ProjectTemplate <a href="http://projecttemplate.net/">http://projecttemplate.net/</a>
- Guidelines:
  - Karl Broman's initial steps toward reproducible research (R, explains python too) https://kbroman.org/steps2rr/
- Reproducible papers:
  - PINGA lab's template (computational science, GitHub, Python, LaTeX) <u>https://www.leouieda.com/blog/paper-template.html</u>
  - Manubot (markdown, git, collaboration) <a href="https://manubot.org">https://manubot.org</a>
  - Akhaghi (C/C++, LaTeX) <a href="https://gitlab.com/makhlaghi/reproducible-paper">https://gitlab.com/makhlaghi/reproducible-paper</a>
- Book:
  - The Practice of Reproducible Research: Case Studies and Lessons from the Data-Intensive Sciences http://www.practicereproducibleresearch.org/



#### Resources (2)

- Short courses/MOOCs:
  - Essential skills for reproducible research computing https://barbagroup.github.io/essential\_skills\_RRC/
  - Reproducible Research using Jupyter Notebooks <a href="https://reproducible-science-curriculum.github.io/workshop-RR-Jupyter/">https://reproducible-science-curriculum.github.io/workshop-RR-Jupyter/</a>
  - Duke UPGG Informatics Orientation Bootcamp <a href="https://duke-gcb.github.io/2019-08-12-">https://duke-gcb.github.io/2019-08-12-</a>
     Duke/
  - Reproducible Research and Data Analysis (under development) https://opensciencemooc.eu/modules/reproducible-research-and-data-analysis/
  - Reproducible research: Methodological principles for a transparent science https://learninglab.inria.fr/en/mooc-recherche-reproductible-principes-methodologiquespour-une-science-transparente/
  - Make (Software Carpentry's lesson) <a href="http://swcarpentry.github.io/make-novice/">http://swcarpentry.github.io/make-novice/</a>



#### Resources (3)

#### Tools:

- Popper <a href="https://github.com/getpopper/popper">https://github.com/getpopper/popper</a>
- Reana <a href="http://www.reanahub.io">http://www.reanahub.io</a>
- ReproZip <a href="https://www.reprozip.org">https://www.reprozip.org</a>
- Sciunit <a href="https://sciunit.run">https://sciunit.run</a>

#### Other:

- Reproducible PI Manifesto <a href="https://lorenabarba.com/gallery/reproducibility-pi-manifesto/">https://lorenabarba.com/gallery/reproducibility-pi-manifesto/</a>
- Computational science example (from FSCI 2018 & 2019): https://github.com/danielskatz/repro-fdtd1d
- Make your code ready for publication (sharable and citable) workshop -<a href="https://gitlab.com/hifis/hifis-workshops/make-your-code-ready-for-publication/workshop-materials">https://gitlab.com/hifis/hifis-workshops/make-your-code-ready-for-publication/workshop-materials</a>
- Software Citation Principles <a href="https://doi.org/10.7717/peerj-cs.86">https://doi.org/10.7717/peerj-cs.86</a>

