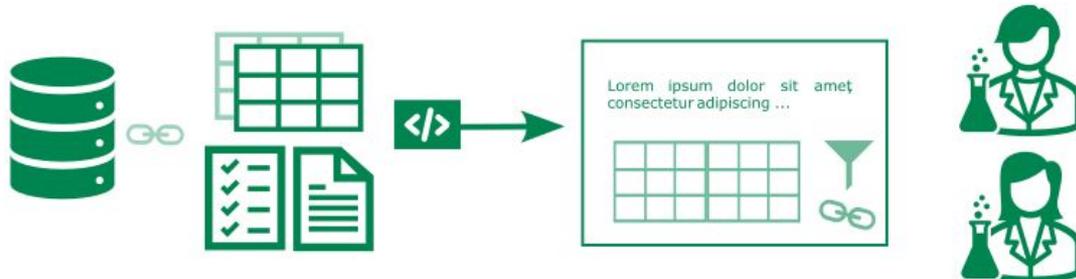


# Where are my data?

## The AFFORD workflow to create your own data index with Git, R and Quarto

Gorka Fraga González



## Goals

- Show researchers ways of improving data management with simple tools
- Enhance data reusability
- Promote workflows to improve computational reproducibility

## Target audience

- Researchers with an interest in data management
- Data stewards
- Requires: basic IT affinity and basic Git and R or Python

## Main Features

Affordable, simple, flexible, uses generalist and open source tools



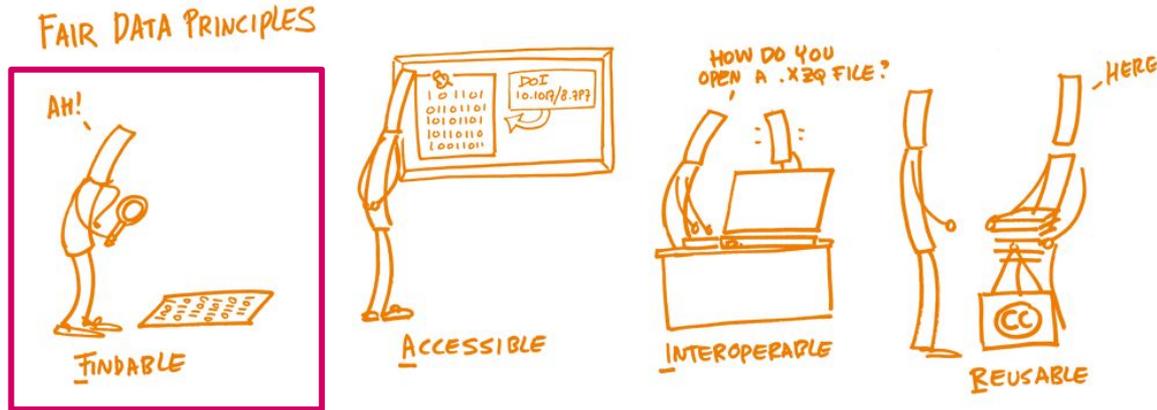


# Background

Findable, Accessible, Interoperable and Reusable (FAIR)

Guiding principles for managing research data. Focus on :

- Machine-readable **metadata**
- **Documentation**
- **Persistent identifiers**
- **Interoperable** formats and vocabularies
- **Licensing**
- **Publishing** in accessible repositories



# Background

## FAIR data is **challenging**

- **Interdisciplinary** differences in technical skills and research culture
- **Data-intensive** research: complex pipelines, heterogeneous file types
- Heterogeneity of resources for data **curation** (e.g., stewards, IT)
- **Fragmented landscape** of tools
- Time-consuming **decision making** and **coordination**

# A data index

*“To share your data you first need to find it” - Anonymous*

**Data index:** **catalogue** that allows browsing through data by using the available metadata.

# A data index in the AFFORD project

in AFFORD our **target** data index should:

- Provide enough **context** (metadata) to the data
- Allow to find where the **source data** files
- **Privacy**: first, it should be possible to share only internally
- Eventually, it should be **public** and useful for all
- **Maintainable** by researchers, **scalable** and **portable**

# A data index in the AFFORD project

## Data



- Large volume stored externally
- Different storage locations
- Subsets expected to be published online
- Ongoing data collection
- Different facilities -> different constraints (e.g., filenames, folder structure)
- Different data types

# A data index in the AFFORD project

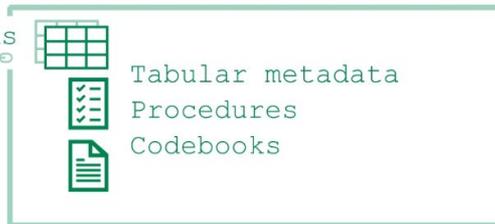
## Data



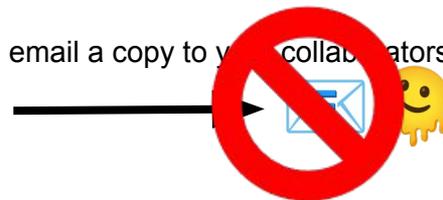
Findable  
Accessible  
Interoperable  
Reusable

## Metadata

links  
∞



email a copy to your collaborators



- Large volume stored externally
- Different storage locations
- Subsets expected to be published online
- Ongoing data collection
- Different facilities -> different constraints (e.g., filenames, folder structure)
- Different data types

### Minimum FAIRNESS requires

- Machine-readable metadata
- Procedures
- Codebooks
- Documentation

# A data index in the AFFORD project

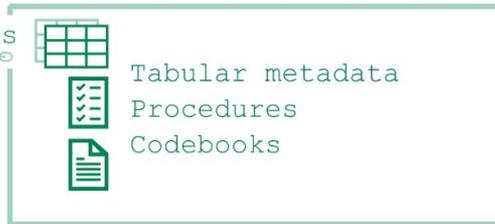
## Data



Findable  
Accessible  
Interoperable  
Reusable

## Metadata

links  
GO



code



## Data Index



collaborators  
navigate



- Large volume stored externally
- Different storage locations
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- Different data types

Minimum FAIRNESS requires

- Machine-readable metadata
- Procedures
- Codebooks
- Documentation

Metadata + simple code to enable SHARING:

- Navigate through metadata
- Filters
- Clickable links to data location
- OPTIONALLY: representative data previews

# A data index in the AFFORD project

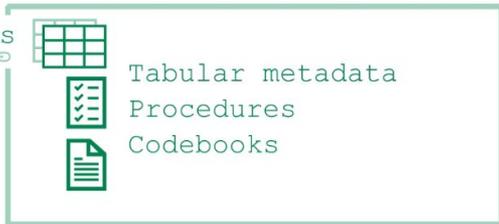
## Data



Findable  
Accessible  
Interoperable  
Reusable

## Metadata

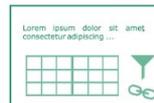
links  
GO



code



## Data Index



collaborators  
navigate



- Large volume stored externally
  - Different storage locations
  - Subsets expected to be published online
  - **FAIR** is much more: full machine-actionability, controlled vocabularies, ontologies, integration in larger data frameworks, etc.
  - But we start with an unambitious, basic level FAIRness that would enable data to be found and understood by other researchers.
- Minimum FAIRNESS requires
- Machine-readable metadata
  - Procedures
- Metadata + simple code to:
- Navigate through metadata
  - Filters
  - Clickable links to data location
  - OPTIONALLY: links to tentative data previews

# The workflow

It uses 3 components important for computational reproducibility:

1. **Dynamic document generation**
2. Repositories with **Git version control**
3. Software **containerization**



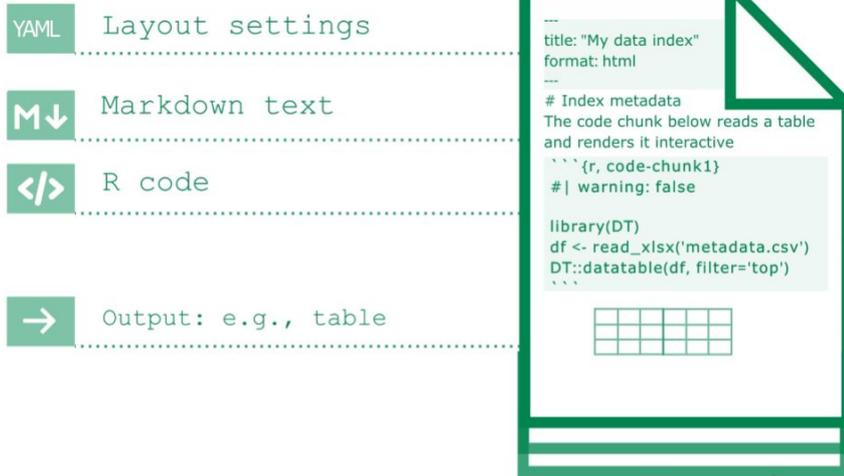
# The workflow

It requires researchers to create first:

- **Metadata tables** in CSV format (comma separated value, favoured for interoperability). Adequately formatted so that is machine readable
- **Codebooks** describing the variables



# The workflow



## 1. Dynamic document generation

We use Quarto, integrated in Rstudio. It can read a file with:

- Header settings/metadata (YAML)
- Markdown text
- Executable Code (R, Python,...)

And print it into multiple formats:

- HTML, pdf, word, etc.

Group docs with different layouts:

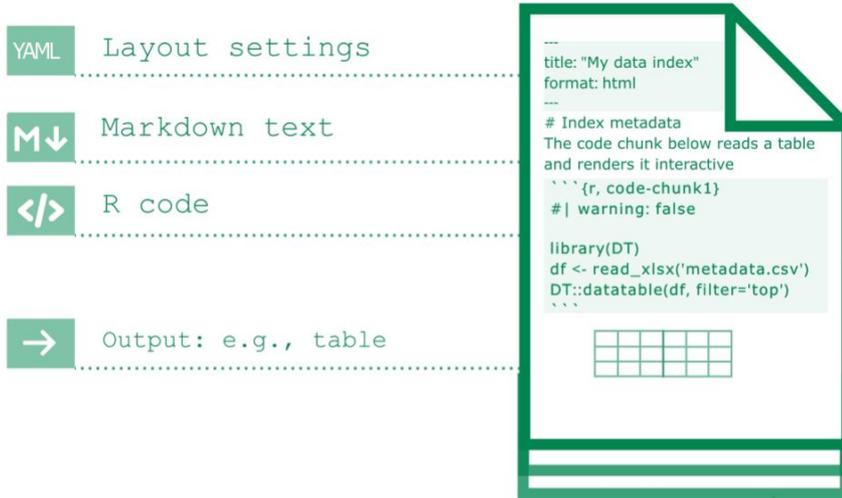
- Websites, book, reports, etc

# The workflow

YAML header. Title, author information, type of output, document , etc.

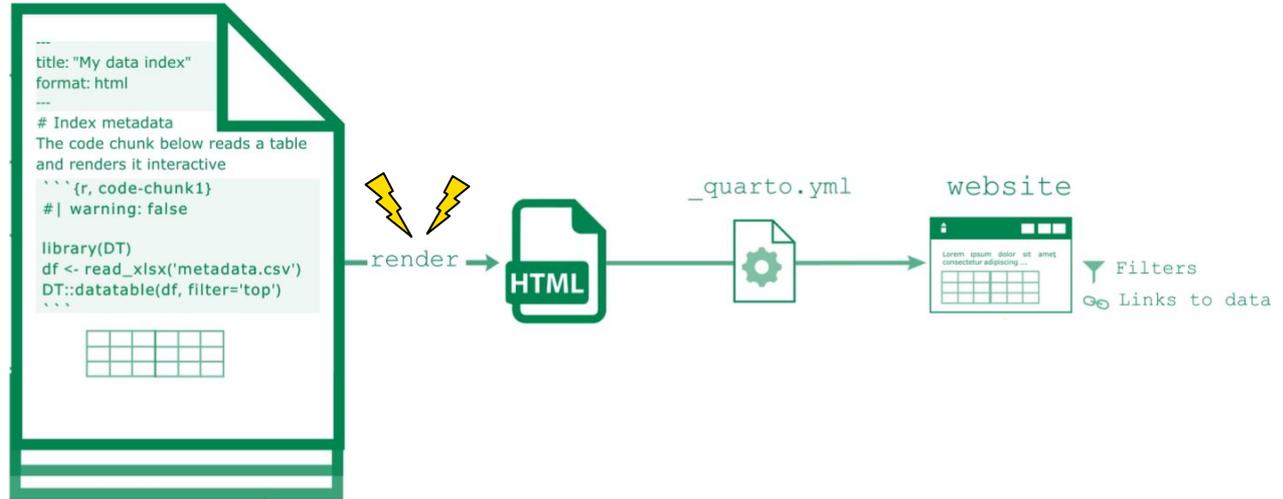
Markdown text. Includes instructions, introductory descriptions, etc.

R code chunk. Reads .CSV tables with metadata and *generates (render)* an interactive table with filters options



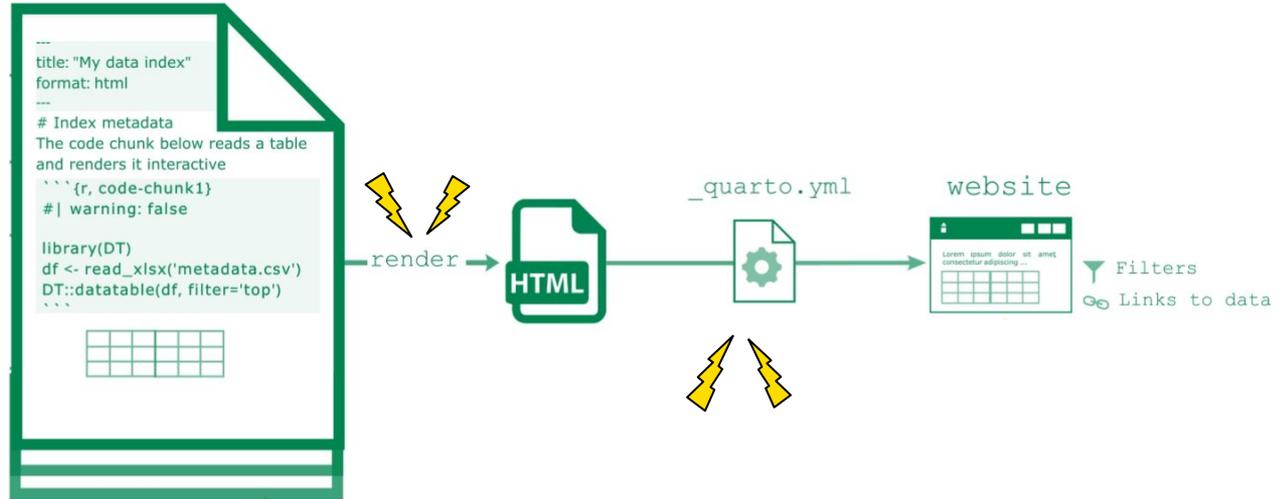
Note: these files have .qmd extension. They can be opened and edited with any text editor (e.g., notepad) as they are plain text. But to execute you need Quarto and R (e.g., in R studio)

# The workflow



When we run the command `quarto render`, the code is executed and the file is saved into a nicely-formatted HTML with an interactive version of the table (with filter panels or any additional panel we want to add in the code)

# The workflow



(RECOMMENDED) configuration file *\_quarto.yml* (*quarto project file*) is also a plain text file with a series of options offered by quarto to define the project type and layout. In this case we have a 'website' that combines multiple HTML files as there are several tables.

# The workflow

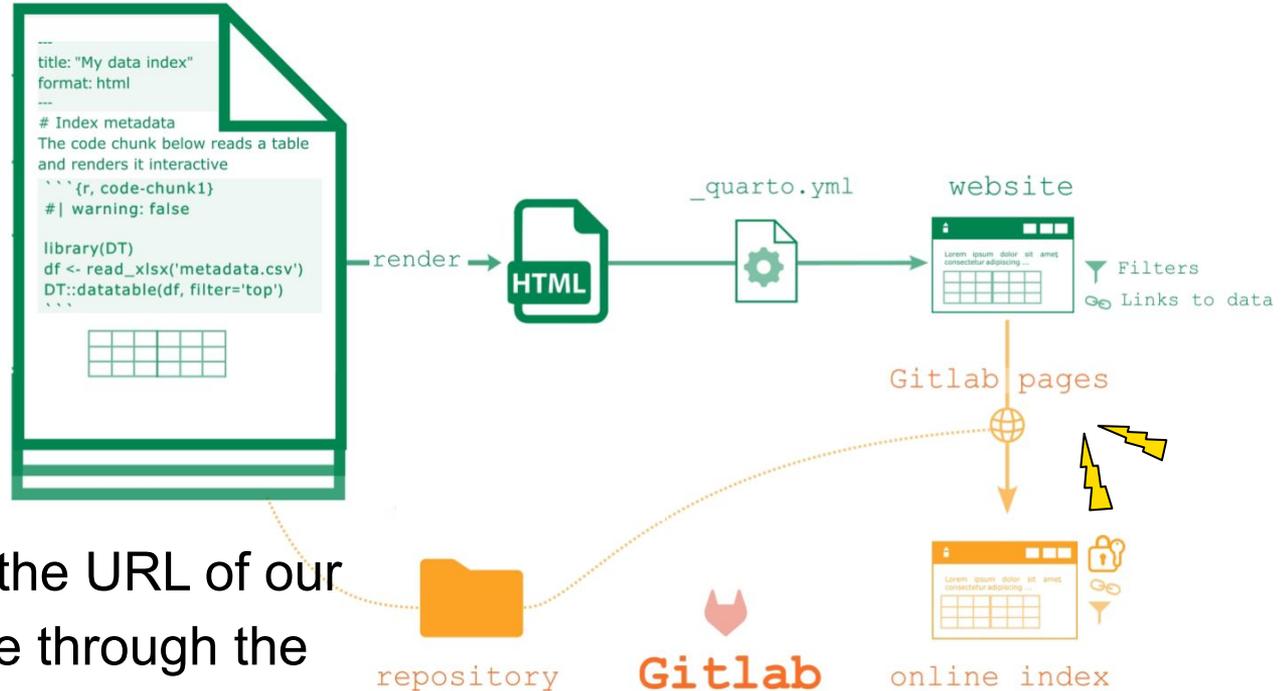
## 2. Version control with Git.

Metadata tables (.CSV) and source Quarto files to generate reports (.QMD) are stored in a Git repository. Git is a decentralized system for tracking changes in files

- Our repository is hosted by [Gitlab](#) (instance managed by UZH)
- Gitlab offer [Gitlab pages](#) a services to make HTML documents accessible from an internet browser
- Gitlab pages can be **private**
- They are **static** pages (all users see only whatever is in the HTML)



# The workflow



Collaborators can go to the URL of our Gitlab page and navigate through the metadata tables

# The workflow

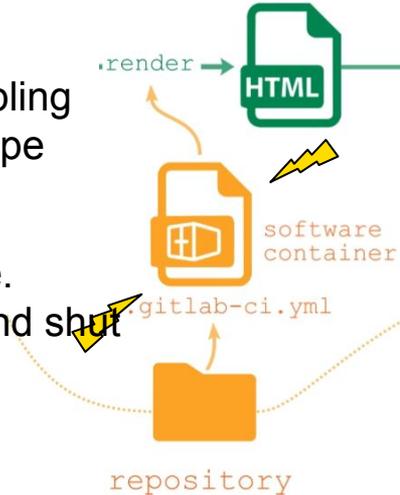
## 3. Software containerization

The code to make the interactive table and website should not break after a package is updated. We run it always through an ‘encapsulated software environment’

 **Recipe.** Human and machine-readable text file. A blueprint of the environment

 **Image.** Machine-readable file assembling and storing all components in the recipe

 **Container.** A working copy of the environment assembled in the image. Meant to be temporary: it is started and shut down after use.



# The workflow

## Gitlab Continuous Integration (Gitlab) file

Every time we change the repository (e.g., new entry on the table). This file executes the code from the software container

```
image: rocker/verse:4.5.0
```

A container image is downloaded the [Rocker](#), a website with open images of R software

```
pages:
  stage: deploy
  before_script:
    # Install additional packages using dependency version control by rocker
    - install2.r kableExtra openxlsx DT here fs stringr

  script:
    # Create site folders
    - mkdir ./public
    # Call python to convert inputs from the CMS into .qmd and add code chunks if needed
    - python3 ./admin/convert_md_to_qmd.py ./webpage_contents/ ./site_qmds/
    # Go to folder with QMDs and the quarto project file, render the site from there
    - cp -r ./webpage_contents/* ./site_qmds/
    - cd ./site_qmds
    - quarto render
    - cp -r ../_site/* ../public/
    - cp -r ../admin/ ../public/
    - cp -r ../public/
```

Extend rocker image with additional packages

in-house code to convert entries as .md into .qmd (see CMS extension in next slides)

Create website's HTML

```
artifacts:
  paths:
    - public

only:
  - main

interruptible: true
```

# The workflow

## Gitlab Continuous Integration (Gitlab) file

Users don't need to be executed from our computer. The application that is executing the code using the container is the [Gitlab runner](#) .

This code is very simple and not computationally demanding so we will not need additional resources to run it.

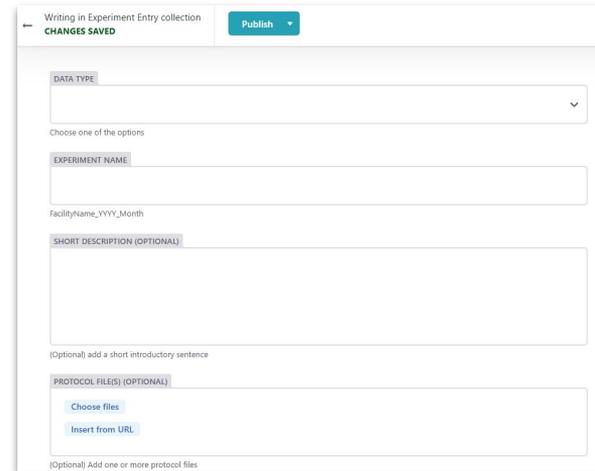


# Variations and extensions

- Example [fabric4.ch](https://fabric4.ch) from the actual research project supported in AFFORD
- The table can also display **pictures** stored in the Repository. Example: [https://crsuzh.pages.uzh.ch/AFFORD\\_website/ORD\\_index/](https://crsuzh.pages.uzh.ch/AFFORD_website/ORD_index/)
- A **simpler** use case: e.g., [crsuzh.pages.uzh.ch/datastew/](https://crsuzh.pages.uzh.ch/datastew/)

## A slightly more complicated workflow:

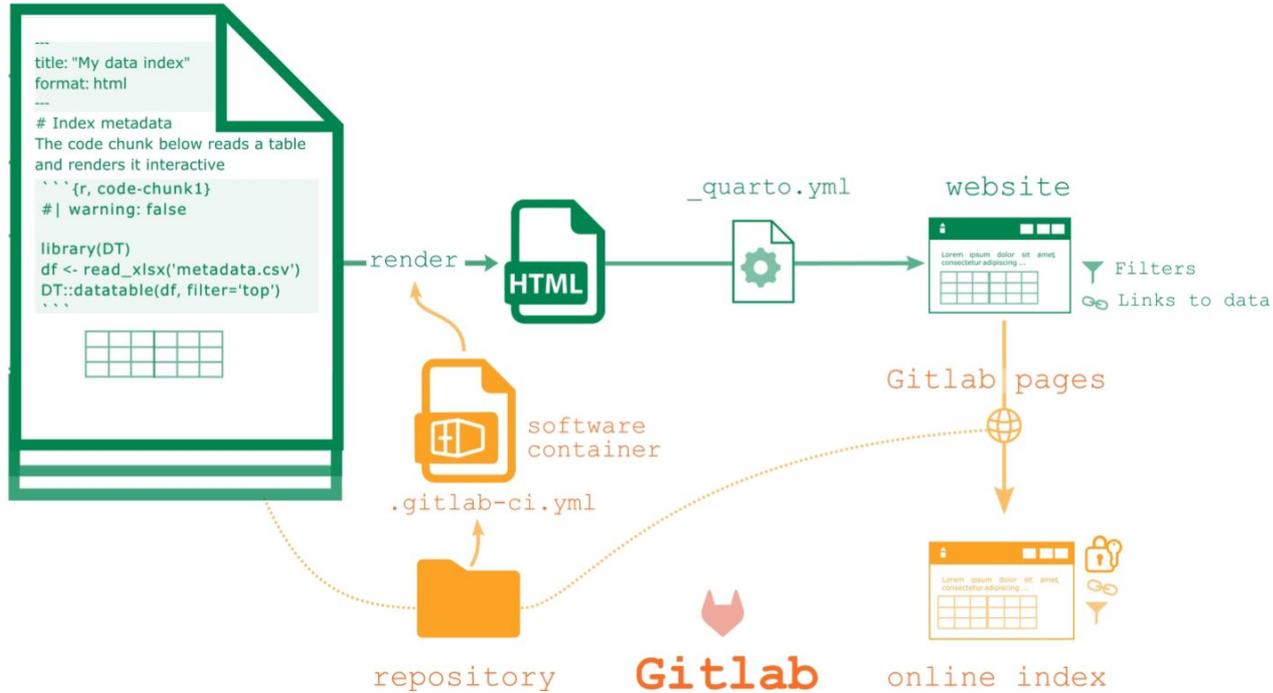
- To facilitate **data entry** for dummies, the application [DECAP CMS](#) can be used to have a simple Content Management System (CMS). A demo repository with this CMS is work in progress



The screenshot shows a web form for entering an experiment entry. At the top, it says "Writing in Experiment Entry collection" and "CHANGES SAVED" with a "Publish" button. The form has several sections:

- DATA TYPE:** A dropdown menu.
- Choose one of the options:** A text input field.
- EXPERIMENT NAME:** A text input field with the placeholder "FacilityName\_YYYY\_Month".
- SHORT DESCRIPTION (OPTIONAL):** A large text area with the instruction "(Optional) add a short introductory sentence".
- PROTOCOL FILES (OPTIONAL):** A section with two buttons: "Choose files" and "Insert from URL". Below it is the instruction "(Optional) Add one or more protocol files".

# The workflow



This code is just for illustration. Ignore the error reading using read\_xlsx for a csv table 🙄(ツ)🙄

# Expected actions from researchers / data stewards

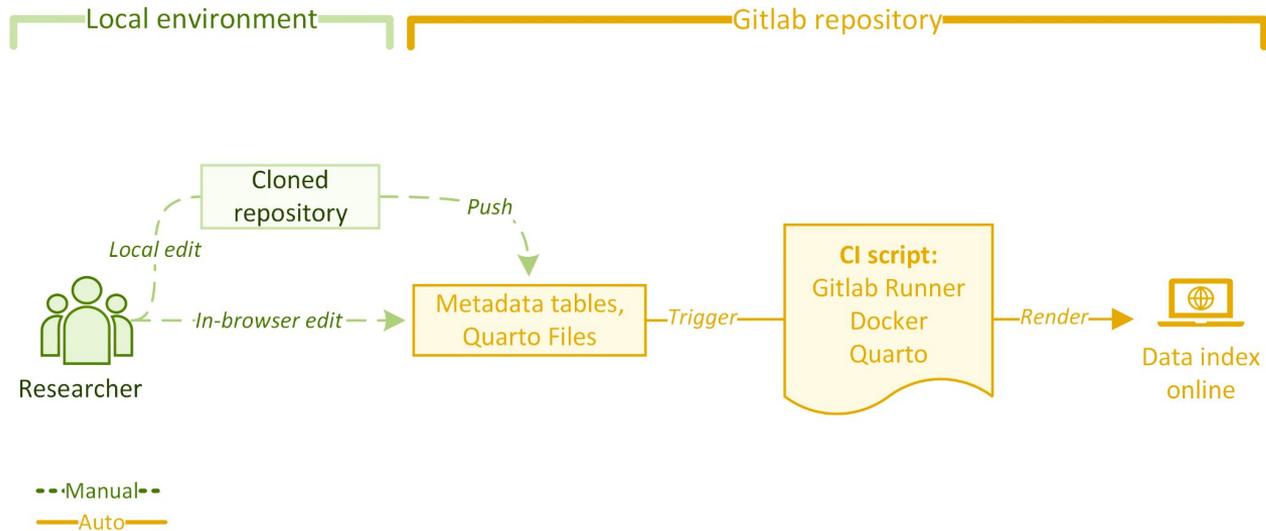
To **set up** this workflow requires familiarity with Git, R and Quarto is needed. You can start by copying this **demo repository** and edit it:  
[https://crsuzh.pages.uzh.ch/afford\\_repos/affordable\\_data\\_index/](https://crsuzh.pages.uzh.ch/afford_repos/affordable_data_index/)

## Recommendations for data stewards

- Explore customizing the workflow
- Keep it simple and easy to maintain in the future
- Consider your collaborators expertise
- Don't make it entirely dependable of the one person that set it up
- Other coding languages and Git platforms enable similar actions
- Similar workflow can be applied to analytic code, visualizations, etc

# The actions from researchers

Collaborators can edit the metadata through the browser (or via a local copy of the repository). The website updates automatically after changing the table



- 1 ACTION: update repository
- 2 Changes trigger the CI pipeline
- 3 HTML is updated
- 4

# Making the Index Citable

- Citing the **URL** to a Gitlab page is **not recommended**
- Make an entry in a platform enabling a Digital Object Identifier (**DOI**) (e.g., Zenodo, Open Science Framework) and link it to the URL
- If the content is stable, upload a copy of the code repository ('release') including the **source metadata tables**
- If not stable, users will still access the website through the **URL at the DOI entry**. If the URL changes , the URL in the DOI entry can be changed without requiring a new DOI

# Challenges

- **well-formatted metadata tables** cannot be taken for granted
- PIs need to **agree** on minimum actions to integrate all groups
- future maintainers will need additional **training** (e.g., Git commits)
- if made too easy for editing (e.g., CMS) it won't be as easy to **maintain code** for a researcher-steward

# Summary and discussion

This workflow

- needs **machine-readable metadata** tables, encouraging FAIR efforts
- uses 3 important elements for **computational reproducibility** (dynamic reporting, Git, containers)
- It is very **flexible**, adaptable to other tools, environments and needs

Considerations

- Find a **balance** between features and maintenance efforts
- The metadata **CSV** tables are the most important. The workflow just facilitates **sharing** in a central, public (or private) platform, **version controlled** and **avoiding duplicates**

# Additional resources

Preprint detailing the workflow: <https://osf.io/preprints/metaarxiv/64fch>

## High-level recommendations and ‘reality checks’ from AFFORD

Fraga-González, G., van de Wiel, H., Garassino, F. *et al.* Affording reusable data: recommendations for researchers from a data-intensive project. *Sci Data* 12, 258 (2025). <https://doi.org/10.1038/s41597-025-04565-0>

## Easy-reads for PIs: the [CRS primers](#)

- on software containers: <https://doi.org/10.5281/zenodo.13757230>
- on digital collaboration: <https://doi.org/10.5281/zenodo.8354375>

# Thank you for your attention!



- Requests? Improvements? Write me! [gorka.fragagonzalez@uzh.ch](mailto:gorka.fragagonzalez@uzh.ch)
- Follow us on [LinkedIn](#) for regular updates
- Check out the blog of [SwissRN](#) working group [Open Research Data](#)



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