





AFFORD deliverables WP2.3.2: Open Research Data (ORD) hub in Gitlab pages

https://crsuzh.pages.uzh.ch/afford_page/

Content

Overview

'Front-end'

- Hub for ORD sharing
- Wiki collaborative documentation

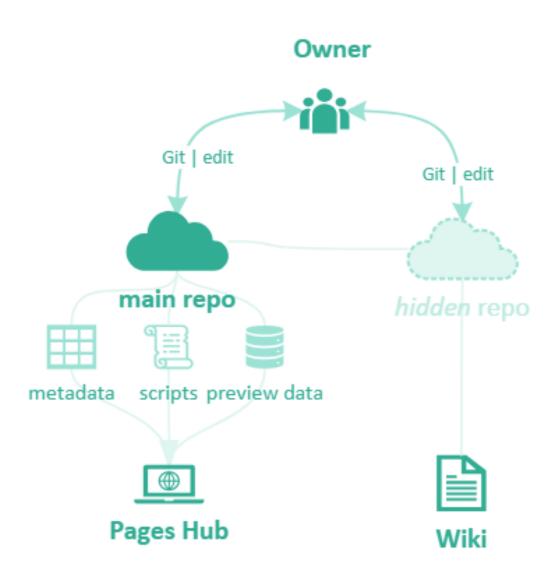
'Back-end'

- Data flow link preview data to sources
- Metadata create standards

Summary & Next steps ...



Overview of Gitlab workflow

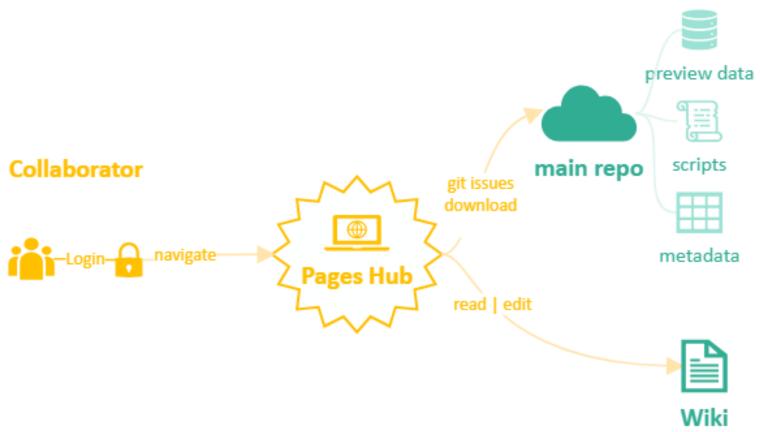


Owner - A lab or researcher managing a project or a series of data packages. They could have a **Gitlab group** set up e.g., *gitlab.uzh.ch/interfacegroup*

Repository scope - e.g., Data from Interface group for the Fluid Dynamics CNS Sinergia project will include several synchrotron data packages from Japan, France, etc.

e.g., gitlab.uzh.ch/interfacegroup/FluDynCNS

Overview of Gitlab workflow



Collaborator - A lab or researcher in direct collaboration with the owner e.g., other members of the Fluid Dynamics CNS project

Hub

What?

- Central hub to metadata, documents, preview data and source data links

Why?

- Improve findability, accessibility, interoperability, showcasing data
- Version control and avoid duplicate files and data transfer

How?

- Gitlab repository with access public or restricted to collaborators



Hub

In-browser Hub Demo: https://crsuzh.pages.uzh.ch/afford_page/

Highlight:

- Arrangement by tab
- Selection and copying of multiple rows
- Simple coding (R Markdown streamlining in progress)
- Data table additional extensions
- Continuous integration (show repo)



Wiki

What?

- Central documentation site (e.g., Fluid dynamics – Interface)

Why?

- Same as hub: version control, findability, accessibility
- Wiki is easy to edit by collaborators
- Markdown keeps formatting simple and with high compatibility

How?

- Markdown files in Wiki through Gitlab repository



Wiki

In-browser Wiki Demo:

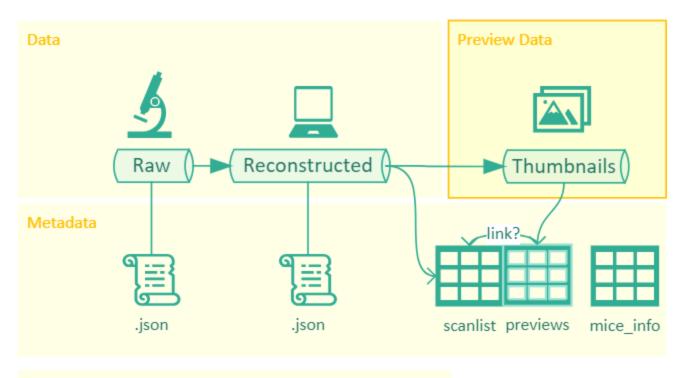
https://gitlab.uzh.ch/crsuzh/afford_page/-/wikis/Home

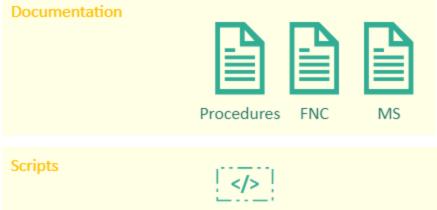
Highlights:

- Track versions history, 'last-edited-by' info
- Rich text editor
- Note: it is a 'hidden' repository separate from Gitlab pages

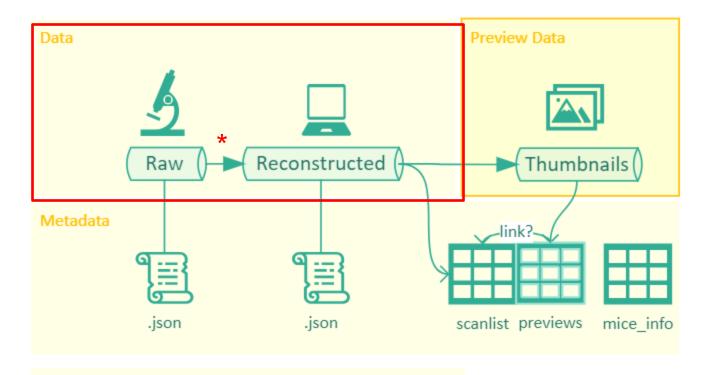


Per data package (e.g., Japan-Spring 2023)





Per data package (e.g., Japan-Spring 2023)

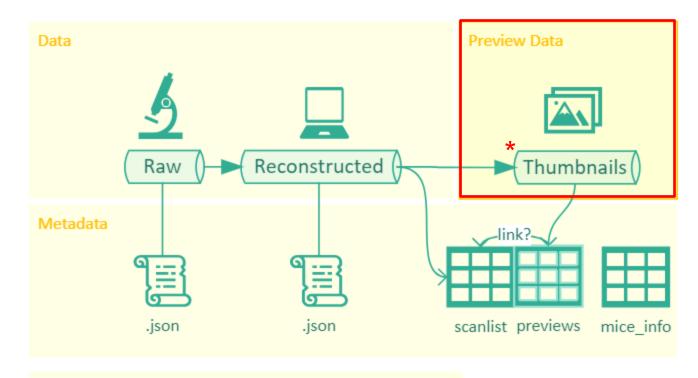


The recorded **data** may undergo minimal preprocessing or **reconstruction** to be useable

- * The script to do this should be provided and documented with all parameters
- * Metadata of reconstructed data should contain the relevant reconstruction parameters (see JSON)



Per data package (e.g., Japan-Spring 2023)



'Preview data' is generated to assess quality, validate, explore results, share with collaborators: that is the data showed in the local HUB

* The **script** to create this data should be provided / documented in the repo

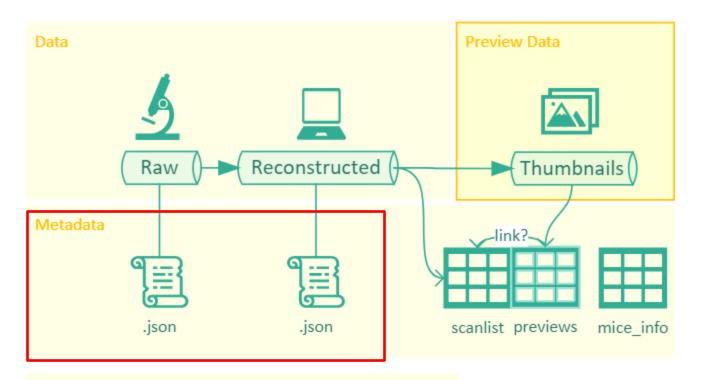
The **link** between the preview file and the source data should be **preserved** (e.g., in metadata tables)

Metadata should also indicate **what exactly** the thumbnail shows (e.g., center slice n. xx)



- e.g., Slice 749 or slice 750 of the reconstructed image? Wiki info vs power shell
 - Now we have 3 scripts to go from reconstructed to preview data: one powershell script to rename(?) some images and then two R scripts to compress them to jpg and extract filename metadata for the hub table.

Per data package (e.g., Japan-Spring 2023)



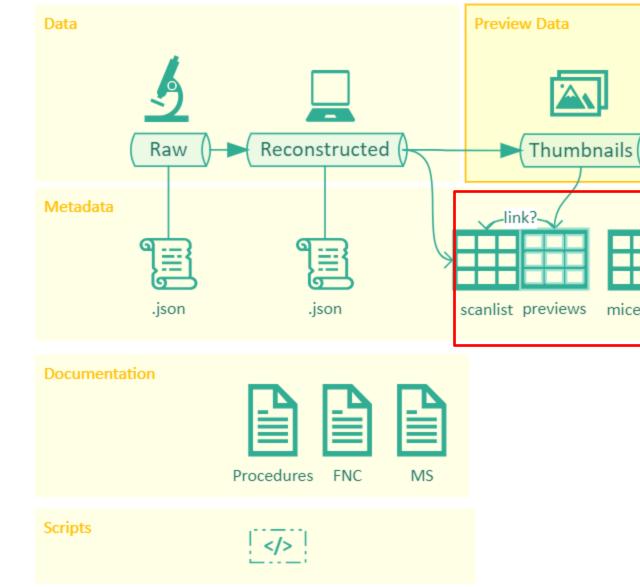
JSON metadata files to accompany each data file

- Basic data **provenance** information (e.g., where, by who, when it was collected)
- More advanced provenance info: Recording parameters, software, hardware, and anything that happened to the file from recording



See JSON format proposals later in this presentation

Per data package (e.g., Japan-Spring 2023)



Metadata tables

mice info

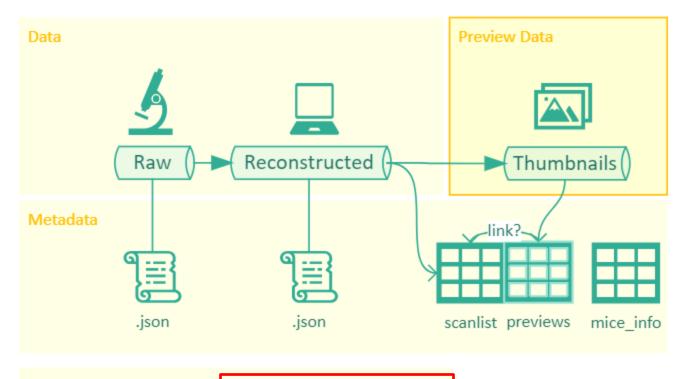
Wide-formatted tables
Summarizing the relevant files of the data package

- Scanlist files describing source recordings
- Previews metadata that can be rendered interactive linking to preview images (HUB)
- Mice_info specimen information (wide format, one specimen per row)
- Additional tables may be necessary for lab tests, control measurements, etc.

A **codebook** should clarify variable names for each table

Located at Gitlab repository: **no duplicates** should exist

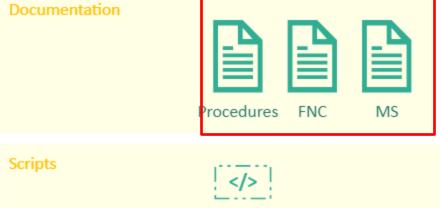
Per data package (e.g., Japan-Spring 2023)



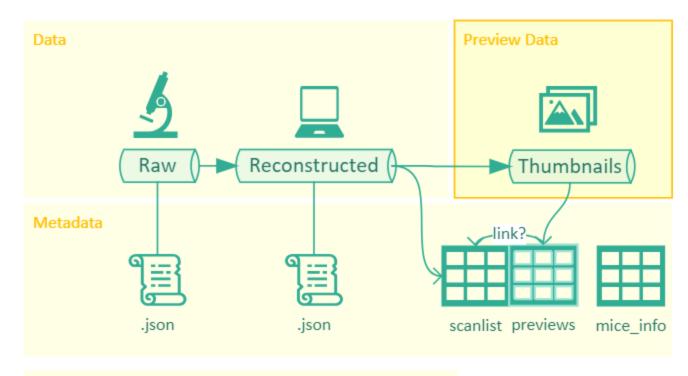
Documentation

Lab procedures, file naming convention and metadata specifications to clarify how data, metadata and scripts are organized

Also in the Gitlab repository



Per data package (e.g., Japan-Spring 2023)





Scripts

Scripts

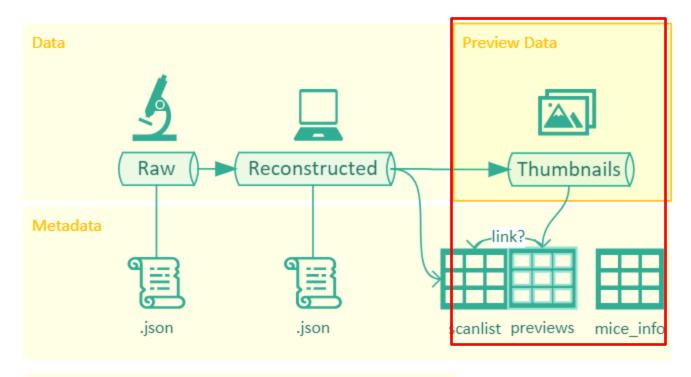
All code to:

- organize the data package
- generate metadata json and tables
- Generate thumbnails
- Rename or do other file edits
- render the Hub

It can be stored in the same Gitlab repository as the website with the Hub

Code snippets or small scripts can also be written in the wiki as part of the documentation

Per data package (e.g., Japan-Spring 2023)



In Gitlab Repo



Summary of a data package content

File	Content	Formatting Issues
Data	Comon	T cimating locate
Raw	Directly measured from instruments	
Reconstructed	Minimally preprocessed so that it can be interpreted	
Thumbnails	Preview images for sharing, inspection, QA, etc	
Documentation		Markdown recommended (e.g., in wiki)
Procedures	How the data was collected and reconstructed	II .
FNC	Filename convention	II .
MS	Metadata specifications: fields descriptions, how tables were generated	n .
Metadata tables		Incl. codebook sheet (variable dictionary) No color coding for interoperability. Wide-format
Scanlist *.xlsx	List of available files and their recording parameters. If applicable, the corresponding thumbnail files	rendered interactive for navigation in the hub
Mice info *.xlsx	Additional info per specimen	
Lab control *.xlsx	Additional info on lab calibration, tests, etc	
hash tables	Archived files and hashes for verification	
Metadata json (per data file)		
<datafilename>.json</datafilename>	To be stored together with each individual data file, basic data provenance tracking (e.g., where, what equipment, who collected, basic rec parameters)	Script-generated: Common info hardcoded, file-specific info derived from data

Metadata standards and JSON files

What?

- Metadata files stored together with recorded datasets

Why?

- Metadata tables are not stored with each data record (e.g., spec-01 folder) and do not contain all info per data object or file
- Why .JSON? Concise, compact, and easy to parse and generate

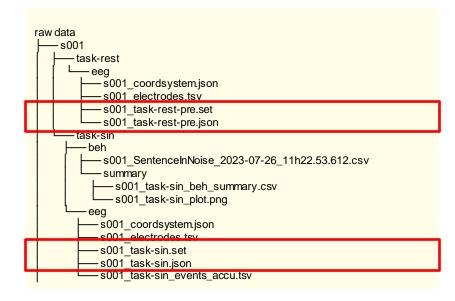
How?

- Manual generation is not recommended (scriptable with any language) :
- Hardcoded info (e.g., project title) + data-derived info (e.g., duration)

Metadata standards and JSON files

Example of an EEG project following BIDS guidelines

Each EEG dataset (.set) comes with a .json file (with same name). Provides common info to all project files as well as file-specific info (e.g., duration)



```
| s001_task-sin_eeg.json ■
            "ProjectName": "Speech in noise EEG",
           "TaskName": "task-sin".
           "TaskDescription": "A version of a speech intelligibility task us:
           "Instructions": "Listen well and click on the images representing
           "InstitutionAddress": "LiRI Linguistic Research Infrastructure, Un
           "InstitutionName": "LiRI Linguistic Research Infrastructure, Unive
           "PowerLineFrequencyHz": 50.
           "Manufacturer": "BIOSEMI",
           "ManufacturersModelName": "ActiveTwo MK2HS",
           "RecordingType": "continuous",
 12
           "RecordingDurationSec": 3727.271,
           "EEGPlacementScheme": "International 1020 system",
 14
           "EEGReference": "CMS/DRL",
           "SamplingFrequencyHz": 2048.
 16
           "EEGChannelCount": 64,
 17
           "MiscChannelCount": 2,
 18
           "TriggerChannelCount": 1,
 19
           "EOGChannelCount": 4,
 20
           "ECGChannelCount": 0.
 21
           "EMGChannelCount": 0,
 22
           "SoftwareFilters": "n/a",
 23
           "HardwareFilters": "n/a",
 24
           "Comments": "Data set created from spliting the source .bdf file
 25
```

Metadata standards and JSON files

Example of an EEG project following BIDS guidelines

Additional json files containing details on the recording system

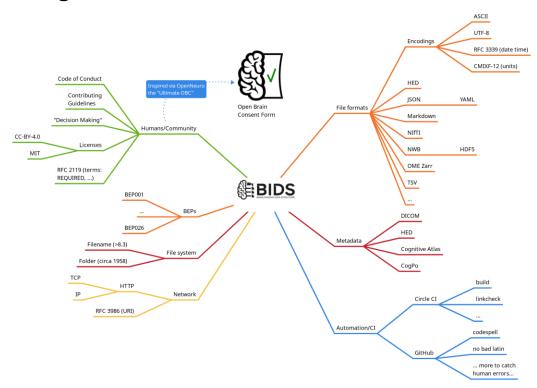
E.g., coordinate system, to interpret the coordinates provided in metadata file 's001_electrodes.tsv'

```
"EEGCoordinateSystem": "EEGlab",
"EEGCoordinateUnits": "mm",
"EEGCoordinateSystemDescription": "https://eeglab.org/tutorials/ConceptsGuide/coordinateSystem.html",
"IntendedFor": "s001_task-sin"
```

- Are there existing standards in the field?

Brain Imaging Data Structure https://bids.neuroimaging.io/

"Most advanced and widely used option for structuring files according to MRI sequences and including standardized metadata" Kalantari et al, 2023



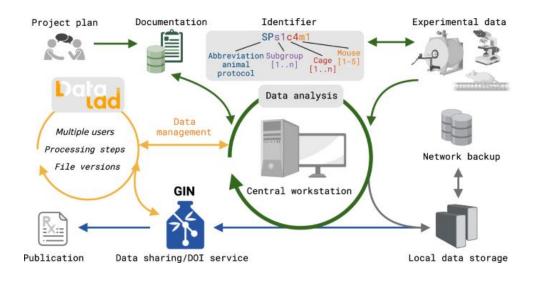
BIDS Extensions Proposals (BEP)

https://bids.neuroimaging.io/get_involved.html#extending-the-bids-specification

- You can propose a BIDS extension to your data modality
- Each proposal is an open google doc where experts can give comments
- Once accepted it will be integrated in the BIDS
- Currently active BEPs: CT, computational models, microscopy, animal electrophysiology, ...
- Integration with BIDS will facilitate visibility, adoption and version tracking

- Publications related to BIDS extensions

Kalantari, A., Szczepanik, M., Heunis, S., Mönch, C., Hanke, M., Wachtler, T., & Aswendt, M. (2023). How to establish and maintain a multimodal animal research dataset using DataLad. Scientific data, 10(1), 357. https://www.nature.com/articles/s41597-023-02242-8



- DataLad opensource software for research data management https://www.datalad.org/
- YODA principles for data analysis https://handbook.datalad.org/en/latest/basics/101-127-yoda.html
- BIDS compatibility of structure and metadata

- There are other publications related to BIDS extensions

Bourget, M. H., Kamentsky, L., Ghosh, S. S., Mazzamuto, G., Lazari, A., Markiewicz, C. J., ... & Salo, T. (2022). Microscopy-BIDS: An extension to the Brain Imaging Data Structure for microscopy data. *Frontiers in Neuroscience*, *16*, 871228. https://www.frontiersin.org/articles/10.3389/fnins.2022.871228/full

Next steps on WP2

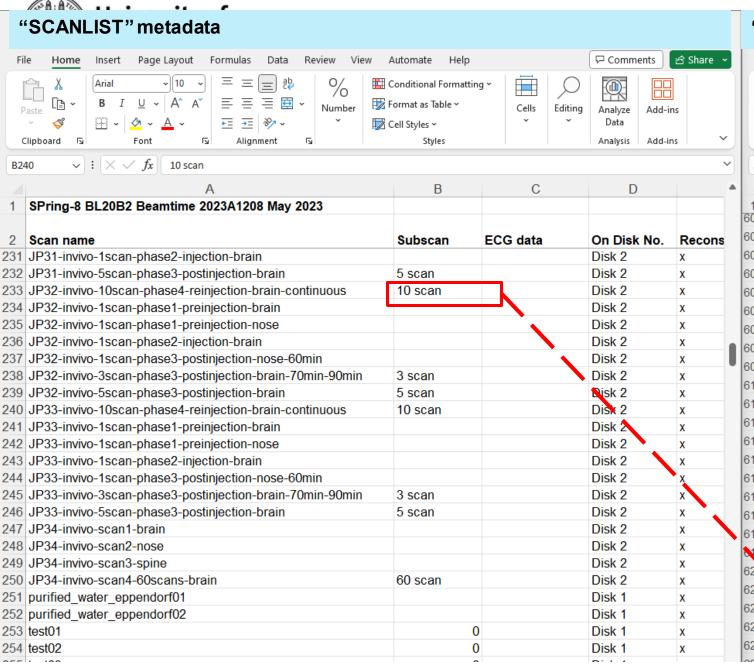
CRS

- Hub table code generalizable (troubleshoot, streamline, documentation)
- Beginner Tutorials (currently collection for beginners)

Interface group (+ CRS)

- Defining JSON fields based on BIDs proposals and own data
- Clarifying workflow from source data to preview files
- Search if other standards (besides BIDS) exist for data types in FDCNS
- Share hub with FDCNS collaborator to attempt a set up from scratch

Thank you



"Preview" files metadata (showed in HUB)

