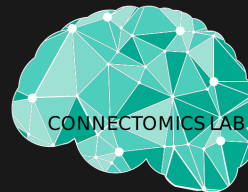


A BIDS MRI DATASET USECASE

NCCR-SYNAPSY Early-Psychosis Dataset

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EARLY-PSYCHOSIS DATASET

- Split into 2 datasets:
 - BIDS compliant dataset created from DICOMs: 346 subjects
 - Dataset of participants with missing DICOMs: 12 subjects
- BIDS dataset created from DICOMs validated with [bids-validator](#):

```
$ bids-validator "/media/localadmin/HagmannHDD/Seb/PsychoDev/NiftiBIDSCompliant" --ignoreSubjectC
bids-validator@1.7.2
```

```
[...]
```

Summary:

9212 Files, 47.25GB

346 - Subjects

6 - Sessions

Available Tasks:

rest

Available Modalities:

MTR

T1w

T2map

dwi

bold

T1map

fieldmap

DATASET HISTORY

- Creation of pre-configured empty Datalad dataset
- Copy the content of the initial BIDS dataset with a few problems such as the use of the acquisition date as a session label.
- Edit the CHANGES

```
0.1 2021-06-05
```

```
- Initial commit of BIDS dataset after reconversion of DICOMs to Niftis using dcm2niix
```

- Save the dataset with version tag 0.1

DATASET HISTORY

- Add and run a script in code/ folder to fix BIDS validation errors
- Validate dataset with the bids-validator
- Edit the CHANGES

0.2 2021-07-07

- Create bidsify_dataset.py in code/ folder
- Run code/bidsify_dataset.py that:
 - * Add CHUV prefix to BIDS participant_id labels
 - * Change session labels from acquisition date to scan1, scan2, ...
 - * Add sub-<label>_sessions.tsv/.json files that stores the participant age in month and the acquisition date
 - * Generate participants.tsv/.json files that stores the participant sex and the group

- Save the dataset with version tag 0.2
- Publish the dataset to our institutional data storage accessible via ssh

DATASET AVAILABILITY

Until now

- Dataset can be installed/cloned with:

```
$ datalad clone ssh://<user>@stockage-horus.chuv.ch:/archive/PRTNR/CHUV/RADMED/phagmann/biopsycho/NiftiBI
```

Note: one needs to be inside CHUV intranet or to use CHUV VPN and have access to **stockage-horus** to be able to clone the dataset

In the near future

- Dataset could be installed/cloned with:

```
$ datalad clone git@github.com:NCCR-SYNAPSY/ds-biopsycho
```

Note: everybody would be able to clone the dataset from GitHub but one still needs to be inside CHUV intranet or to use CHUV VPN and have access to **stockage-horus** to be able to retrieve the content of the files

TIPS, TRICKS AND DEMOGRAPHIC REQUIREMENTS

HOW TO DEAL WITH LONGITUDINAL AND MULTI-SITE STUDIES

BIDS-compliant solution

- Add an extra layer of directories and file names in the form of **ses-<label>** to encode multiple sessions (visits)
- Use a subject label prefix identifying the site (CHUV here)

```
├── sub-CHUVA001
│   ├── ses-scan1
│   │   ├── anat
│   │   │   ├── sub-CHUVA001_ses-scan1_run-1_T1w.json
│   │   │   ├── sub-CHUVA001_ses-scan1_run-1_T1w.nii.gz
│   │   │   └── ...
```

Reference: [Longitudinal and multi-site studies](#) section of the specifications

HOW TO DEAL WITH SIMILAR MRI ACQUIRED DURING THE SAME SESSION

BIDS-compliant solution

- Use BIDS `_run-<label>` entity to differentiate scans with same acquisition parameters
- Use BIDS `_acq-<label>` entity to differentiate similar scans with different acquisition parameters

```
sub-CHUVA022
├── ses-scan1
│   ├── anat
│   │   ├── sub-CHUVA022_ses-scan1_run-1_T1w.json
│   │   ├── sub-CHUVA022_ses-scan1_run-1_T1w.nii.gz
│   │   ├── sub-CHUVA022_ses-scan1_run-2_T1w.json
│   │   ├── sub-CHUVA022_ses-scan1_run-2_T1w.nii.gz
│   │   └── ...
│   └── dwi
│       ├── sub-CHUVA022_ses-scan1_acq-dsiNdir129_run-1_dwi.bval
│       ├── sub-CHUVA022_ses-scan1_acq-dsiNdir129_run-1_dwi.bvec
│       ├── sub-CHUVA022_ses-scan1_acq-dsiNdir129_run-1_dwi.json
│       ├── sub-CHUVA022_ses-scan1_acq-dsiNdir129_run-1_dwi.nii.gz
│       ├── sub-CHUVA022_ses-scan1_acq-hardiNdir147Bval3000_run-1_dwi.bval
│       ├── sub-CHUVA022_ses-scan1_acq-hardiNdir147Bval3000_run-1_dwi.bvec
│       ├── sub-CHUVA022_ses-scan1_acq-hardiNdir147Bval3000_run-1_dwi.json
│       └── sub-CHUVA022_ses-scan1_acq-hardiNdir147Bval3000_run-1_dwi.nii.gz
```

Reference: [Entities](#) section of the specifications

REQUIREMENTS FOR DEMOGRAPHIC INFORMATION

- At least `participant_id`, `age`, `sex` and `group` MUST BE REPORTED
- If the study IS longitudinal:
 - `participant_id`, `sex` and `group` MUST be reported in the `participants.tsv` file
 - `age` MUST be reported in the `session.tsv` file
- If the study IS NOT longitudinal:
 - `participant_id`, `age`, `sex`, `age`, and `group` MUST be reported in the `participants.tsv` file
- Age MUST BE REPORTED in months to account for cases with multiple acquisition sessions in a given year
- All columns (except `participant_id`) MUST BE DESCRIBED in the `participants.json` / `session.json`

SOME OTHER RECOMMENDATIONS

- **Never keep the acquisition date in the session label:** this might enable the identification of the scanned individual even if MRI was anonymized or de-identified
- **Keep the README of the dataset the most informative:** it is the first information a user will have access to.
- **De-identify MRI with a defacing tool** such as [pydeface](#) if the intent is to release the dataset publicly.

Please check [this paper](#) (Athena E. Theyers et al, 2021) for a comparison of multiple defacing tools.

- **Install bids-validator locally if the dataset is big;** the online validator is very slow. See the [bids-validator Github repo](#) for more details.