# BIDS: My data organization is as good as yours ULiège Open Science Day

Christophe Phillips<sup>1</sup> Nikita BELIY<sup>1</sup>

<sup>1</sup>GIGA CRC, Liège University, Belgium

November 6, 2024







Introduction I

Introduction I: Data acquisition

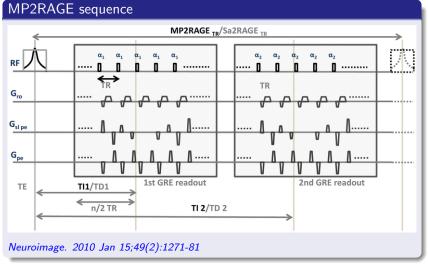
## MRI acquisition

- MRI principles
  - $B_0$  Proton spin alignment along z-axis
  - Repeated RF pulses to move proton spins out of equilibrium
  - Disturbed protons generate a electro-magnetic field used to reconstruct image
- Pulse parameters and sequences define different modalities
  - anatomical images
  - functional images (time-dependent blood oxygenation)
  - diffusion images (water molecules)
- Acceleration and noise suppression techniques add complexity

Knowledge of parameters needed to interpret images



## MRI acquisition



### DICOM data

Too much metadata

https://www.dicomstandard.org/current

- Created in 80s
- File format and TCP/IP protocols
- Common format for different manufacturers
- Format of out-of scanner raw image

### **Difficulties**

- Image and metadata manipulation
- Hundreds of defined metadata fields
- Manufacturer private fields
- Protocol private fields

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- Manufacturer private fields
- Protocol private fields
- Cryptic fields

### NIfTI

Not enought metadata

https://nifti.nimh.nih.gov/

### A (Sort of) New Image Data Format Standard: NIfTI-1

Robert W Cox1, John Ashburner2, Hester Breman3, Kate Fissell4, Christian Haselgrove5, Colin J Holmes6, Jack L Lancaster7, David E Rex8, Stephen M Smith9, Jeffrey B Woodward10, Stephen C Strother11 1SSCC/NIMH/NIH/DHHS/Bethesda, 2FIL/London, 3Brain Innovation/Maastricht, 4U Pittsburgh/Pittsburgh, 5MGH/Charlestown. SGI/Mountain View, 7RIC/UTHSCSA/San Antonio, 8LONI/UCLA/Los Angeles, 9FMRIB/Oxford, 19Dartmouth College/Hanover. 11U Minnesota/Minneapolis and NIfTI-DFWG Chair

- NIfTI = Neuroimaging Informatics Technology Initiative NIH-sponsored working group to promote interoperability of functional neuroimaging software tools
- DFWG = Data Format Working Group within NIfTI to deal with data interoperability
  - e.g., make it easier to interchange image (etc.) data between analysis packages
- Near-term efforts: extend ANALYZE™-7.5 file format (.hdr/.img file pairs) to add features the DFWG agreed were highly desirable for FMRI analysis = the NIfTI-1 format New features fit into unused/little-used ANALYZE fields

#### **Current Status**

- DFWG has approved NIfTI-1 format
  - Major software packages (AFNI, BrainVoyager, FSL, SPM) agree to read NIfTI-1 files by July 31, 2004 and to be able to write them by Dec 31, 2004
  - NIfTI-1 specification is in the form of a very heavily commented C header file, laving out the fields and their interpretations:

http://nifti.nimh.nih.gov/dfwg/

#### **OHBM 2004**

- (Almost) no acquisition metadata stored
- Metadata dumped at conversion into JSON file ( ) SON f



Introduction II: Data usage

## Naive data organisation

### CRC experience

- Subject's files in one folder
  - Not always
- Can be thousands of files in folder
  - In complex studies
- Protocols identified by "Series number"
  - Unreliable and unclear
- Selection by regexp or manually
  - Error prone
- Processing output in the same folder (sometimes)

```
-0004-00001-000001-01.ison
-0004-00001-000001-01.nii
-0004-00002-000002-01.json
-0004-00002-000002-01.nii
-0004-00003-000003-01.ison
-0004-00003-000003-01.nii
-0004-00004-000004-01.json
-0004-00004-000004-01.nii
-0004-00005-000005-01.json
-0004-00005-000005-01.nii
-0004-00006-000006-01.json
0004 00006-000006-01.nii
0004 00007-000007-01.json
0004 00007-000007-01.nii
0004 00008-000008-01.json
0004 00008-000008-01.nii
0004 00009-000009-01.json
0004 00009-000009-01.nii
0004 00010-000010-01.json
0004 00010-000010-01.nii
0005 00001-000001-01.json
0005 00001-000001-01.nii
0005 00002-000002-01.json
0005 00002-000002-01.nii
0005 00003-000003-01.json
```



## Sharing data

Running own pipeline on foreign data

- Dataset under their hand
- No or minimal effort to organize/document
- Often raw data without any organization
- Difficulty to obtain additional information

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### Personal experience (pathological cases)

- Non-organized data
  - Just one big folder with all files
- Converted data without any metadata
- Different protocol from what is claimed



## Sharing code

Running foreign pipeline on own data

- Between colleagues
  - Similar dataset structure with small adjustments
  - Low quality code for personal use
  - Little to no documentation
  - Hard-coded paths and file names in surprising places
  - Cryptic metadata retrieval even author don't remember what and why
  - Hard-coded metadata

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  - Little to no documentation
  - Hard-coded paths and file names in surprising places
  - Cryptic metadata retrieval even author don't remember what and why
  - Hard-coded metadata
- Between institutions/open source
  - Some effort on documentation
  - Limited or no hard-coded paths
  - Issues/bugs follow up
  - Maybe incompatible data structure



## BIDS:

Brain Imaging Data Structure

## BIDS

### Brain Imaging Data Structure

https://bids.neuroimaging.io

- Community effort
  - Started in 2015
  - Current version 1.10.0

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## The brain imaging data structure, a format for organizing and describing outputs of neuroimaging experiments

Krzysztof J. Gorgolewski <sup>Cl.</sup>, Tibor Auer, Vince D. Calhoun, R. Cameron Craddock, Samir Das, Eugene P. Duff, Guillaume Flandin, Satrajit S. Ghosh, Tirstan Glatard, Yaroslav Cl. Halchenko, Daniel A. Handwerker, Michael Hanke, David Keator, Xiangrui Li, Zachary Michael, Camille Maumet, B. Nolan Nichols, Thomas E. Nichols, John Pellman, Jean-Bagtiste Poline, Ariel Rokem, Gunnar Schaefer, Yanessa Sochat, William Irriplett, ... Russell A. Poldrack + Show authors

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- Community effort
  - Started in 2015
  - Current version 1.10.0
- Human readable
  - Minimized curation
  - Error reduction
- Computer readable
  - Optimized usage of data analysis software
  - Development of automated tools

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## File naming rules

**Findable** 

GitHub:bids-examples

```
sub-01
ses-1
sub-01 ses-1
sub-01 ses-1
task-rest acq-fullbrain un-1
sub-01 ses-1
task-rest acq-fullbrain un-2
sub-01 ses-1
task-rest acq-fullbrain un-2
sub-01 ses-1
task-rest acq-fullbrain un-2
sub-01 ses-1
task-rest acq-prefrontal
bold.nii.gz
sub-01 ses-1
task-rest acq-prefrontal
physio.tsv.gz
```

### Directories:

- sub-<label>: per subject
- *ses-<label>*: per session (optional)
- < data type>: group of different types of data

### Names:

- <suffix>: defines modality ("kind" of image)
- < entity>-< label>: defines acquisition parameters of image

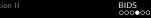


### Metadata definitions

#### Interoperable

Key name	Requirement Level	Data type	Description
EchoTime	RECOMMENDED, but	number or	The echo time (TE) for the acquisition
	REQUIRED if	array of	specified in seconds. Corresponds to
	corresponding	numbers	DICOM Tag 0018, 0081 Echo Time
	fieldmap data is		(please note that the DICOM term is in
	present, or the data		milliseconds not seconds). The data

- Stored in JSON file
- Strict definition: conventions and units
- Requirement levels:
  - REQUIRED: needed to interpret data
  - RECOMMENDED: will improve interpretation
  - OPTIONAL: might be useful



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### Sidecar JSON file

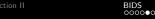
```
"CogAtlasID": "https://www.cognitiveatlas.org/id/trm_4c8a834779883",
"EchoTime": 0.017,
"EffectiveEchoSpacing": 0.0003333262223739227,
"PhaseEncodingDirection": "j-",
"RepetitionTime": 3.0,
"SliceEncodingDirection": "k",
```



## Modality agnostic (top-level) files

#### Reusable

- dataset\_description.json
  - Dataset name, BIDS version, authors, DOI, etc...
- README(.md, .txt, .rts)
  - Free text detailed description and notes on dataset
- CITATION.cff/CHANGES/LICENSE



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#### participants.tsv - participants description participant id age number handedness sub-01 29 17 100 sub-02 100 23 sub - 03 25 18 86 sub-04 26 100

## Expandable to other data types

**BIDS Expansion Proposals** 

https://bids.neuroimaging.io/get\_involved.html

- Magnetoencephalography (MEG) 2018
   Sci Data 5, 180110 (2018)
- Electroencephalography (EEG/iEEG) 2019
   Sci Data 6, 103 (2019), Sci Data 6, 102 (2019)
- Positron emission tomography (PET) 2022
   Sci Data 9, 65 (2022)
- Quantitative MRI (qMRI) 2022
   Sci Data 9, 517 (2022)
- Microscopy 2022
   Front Neurosci, 16 (2022)
- Near-Infrared Spectroscopy (NIRS) 2023
   PsyArXiv. doi:10.31219/osf.io/7nmcp
- Magnetic Resonance Spectroscopy (MRS) 2024 (publication forthcoming)



Conclusion: Why it works

### Human perspective

### Worst thing ever!

- Considerable effort to organize data
- Sometimes confusing and contradictory descriptions
- Need to integrate all acquisition data

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### Best thing ever!

- Easy to retrieve information
- Easy to run pipelines

### Computer perspective

### Best thing ever!

- Easy to retrieve data and metadata
  - bids-matlab, pybids query based data retrieval
- Easy to patch errors
- Easy to write pipelines
  - qmri, fmriprep BIDS-based preprocessing pipelines
- Modular composition (BIDS in, BIDS out)

## Computer perspective

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- Easy to retrieve data and metadata
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### Worst thing ever!

- Rare case of non-BIDS metadata
- Cases of modalities not included in BIDS
- No strict regulation of pipeline outputs (derivatives)



## Thanks for your attention!