



# **What is the Brain Imaging Data Structure and why you should know about this!**

16th May 24 – Open Science Series

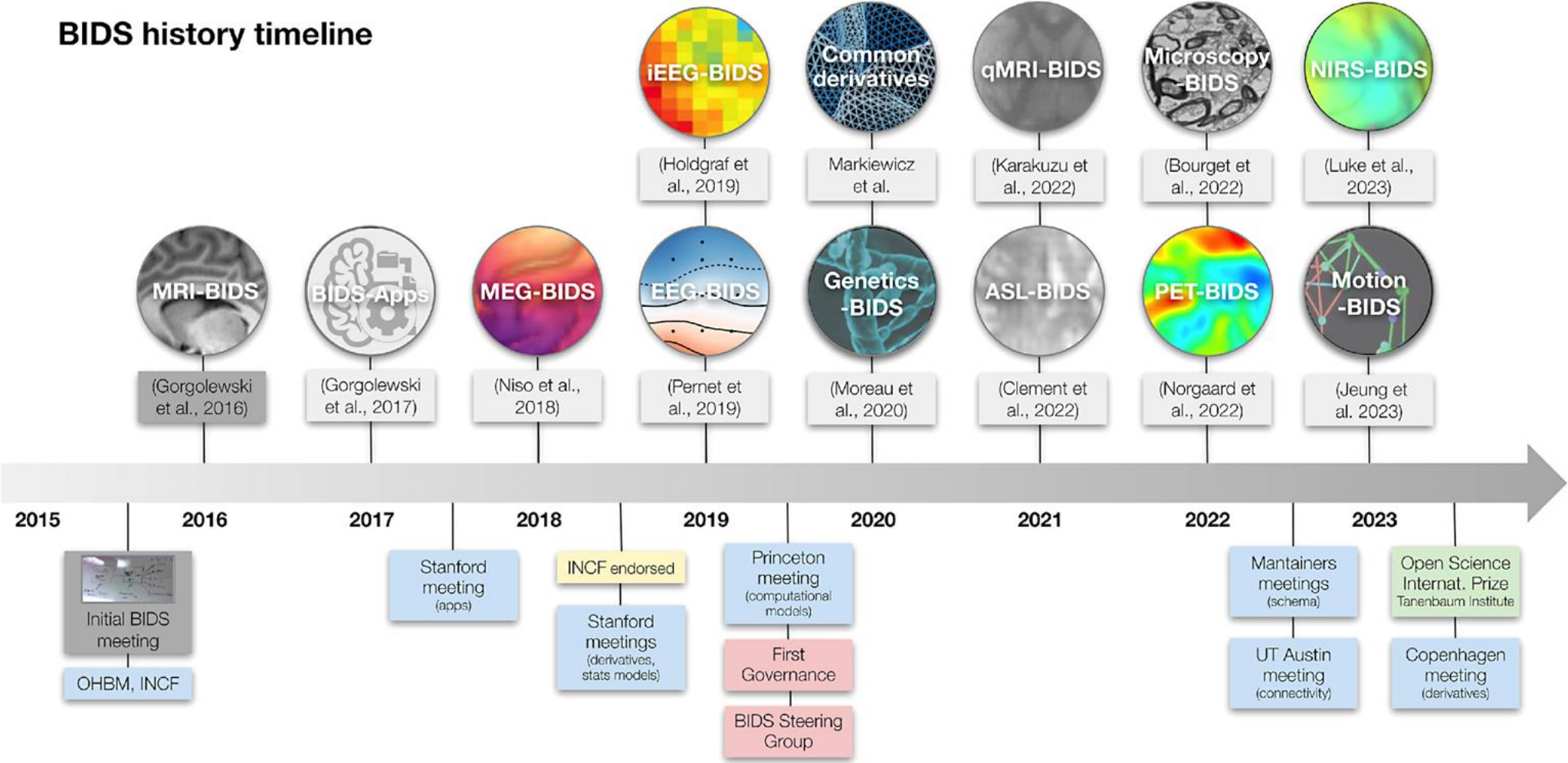


```
|— README.md
|— dataset_description.json
|— participants.json
|— participants.tsv
└─ sub-001
    |— ses-01
        |— eeg/
        └─ motion/
    └─ ses-02
└─ sub-002
```

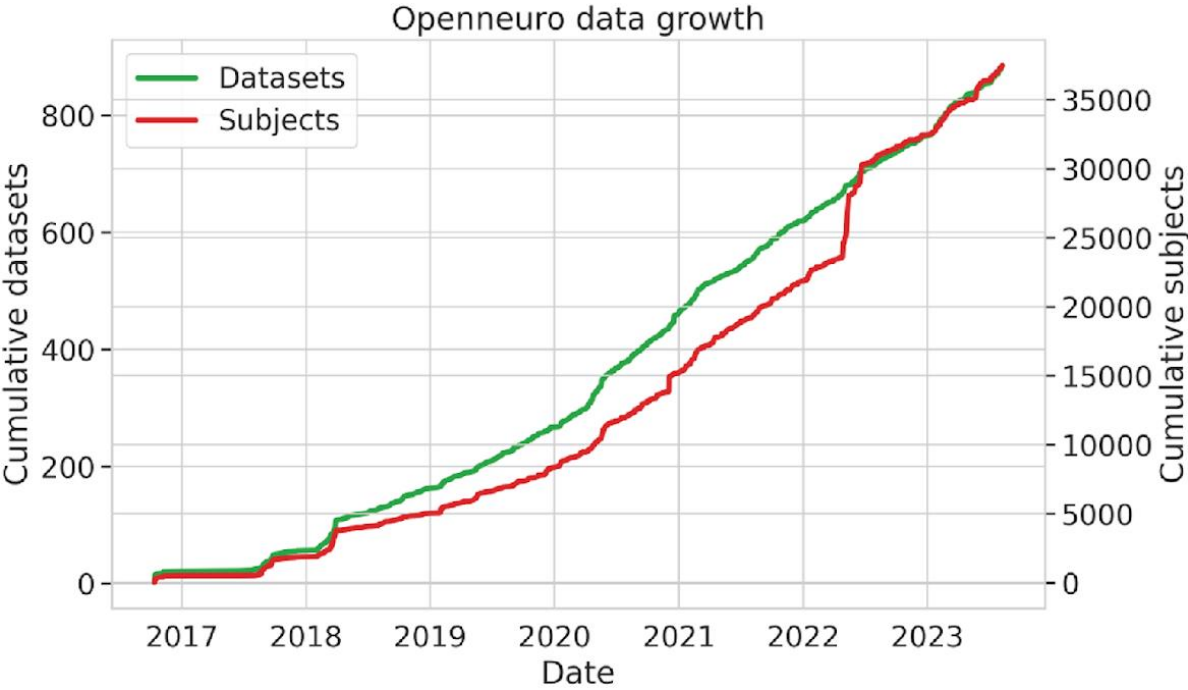
# **What is the Brain Imaging Data Structure** and why you should know about this!

# The Past, Present, and Future of BIDS

BIDS history timeline



# The Past, Present, and Future of BIDS



**OpenNeuro EEG**

OpenNeuro added support for EEG datasets in 2019 when EEG was incorporated into the BIDS standard.

**7.926** Participants      **200** Public Datasets

## Search EEG Portal

Search at the participant-level with Neurobagel ?

These filters return **70** datasets:

Keywords ?

KEYWORD:

MODALITY:



gait X

EEG X

# What is BIDS?



About 20+ different EEG systems  
record in different file formats





**Remi Gau**

Remi-Gau

Follow

I wanted to understand the brain, now I tell people how to name files, folders and variables.

# What is BIDS?



- BIDS is based on simple file formats and folder structures

## DATA STRUCTURE

- Which file formats to use, BIDS is **NOT** a file format
- Naming convention for files and directories

## METADATA

- Prevents metadata getting lost
- Some metadata is better than no metadata (80/20 rule)
- Stored in **json** files, readable by both humans and machines

A screenshot of the Brain Imaging Data Structure v1.9.0 website. The page has a dark blue header with the title "Brain Imaging Data Structure v1.9.0" and a search bar. The main content area is white and features a navigation menu on the left with items like "The BIDS Specification", "Introduction", "Common principles", etc. The main text area is titled "The Brain Imaging Data Structure" and contains a large, bold heading "JavaScript Object Notation" with a code block showing a JSON object: 

```
{  
  "key": "value",  
  "name": "Julius",  
}
```

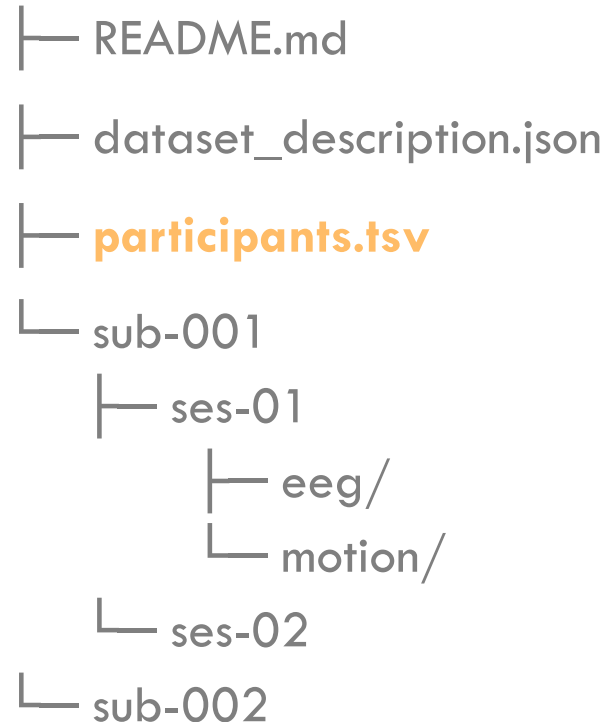
 Below the code block, there is a link "also be downloaded as PDF." and some partially visible text on the right side of the page.



# Requirement levels

Level of requirement **for** files and **within** files:

1. **REQUIRED**
2. **RECOMMENDED**
3. **OPTIONAL**



participant_id	age	group
sub-001	34	C
sub-002	12	P
sub-003	33	M

## Dataset info



```
├─ README.md
├─ dataset_description.json
├─ participants.json
├─ participants.tsv
├─ sub-001
│   └─ ses-01
│       ├── eeg/
│       └─ motion/
├─ ses-02
└─ sub-002
```

# Modality agnostic data

# dataset\_description.json

```
|— README.md
|— dataset_description.json
|— participants.json
|— participants.tsv
├— sub-001
  |— ses-01
  └— ses-02
└— sub-002
```



```
{
  "Name": "The mother of all experiments",
  "BIDSVersion": "1.4.0",
  "DatasetType": "raw",
  "License": "CC0",
  "Authors": [ "Paul Broca", "Carl Wernicke" ],
  "Funding": [ „NIH F37823MFH1" ],
  "EthicsApprovals": [„HRPO (Protocol AR0928" ],
  "DatasetDOI": "10.0.2.3/dfjj.10"
}
```

# participants

- └ README.md
- └ dataset\_description.json
- └ **participants.tsv**
- └ **participants.json**
- └ sub-001
  - └ ses-01
  - └ ses-02
- └ sub-002

## participants.tsv

<b>participant_id</b>	<b>age</b>	<b>group</b>
sub-001	34	C
sub-002	12	P
sub-003	33	M

## participants.json

```
"age": {  
  "Description": "age of  
the participant",  
  "Units": "years"  
},  
"group": {  
  "Description": "assigned  
group",  
  "Levels": {  
    "C": "control",  
    "M": "medication",  
    "P": "placebo",  
  }  
}
```

## Dataset info

### EEG



```
├─ README.md
├─ dataset_description.json
├─ participants.json
├─ participants.tsv
├─ sub-001
│   └─ ses-01
│       ├── eeg/
│       └─ motion/
├─ ses-02
└─ sub-002
```

# Modality specific data

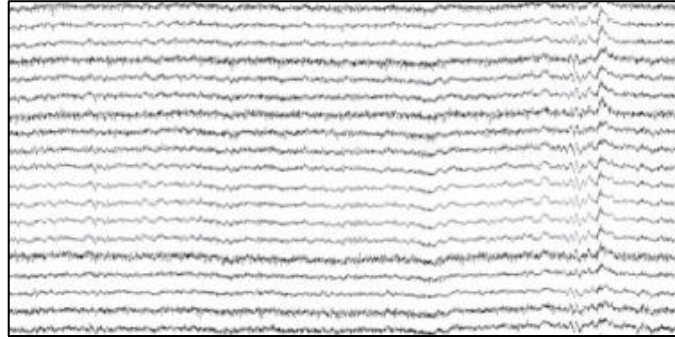
# Directories and file naming structure

- Data for each subject are organized in subdirectories labeled "**sub-<label>**", with "<label>" representing the unique identification label of each subject
- If subjects participated in multiple sessions, subdirectories labeled "**ses-<label>**" contain session-specific data within the subject directory
- Each session subdirectory (or the subject subdirectory if no session data exists) contains subdirectories for different data types, only defined if files are present for that type
- For a data file that was collected in a given **session** from a given **subject**, the file name **MUST** begin with the string **sub-<label>[\_ses-<label>]<sub>task-<label></sub>**

```
├─ README.md
├─ dataset_description.json
├─ participants.json
├─ participants.tsv
├─ sub-001
│   └─ ses-01
│       └─ eeg/
│           └─ sub-001_ses-01_task-GoNoGo.eeg
```

# Files | eeg

└─ sub-001  
└─ ses-01  
└─ eeg/



```
{  
  "TaskName": "TASKNAME",  
  "SamplingFrequency": 1000,  
  "SoftwareFilters": "n/a",  
  "EEGChannelCount": 4,  
  "EOGChannelCount": 1,  
  "EEGReference": "placed on Cz",  
  "PowerLineFrequency": 50  
}
```

└─ sub-001\_ses-01\_task-<label>\_eeg.<extension>

└─ sub-001\_ses-01\_task-<label>\_eeg.json

└─ sub-001\_ses-01\_task-<label>\_channels.tsv

└─ sub-001\_ses-01\_task-<label>\_electrodes.tsv

└─ sub-001\_ses-01\_task-<label>\_electrodes.tsv

name	type	units	status	status_description
CP5	EEG	microV	good	n/a
FC5	EEG	microV	bad	high freq noise
FC1	EEG	microV	good	n/a
C3	EEG	microV	good	n/a
VEOG	EOG	microV	good	n/a

name	x	y	z	impedance
CP5	-0.77	-0.30	0.57	8
FC5	-0.77	0.30	0.57	12
FC1	-0.29	0.31	0.91	2
C3	-0.59	0.00	0.81	5
VEOG	n/a	n/a	n/a	n/a

```
{  
  "EEGCoordinateSystem": "T1w",  
  "EEGCoordinateUnits": "mm",  
  "AnatomicalLandmarkCoordinates": {  
    "LPA": [-0.067, 1.736e-09, -3.844e-09],  
    "NAS": [-4.11e-09, 0.091, -4.541e-10],  
    "RPA": [0.064, -6.435e-09, -4.566e-09]  
  },  
  "AnatomicalLandmarkCoordinateSystem": "T1w",  
  "AnatomicalLandmarkCoordinateUnits": "mm",  
  "IntendedFor": "sub-01_T1w.nii.gz"  
}
```

# Metadata | eeg.json

## REQUIRED

TaskName, SamplingFrequency,  
PowerlineFrequency, EEGReference,  
SoftwareFilters

## RECOMMENDED

TaskDescription, Manufacturer,  
EEGChannelCount,  
EOGChannelCount, ...

## OPTIONAL

-

```
{  
  TaskName: Seeing stuff,  
  SamplingFrequency: 2400,  
  Manufacturer: Brain Products,  
  ManufacturersModelName: BrainAmp DC,  
  EEGChannelCount: 87,  
  EOGChannelCount: 2,  
  TriggerChannelCount: 1,  
  PowerLineFrequency: 50,  
  EEGPlacementScheme: 10 percent system,  
  EEGReference: single electrode placed on FCz,  
  SoftwareFilters: {  
    Anti-aliasing filter: {  
      half-amplitude cutoff (Hz):  
      500,  
      Roll-off: 6dB/Octave  
    }  
  }  
}
```



# Metadata | channels.tsv

## REQUIRED

name, type, units

## RECOMMENDED

-

## OPTIONAL

description, sampling\_frequency

<b>name</b>	<b>type</b>	<b>units</b>	<b>description</b>	<b>reference</b>	<b>status</b>	<b>status_description</b>
VEOG	VEOG	uV	left eye	VEOG-, VEOG+	good	n/a
FDI	EMG	uV	left first dorsal interosseous	FDI-, FDI+	good	n/a
Cz	EEG	uV	n/a	REF	bad	high frequency noise
UADC001	MISC	n/a	envelope of audio signal	n/a	good	n/a

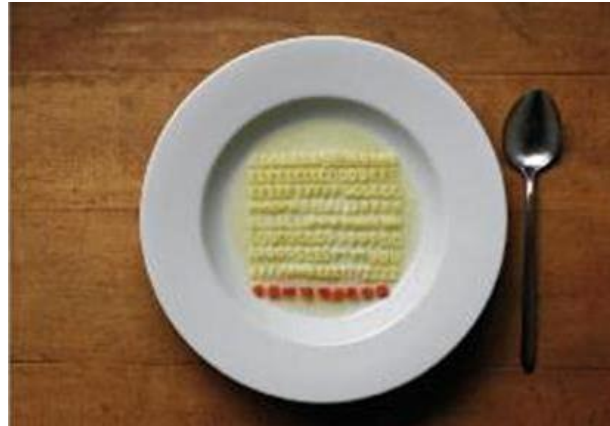
# Metadata | electrodes.tsv

name	x	y	z	type	material	color
VEOG+	n/a	n/a	n/a	cup	Ag/AgCl	blue
VEOG-	n/a	n/a	n/a	cup	Ag/AgCl	white
FDI+	n/a	n/a	n/a	cup	Ag/AgCl	red
FDI-	n/a	n/a	n/a	cup	Ag/AgCl	red
GND	-0.07	0.00	-0.070	clip-on	Ag/AgCl	pink
Cz	0.00	0.07	0.06	cup	Ag/AgCl	yellow
REF	-0.07	-0.02	-0.01	cup	Ag/AgCl	grey

Column name	Requirement Level	Data type	Description
name	REQUIRED	string	Name of the electrode contact point. Values in <code>name</code> MUST be unique. This column must appear <b>first</b> in the file.
x	REQUIRED	number	Recorded position along the x-axis. This column must appear <b>second</b> in the file.
y	REQUIRED	number	Recorded position along the y-axis. This column must appear <b>third</b> in the file.
z	REQUIRED	number or "n/a"	Recorded position along the z-axis. This column must appear <b>fourth</b> in the file.
type	RECOMMENDED	string	Type of the electrode (for example, cup, ring, clip-on, wire, needle). This column may appear anywhere in the file.
material	RECOMMENDED	string	Material of the electrode (for example, Tin, Ag/AgCl, Gold). This column may appear anywhere in the file.
impedance	RECOMMENDED	number	Impedance of the electrode, units MUST be in kOhm. This column may appear anywhere in the file.
<b>Additional Columns</b>	OPTIONAL	n/a	Additional columns are allowed if they are defined in the associated metadata file.

## Dataset info

### Motion



```
├─ README.md
├─ dataset_description.json
├─ participants.json
├─ participants.tsv
├─ sub-001
│   └─ ses-01
│       └─ eeg/
│           └─ motion/
├─ ses-02
└─ sub-002
```

# Modality specific data

# Files | motion

```
└─ sub-001/  
  └─ ses-01/  
    └─ eeg/  
      ...  
    └─ motion/
```

```
└─ sub-001_ses-01_task-<label>_tracksys-<label>_motion.tsv  
└─ sub-001_ses-01_task-<label>_tracksys-<label>_motion.json  
└─ sub-001_ses-01_task-<label>_channels.tsv
```

0,2634511	0,092295	0,0086682	0,9305117	0,690106	0,8098815
0,694520	0,1918243	0,8437273	0,3975710	0,885496	0,8952724
0,0766395	0,2587211	0,5434792	0,2822837	0,2789791	0,2326254
0,5779993	0,0456141	0,0490745	0,9408899	0,1533421	0,6683652
0,054556	0,7915927	0,5871733	0,4669577	0,9754468	0,0480541
0,966024	0,1962834	0,7114406	0,3389448	0,7194495	0,4384892
0,984172	0,5079461	0,1180168	0,7966978	0,1753768	0,4886533
0,9883907	0,1557346	0,8002013	0,6334882	0,7526906	0,8529441
...	...	...	...	...	...

```
{  
  "SamplingFrequency": 60,  
  "SamplingFrequencyEffective": 60.19,  
  "TaskName": "BIDS Motion fictive example",  
  "TrackingSystemName": "imu1",  
  "TaskDescription": "walking and talking",  
  "MotionChannelCount": 6,  
  "SubjectArtefactDescription": "n/a",  
  "TrackedPointsCount": 2,  
  "ACCELChannelCount": 3,  
  "GYROChannelCount": 3,  
  "Manufacturer": "BWSensing",  
  "ManufacturersModelName": "BW-imu600",  
}
```

name	component	type	tracked_point	units	placement
imu1_rf_acc_x	x	ACCEL	rf	m/s <sup>2</sup>	right_foot
imu1_rf_acc_y	y	ACCEL	rf	m/s <sup>2</sup>	right_foot
imu1_rf_acc_z	z	ACCEL	rf	m/s <sup>2</sup>	right_foot
imu1_rf_gyro_x	x	GYRO	rf	rad/s	right_foot
imu1_rf_gyro_y	y	GYRO	rf	rad/s	right_foot
imu1_rf_gyro_z	z	GYRO	rf	rad/s	right_foot

# Metadata | motion.json

## REQUIRED

TaskName, SamplingFrequency

## RECOMMENDED

TaskDescription, RecordingDuration,  
MotionChannelCount, <type>ChannelCount,  
SoftwareFilters, ...

## OPTIONAL

Manufacturer, RecordingSoftware, ...

```
{  
  "SamplingFrequency": 60,  
  "TaskName": "BIDS Motion fictive example",  
  "TrackingSystemName": "IMU Right Hand",  
  "TaskDescription": "walking and talking",  
  "InstitutionAddress": "Fictive address",  
  "InstitutionName": "Fictive Institution",  
  "MotionChannelCount": 18,  
  "RecordingDuration": 4667.641106,  
  "SubjectArtefactDescription": "n/a",  
  "TrackedPointsCount": 2,  
  "ACCELChannelCount": 6,  
  "GYROChannelCount": 6,  
  "MAGNChannelCount": 6,  
  "Manufacturer": "BWSensing",  
  "ManufacturersModelName": "BW-IMU600",  
}
```

# Metadata | channels.tsv

## REQUIRED

name, component, type, tracked\_point, units

## RECOMMENDED

placement, reference\_frame

## OPTIONAL

status, status\_description, sampling\_frequency

### Restricted keyword list for channel type

Restricted keyword list for column `type` in alphabetic order. Note that upper-case is REQUIRED:

Keyword	Description
ACCEL	Accelerometer channel, one channel for each spatial axis. Column component for the axis MUST be added to the *_channels.tsv file (x, y, or z).
ANGACCEL	Angular acceleration channel, one channel for each spatial axis. Column component for the axis MUST be added to the *_channels.tsv file (x, y, or z).
GYRO	Gyrometer channel, one channel for each spatial axis. Column component for the axis MUST be added to the *_channels.tsv file (x, y, or z).
JNTANG	Joint angle channel between two fixed axis belonging to two bodyparts. Angle SHOULD be defined between proximal and distal bodypart in deg.

name	component	type	tracked_point	units	reference_frame
t1_acc_x	x	ACCEL	LeftFoot	m/s <sup>2</sup>	global
t1_acc_y	y	ACCEL	LeftFoot	m/s <sup>2</sup>	global
t1_acc_z	z	ACCEL	LeftFoot	m/s <sup>2</sup>	global
t1_gyro_x	x	GYRO	LeftFoot	rad/s	global
t1_gyro_y	y	GYRO	LeftFoot	rad/s	global
t1_gyro_z	z	GYRO	LeftFoot	rad/s	global
...					

## Dataset info

### Events



```
├─ README.md
├─ dataset_description.json
├─ participants.json
├─ participants.tsv
├─ sub-001
│   ├── ses-01
│   │   ├── eeg/
│   │   └─ motion/
│   └─ ses-02
├─ sub-002
```

# Modality specific data

# Task events

sub-<label>/

<data\_type>/

<matches>\_events.tsv

<matches>\_events.json

```
{
  "trial_type": {
    "LongName": "Event category",
    "Description": "Indicator of type of action that is expected",
    "Levels": {
      "start": "A red square is displayed to indicate starting",
      "stop": "A blue square is displayed to indicate stopping"
    }
  },
  "channel": {
    "Description": "Channel(s) associated with the event"
  },
  "annots": {
    "LongName": "Annotations",
    "Description": "Annotations associated with channels indicated in the channel column.",
    "Levels": {
      "musc": "Muscle artifact. A very common, high frequency, sharp artifact that corresponds with agitation/nervousness in a patient."
    }
  }
}
```

onset	duration	trial_type	response_time	stim_file	channel	annots
1.23	0.65	start	1.435	red_square.jpg	n/a	n/a
5.65	0.65	stop	1.739	blue_square.jpg	n/a	n/a
12.1	2.35	n/a	n/a	n/a	Cz	musc







What is the Brain Imaging  
Data Structure and **why you  
should know about this!**

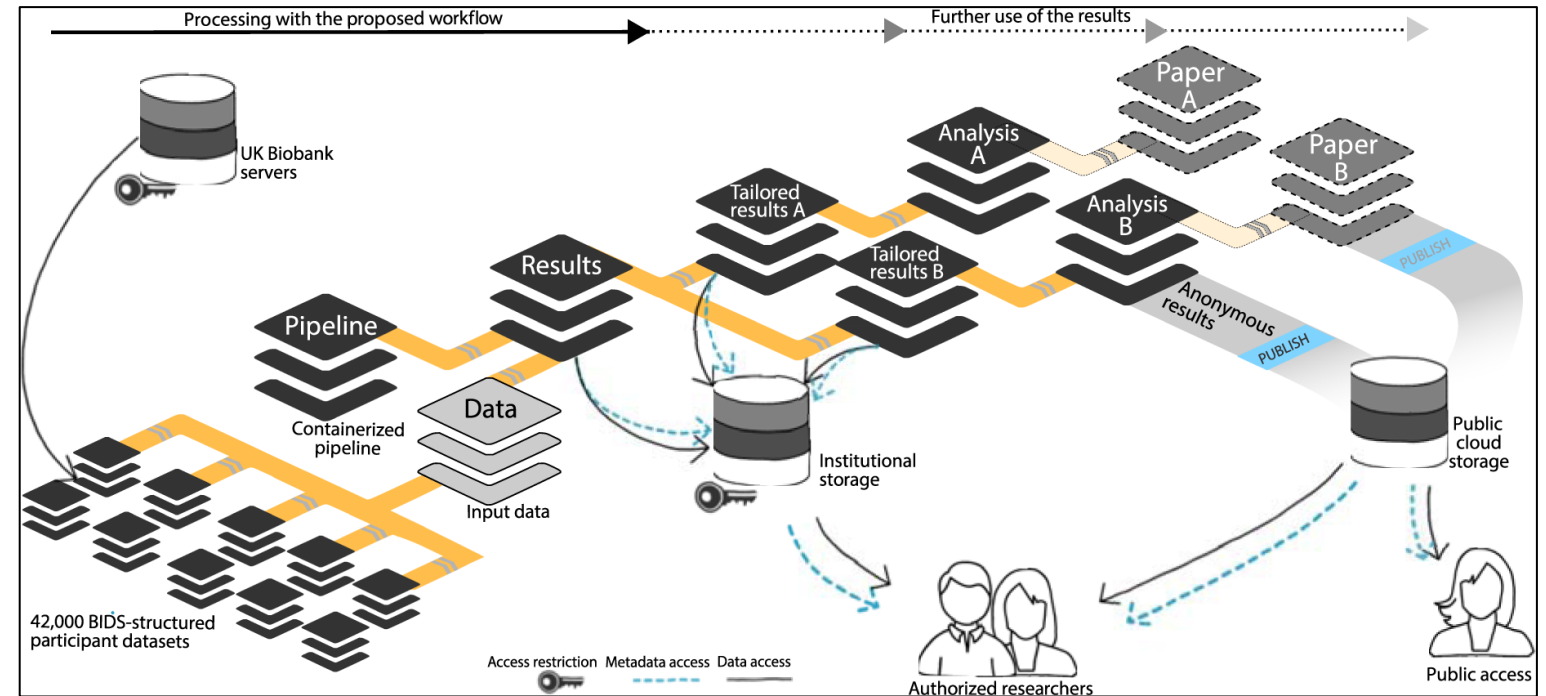
# For research

- Validation and optimizing pipelines

- Building a sustainable research environment

Article | [Open access](#) | Published: 09 February 2023  
**EEG is better left alone**  
[Arnaud Delorme](#)   
[Scientific Reports](#) **13**, Article number

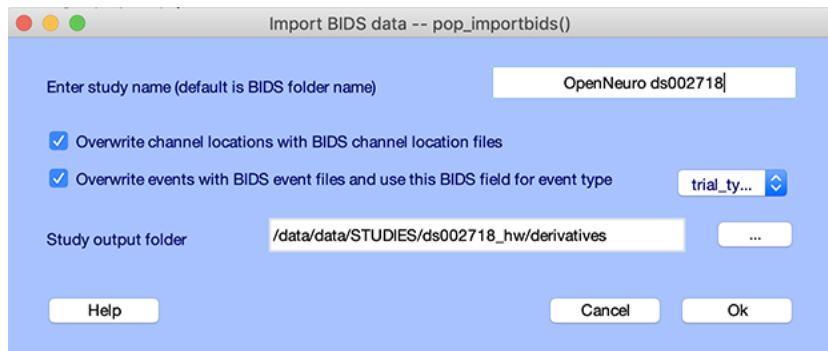
**Preprocessing Choices for P3 Analyses with Mobile EEG: A Systematic Literature Review and Interactive Exploration**  
 Nadine S. J. Jacobsen,  Daniel Kristanto, Suong Welp, Yusuf Cosku Incel,  Stefan Debener  
doi: <https://doi.org/10.1101/2024.04.30.591874>



# For you

## Load public data from the BIDS format

- EEGLab, FieldTrip and MNE provide importers



## Converting your data to BIDS

- Reuse data from your Lab and others
- Foster collaborations

```
sub = {'01', '02', '03', '04', '05', '06', '07', '08', '09', '10'};

% for subject 3 the age is unknown, for subject 2 the sex is not specified
age = [11 96 nan 77 82 87 18 40 26 80];
sex = {'f' [] 'f' 'f' 'f' 'm' 'm' 'm' 'm' 'm'};

for subindx=1:numel(sub)

    cfg = [];
    cfg.datatype = 'eeg';

    % Load your data
    cfg.dataset = sub{subindx} + '.edf';

    % specify the output directory
    cfg.bidsroot = 'bids';

    cfg.sub = sub{subindx};

    % specify the information for the participants.tsv file
    % this is optional, you can also pass other pieces of info
    cfg.participants.age = age(subindx);
    cfg.participants.sex = sex(subindx);

    % provide the mnemonic and long description of the task
    cfg.TaskName = 'changedetection';
    cfg.TaskDescription = 'Subjects were responding as fast as possible upon a
                           change in a visually presented stimulus.';

    % these are EEG specific
    cfg.eeg.PowerLineFrequency = 50; % since recorded in the EU
    cfg.eeg.EEGReference = 'M1'; % left mastoid

    data2bids(cfg);

end
```

# BIDS Starterpack

## Website

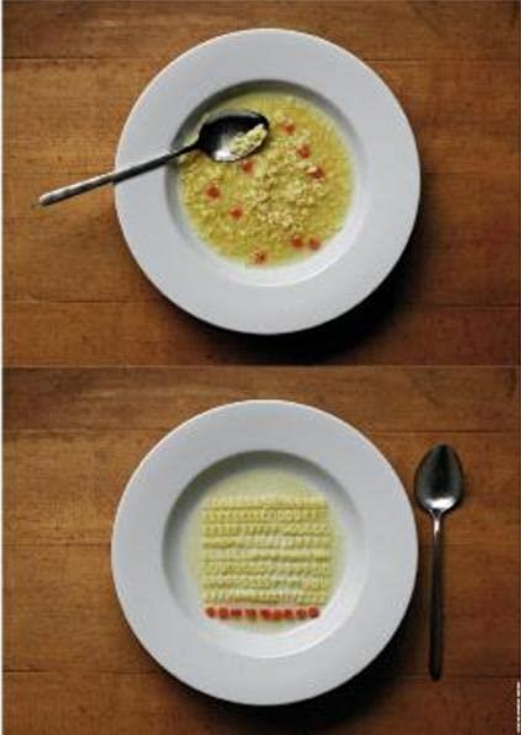
[<https://bids-specification.readthedocs.io/en/stable/>]

## Validator

[<https://bids-standard.github.io/bids-validator/>]

The screenshot shows the BIDS Validator v1.14.6 interface. At the top, there is a dark header with the text "BIDS Validator v1.14.6". Below this is a search bar with the placeholder text "Select a BIDS dataset to validate". The main navigation bar is blue and contains the text "Brain Imaging Data Structure v1.9.0" on the left, a search icon and "Search" in the center, and a GitHub logo with "v1.9.0", "255", and "153" on the right. The main content area is split into two columns. The left column is a table of contents with the following items: "Brain Imaging Data Structure v1.9.0", "The BIDS Specification", "Introduction", "Common principles", "Modality agnostic files", "Modality specific files", "Derivatives", "Longitudinal and multi-site studies", "Glossary", "BIDS Extension Proposals", "Appendix", "Changelog", "The BIDS Starter Kit", "Website", "Tutorials", and "GitHub repository". The right column is titled "The Brain Imaging Data Structure" and contains the following text: "The Brain Imaging Data Structure (BIDS) is a simple and intuitive way to organize and describe data.", "This document defines the BIDS specification, which provides many details to help implement the standard. It includes the core specification as well as many extensions to specific brain imaging modalities, and increasingly also to other kinds of data.", "If BIDS is new to you, and you would like to learn more about how to adapt your own datasets to match the BIDS specification, we recommend exploring the [BIDS Starter Kit](#). Alternatively, to get started please read [the introduction to the specification](#).", and "For an overview of the BIDS ecosystem, visit the [BIDS homepage](#). The entire specification can also be [downloaded as PDF](#)." At the bottom of the page, there is a dark footer with the text "Copyright © BIDS Contributors - CC BY 4.0" and social media icons for GitHub, Twitter, and LinkedIn.

# Conclusions



# Thank you for listening carefully



Thanks to **Sein Jeung** for pushing this to completion over the past three years  
Thanks to all of the **BIDS Maintainers and Devs** who help to implement this  
Thanks to my **working group**, who have given me the freedom to work in this  
Thanks to all **participants** who ever provided data which is now in BIDS

**Dataset info**



**Example**