



No country for old data: Increasing FAIR-ness of research outcomes through standardization and community-driven development

CSPD 2020

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- BRAMS

Member - BIDS, ReproNim, Brainhack, UNIQUE



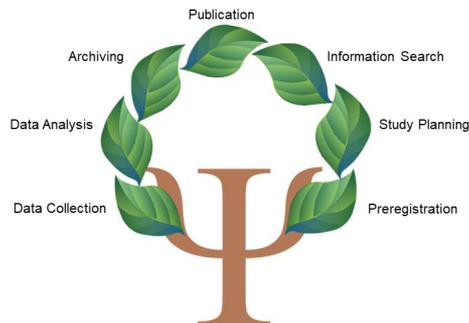
@peerherholz



A few things before we start ...

- to the folks with hearing impairments:
 - There will be live captions at the bottom of the screen. If something doesn't work, please let me/us know in the chat.
- to the folks with vision impairments:
 - All graphics (except those taken from publications) should be high in contrast and color-blind friendly. If there are problems, please let me/us know in the chat.
- to everyone:
 - I will try my best to speak loud and clearly. If you have troubles understanding me, please say so.
 - If you feel uncomfortable doing this/asking questions via the public chat, please send me a direct message.
- to everyone (continued):
 - I sincerely hope you feel comfortable indicating your pronouns, thus please add them to your name if you want to.
 - During the discussion/questions, please say your name as I (and others) would like to avoid mispronouncing it.
 - I try my best to present an objective overview. However, there's inherent bias to such presentations based on personal opinions and experience. Thus, I'm looking forward to openly discuss every point/comment with you all.
 - Let's all give our best to create an open, supportive and welcoming atmosphere for everyone.
- special thanks for pioneering work go to:
 - Kirstie Whitaker, Fernando Perez, Elizabeth DuPre, Gael Varoquaux

A few things before we start ...



Towards the sustainable use of
psychological research data

Who here has reused data?

If so, what kind?

A data sharing nightmare in three parts ... or “A data sharing carole”



<https://youtu.be/N2zK3sAtr-4>

It gets better! Does it really?

the classic

the solution?

Data availability

The data that support the findings of this study are available from the corresponding author upon reasonable request. **Code availability**

DATA AVAILABILITY STATEMENT

Code used to generate the analyses are available from the corresponding

The data that support the findings of this study are available from the corresponding author upon reasonable request.

Data Availability Statement

Go to: ☺

The datasets generated for this study are available on request to the corresponding author.

The screenshot shows the IBM SPSS Statistics Data Editor interface. The title bar reads "bcopen.sav [DataSet1] - IBM SPSS Statistics Data Editor". The menu bar includes File, Edit, View, Data, Transform, Analyze, Direct Marketing, Graphs, Utilities, Add-ons, Window, and Help. The toolbar contains various icons for file operations, data manipulation, and analysis. The main window displays a data grid with 20 rows and 11 columns. The first column is labeled "rowlabel" and contains row numbers 1 through 20. The second column is labeled "rowlabel" and contains the ID number 61302140. The remaining columns are labeled "sex", "age", "livharm1", "ethgrp2", "educat3", "work", "yrsarea", "resyrago", and "tenur". The data is as follows:

| rowlabel | rowlabel | sex | age | livharm1 | ethgrp2 | educat3 | work | yrsarea | resyrago | tenur |
|----------|----------|--------|-----|------------|----------------|----------------|------|----------------|----------|-------------|
| 1 | 61302140 | female | 36 | married | white | none | yes | 10 years b... | | buying i... |
| 2 | 61384060 | male | 44 | separated | white | apprentice... | yes | less than 1... | no | |
| 3 | 63684260 | male | 43 | married | white | o level/gcse | yes | 2 years but... | | buying i... |
| 4 | 63790220 | female | 27 | single | black or bl... | apprentice... | yes | less than 1... | no | |
| 5 | 63843180 | male | 38 | married | white | degree or d... | yes | 10 years b... | | buying i... |
| 6 | 71105020 | female | 18 | single | asian or as... | o level/gcse | yes | 10 years b... | | |
| 7 | 71105140 | male | 65 | single | asian or as... | degree or d... | yes | 20 years or... | | buying i... |
| 8 | 71106020 | male | 58 | divorced | white | none | no | 20 years or... | | |
| 9 | 71106060 | female | 38 | single | white | none | no | 2 years but... | no | |
| 10 | 71108100 | male | 29 | single | black or bl... | degree or d... | no | 3 years but... | | own it o... |
| 11 | 71110140 | male | 38 | cohabiting | white | degree or d... | yes | 3 years but... | | own it o... |
| 12 | 71110260 | male | 67 | single | white | none | yes | 10 years b... | | |
| 13 | 71114020 | male | 21 | single | white | apprentice... | yes | 20 years or... | | live her... |
| 14 | 71115060 | female | 37 | separated | black or bl... | o level/gcse | no | 5 years but... | | |
| 15 | 71119020 | female | 23 | single | mixed | degree or d... | yes | 3 years but... | | |
| 16 | 71119060 | male | 57 | married | chinese or ... | other | no | 20 years or... | | own it o... |
| 17 | 71119140 | female | 34 | single | black or bl... | apprentice... | no | 5 years but... | yes | |
| 18 | 71119180 | male | 55 | single | white | degree or d... | yes | 10 years b... | | buying i... |
| 19 | 71119220 | female | 27 | single | white | degree or d... | no | 2 years but... | | |
| 20 | 71119260 | female | 37 | divorced | white | none | no | 10 years b... | | buying i... |

https://bitly247.files.wordpress.com/2014/04/data-editor-screens.png

“upon (reasonable) request” =
you might get it if (a) can find it and (b) feel like it

if “data” is shared = here you go, have fun

The great baking analogy...

data as shared
by author



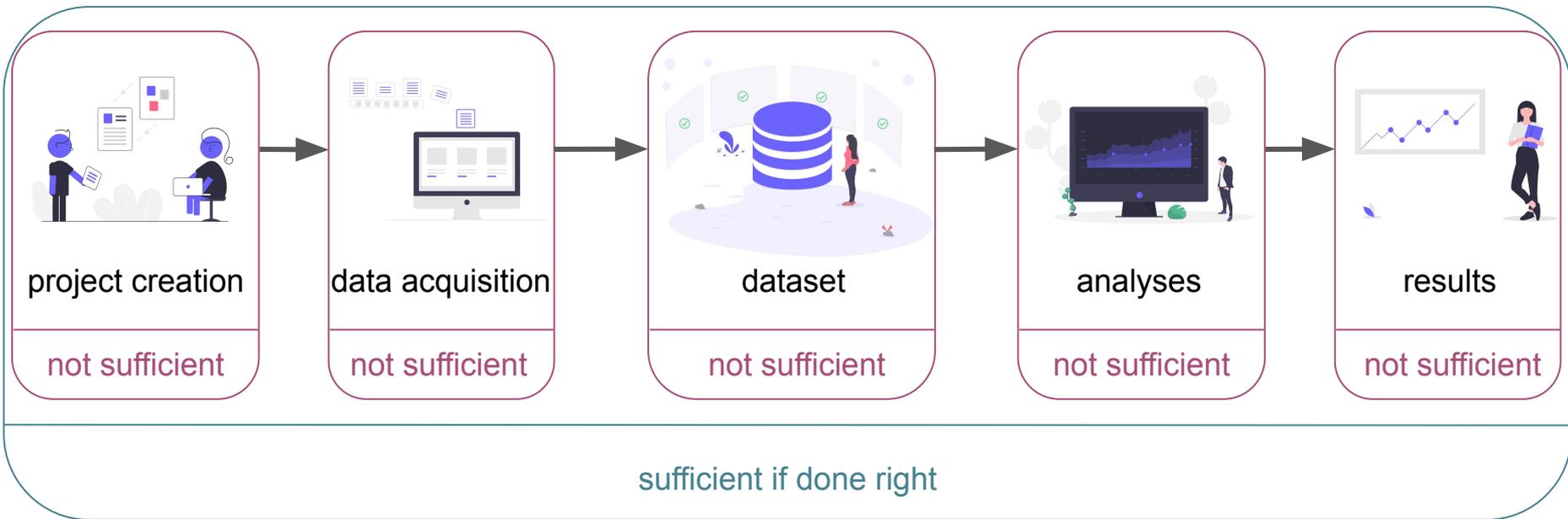
folks interested
in the data



data/results in
fancy journal



The great baking analogy...every ingredient AND the recipe are required



Please make sure to check <https://undraw.co/>

Findable Interoperable Accessible Reusable research outcomes

Findable

- data and metadata easily findable for both humans and machines (machine readable metadata *essential* for automatic discovery)

Accessible

- data can be easily obtained by humans as well as by machines, through well-defined and ideally standardized protocols
- conditions must be clearly established (license, reuse rights, etc.).
- when the data is inaccessible, the metadata should remain so in a way that the dataset remains discoverable

Interoperable

- possible to integrate different data into a single interface
- use of a common programming language, the use of controlled vocabulary, as well as non-proprietary formats
- can also be applied to metadata as well as references to other (meta)data

Re-Usable

- data and metadata following best practices -> optimized for reuse
- data and metadata must be richly detailed, as well as user licenses
- any information specific to the field of interest of the data in question must be made available as well.



<https://ogsi.ca/wp-content/uploads/Fair-rectangle-en.png>

- <https://www.force11.org/fairprinciples>
- <https://www.go-fair.org/fair-principles/>
- <https://www.andis.org.au/working-with-data/fairdata>
- <https://www.nature.com/articles/sdata201618.pdf>

Findable Interoperable Accessible Reusable research outcomes

- FAIR is *fuzzy per design*
- a *general set of principles* that should *guide* research *across domains*
- FAIR has *natural variation across domains* (e.g. metadata in Psychology or Neuroscience)
- each domain *needs* to develop its own FAIR *standards and terms*, as no solution can fit all



<https://ogsi.ca/wp-content/uploads/Fair-rectangle-en.png>

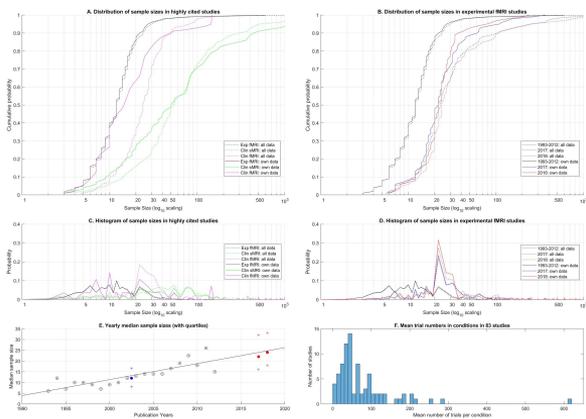
- <https://www.force11.org/fairprinciples>
- <https://www.go-fair.org/fair-principles/>
- <https://www.andis.org.au/working-with-data/fairdata>
- <https://www.nature.com/articles/sdata201618.pdf>

The four horsemen of ... unFAIR research outcomes

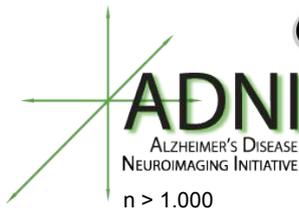
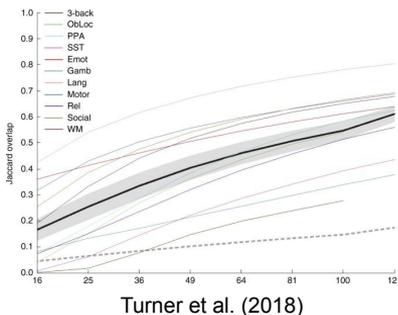
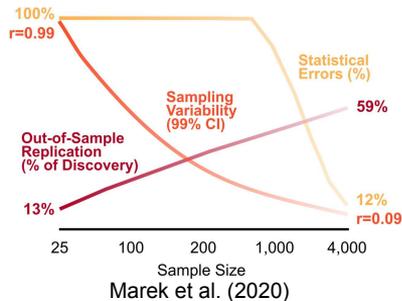


Source: Public domain

We need more data and we need it now, but FAIR



Szucs & Ioannidis (2020)



$n = 12.000$



CONNECTOME
COORDINATION FACILITY
 $n \sim 10.000$



| Category ID | Description | Items |
|-------------|--|-------|
| 1014 | Brain MRI | 922 |
| 1005 | Cognitive function summary | 5 |
| 1004 | Diet and alcohol summary | 321 |
| 1002 | Early life | 13 |
| 1007 | Education and employment | 16 |
| 1017 | Genomics | 78 |
| 100113 | Geographical and location | 13 |
| 1015 | Heart MRI | 39 |
| 1019 | Linked health outcomes | 75 |
| 1016 | Main abdominal MRI fields likely to be of interest to researchers. | 22 |
| 1018 | Mental health | 186 |
| 1006 | Physical measure summary | 66 |
| 1001 | Primary demographics | 8 |
| 1003 | Self-reported medical conditions | 118 |

$n = 40.000$

FAIR data through standardization ... with Marie Kondo



IBM SPSS Statistics Data Editor

| rowlabel | sex | age | lutham1 | ethgrp2 | educat3 | work | yearsa | resyago | tenur |
|----------|----------|--------|---------|-----------|----------------|---------------|--------|----------------|----------|
| 1 | 61302140 | female | 36 | married | white | none | yes | 10 years b. | buying |
| 2 | 61384060 | male | 44 | separated | white | appreciate... | yes | less than 1. | no |
| 3 | 63884800 | male | 43 | married | white | o level/gcse | yes | 2 years but... | buying |
| 4 | 63790220 | female | 27 | single | black or bl. | appreciate... | yes | less than 1. | no |
| 5 | 63843180 | male | 38 | married | white | degree or d. | yes | 10 years b. | buying |
| 6 | 71105020 | female | 18 | single | asian or as... | o level/gcse | yes | 10 years b. | buying |
| 7 | 71105140 | male | 65 | single | asian or as... | degree or d. | yes | 20 years or... | buying |
| 8 | 71106020 | male | 58 | divorced | white | none | no | 20 years or... | no |
| 9 | 71105060 | female | 38 | single | white | none | no | 2 years but... | no |
| 10 | 71108100 | male | 29 | single | black or bl. | degree or d. | no | 3 years but... | own it c |
| 11 | 71110140 | male | 38 | cohabbing | white | degree or d. | yes | 3 years but... | own it c |
| 12 | 71110260 | male | 67 | single | white | none | yes | 10 years b. | own it c |
| 13 | 71114020 | male | 21 | single | white | appreciate... | yes | 20 years or... | line her |
| 14 | 71115060 | female | 37 | separated | black or bl. | o level/gcse | no | 5 years but... | no |
| 15 | 71119020 | female | 23 | single | mixed | degree or d. | yes | 3 years but... | no |
| 16 | 71119060 | male | 57 | married | chinese or ... | other | no | 20 years or... | own it c |
| 17 | 71119140 | female | 34 | single | black or bl. | appreciate... | no | 5 years but... | yes |
| 18 | 71119180 | male | 55 | single | white | degree or d. | yes | 10 years b. | buying |
| 19 | 71119220 | female | 27 | single | white | degree or d. | no | 2 years but... | buying |
| 20 | 71119260 | female | 37 | divorced | white | none | no | 10 years b. | buying |

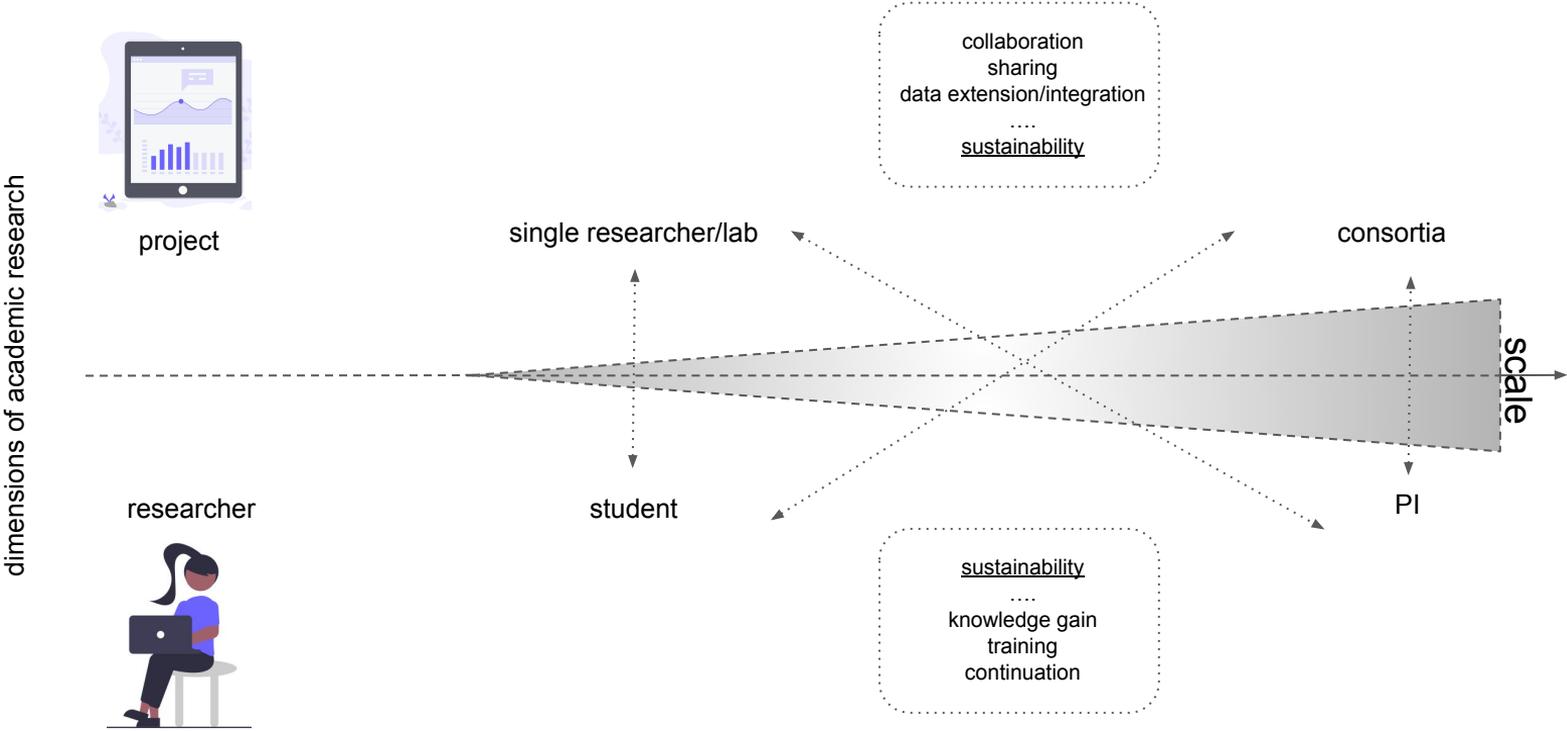
IBM SPSS Statistics Processor is ready

<https://bizlib247.files.wordpress.com/2014/04/data-editor-labels.png>

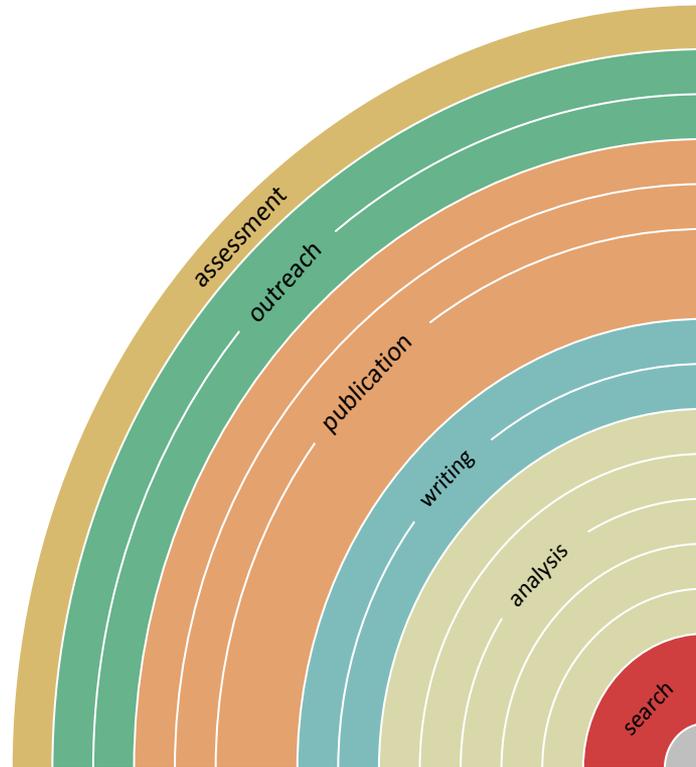
```
sub-control01/
ana/
sub-control01_T1w.nii.gz
sub-control01_T1w.json
sub-control01_T2w.nii.gz
sub-control01_T2w.json
func/
sub-control01_task-nback_bold.nii.gz
sub-control01_task-nback_bold.json
sub-control01_task-nback_events.tsv
sub-control01_task-nback_physio.tsv.gz
sub-control01_task-nback_physio.json
sub-control01_task-nback_sbref.nii.gz
dwi/
sub-control01_dwi.nii.gz
sub-control01_dwi.bval
sub-control01_dwi.bvec
fmap/
sub-control01_phasediff.nii.gz
sub-control01_phasediff.json
sub-control01_magnitudel1.nii.gz
sub-control01_scans.tsv
code/
deface.py
derivatives/
README
participants.tsv
dataset_description.json
CHANGES
```

```
{
  "test": {
    "LongName": "Education level",
    "Description": "Education level, self-rated by participant",
    "Levels": {
      "1": "Finished primary school",
      "2": "Finished secondary school",
      "3": "Student at university",
      "4": "Has degree from university"
    }
  },
  "bmi": {
    "LongName": "Body mass index",
    "Units": "kg/m^2",
    "TermURL": "http://purl.bioontology.org/ontology/SNOMEDCT/60621009"
  }
}
```

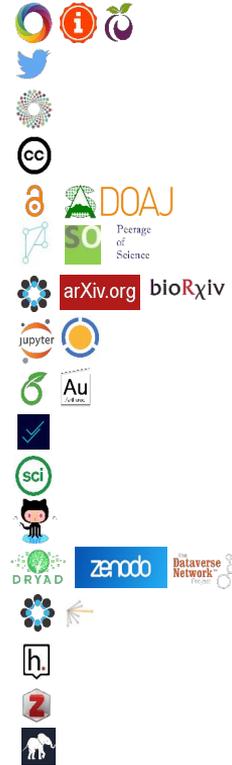
FAIR for all? YES!



FAIR by default, from the beginning to the end and beyond



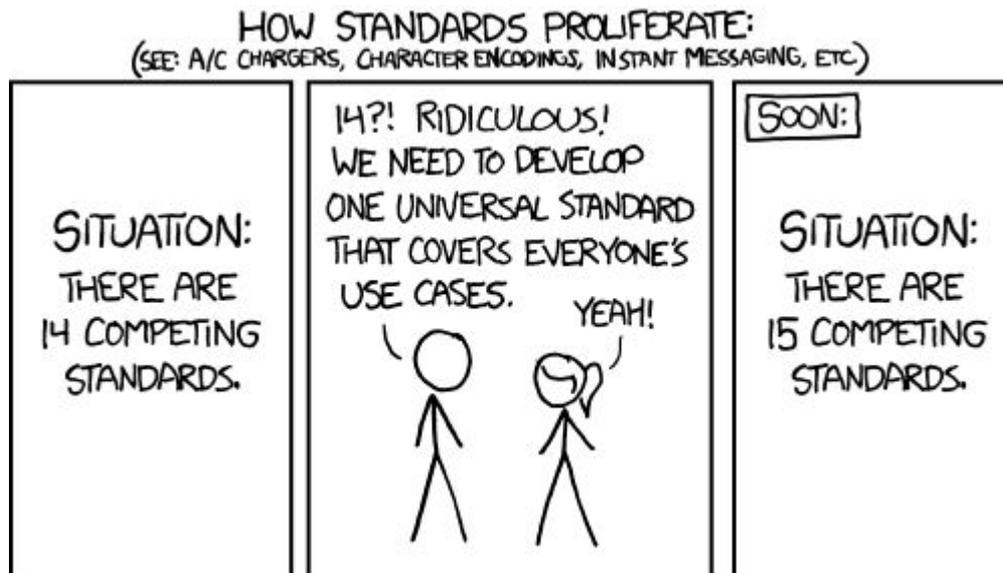
- adding alternative evaluation, e.g. with altmetrics
- communicating through social media, e.g. Twitter
- sharing posters & presentations, e.g. at FigShare
- using open licenses, e.g. CC0 or CC-BY
- publishing open access, 'green' or 'gold'
- using open peer review, e.g. at journals or PubPeer
- sharing preprints, e.g. at OSF, arXiv or bioRxiv
- using actionable formats, e.g. with Jupyter or CoCalc
- open XML-drafting, e.g. at Overleaf or Authorea
- sharing protocols & workfl., e.g. at Protocols.io
- sharing notebooks, e.g. at OpenNotebookScience
- sharing code, e.g. at GitHub with GNU/MIT license
- sharing data, e.g. at Dryad, Zenodo or Dataverse
- pre-registering, e.g. at OSF or AsPredicted
- commenting openly, e.g. with Hypothes.is
- using shared reference libraries, e.g. with Zotero
- sharing (grant) proposals, e.g. at RIO



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

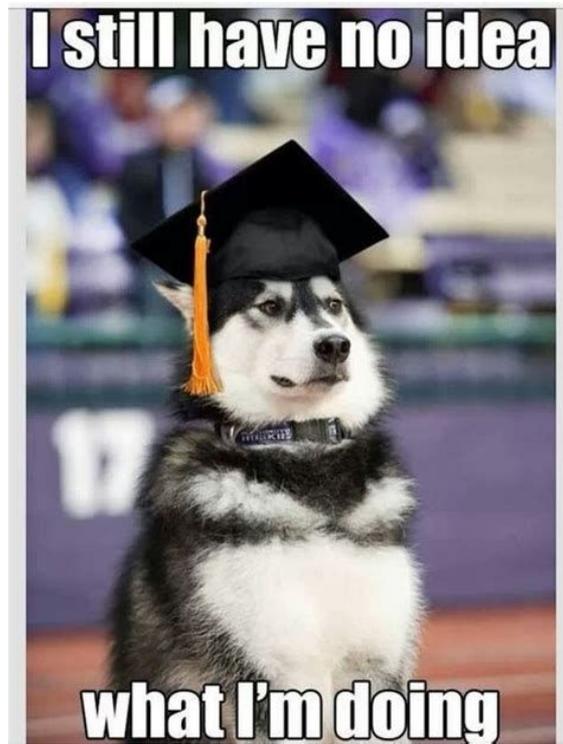
[DOI: 10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR data through standardization - problem I



<https://imgs.xkcd.com/comics/standards.png>

FAIR data through standardization - problem II



“The broad adoption of computers in experimental science has both enabled and obscured research” (Marwick, 2015)

- digital literacy, project/data management, FAIR, open science, etc. not part of most curricula (independent of field of study)
- vast majority across levels not trained in practical skills needed for sufficient and FAIR research

FAIR data through standardization - problem III



* researchers (looking at you students) are frustrated because their programme did not provide them with an realistic point of view regarding research workflows
→ the folks who do the analyses



<http://ggh.is/1H5ABV5>

* the people at the top (looking at you PIs) often expect comprehensive and exhaustive skills without providing the means and opportunities to learn them
→ the folks who don't do the analyses

FAIR data through standardization - problem III

the year 2020...



* there's a ton of open data out there with more participants than folks could have acquired themselves, there's open source software for everything, there are open and free cloud/remote computing possibilities, there are open and free lecture/tutorials/workshops/etc.
→ no one trained to use these opportunities

* PIs/institutes/departments "force" researchers to go the office, acquire data, etc.

* researchers (looking at you students) are frustrated because they can't acquire data, can't go to the lab to run analyses, can't discuss work with peers
→ a lot of projects/theses are effectively on hold since March

* PIs/institutes/departments not prepared for any of this
→ management problems get worse, major problems re grant applications, etc.

FAIR data through standardization - problem III (spoilers)

Offene Wissenschaft in der Zeit von Covid-19 – Ein Blaupause für die psychologische Forschung?

Julia Beitner¹, Garvin Brod^{2,3}, Benjamin Gagl¹, Dominik Kraft⁴, und Martin Schultze⁵

¹Allgemeine Psychologie I, Goethe-Universität Frankfurt am Main. Theodor-W.-Adorno-Platz 6, D-60323 Frankfurt am Main.

²DIPF | Leibniz-Institut für Bildungsforschung und Bildungsinformation, Rostocker Str. 6, D-60323 Frankfurt am Main.

³Pädagogische Psychologie, Goethe-Universität Frankfurt am Main. Theodor-W.-Adorno-Platz 6, D-60323 Frankfurt am Main.

⁴Neurokognitive Psychologie, Goethe-Universität Frankfurt am Main. Theodor-W.-Adorno-Platz 6, D-60323 Frankfurt am Main.

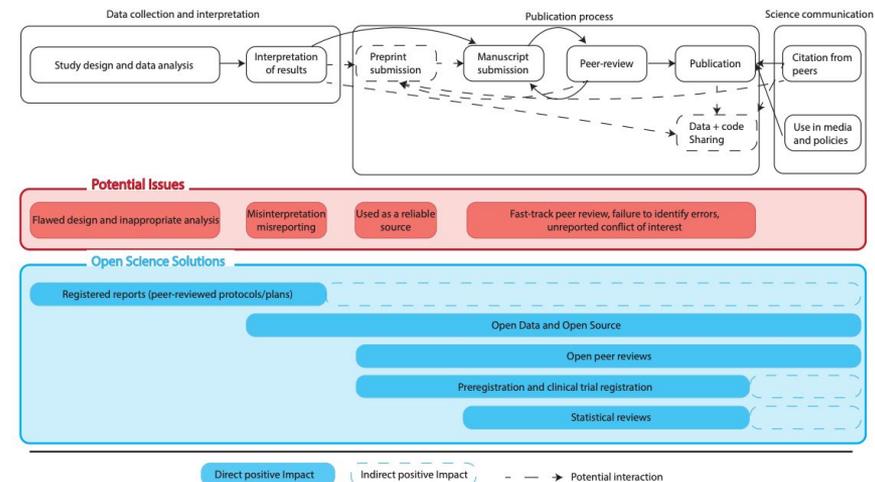
⁵Psychologische Methoden mit interdisziplinärer Ausrichtung, Goethe-Universität Frankfurt am Main. Theodor-W.-Adorno-Platz 6, D-60323 Frankfurt am Main.

Alle Open Science Initiative Frankfurt am Main. Autoreihenfolge alphabetisch.

<https://doi.org/10.31234/osf.io/sh8xg>

Open Science Saves Lives: Lessons from the COVID-19 Pandemic

Lonni Besançon^{1,2*}, Nathan Peiffer-Smadja^{3,4}, Corentin Segalas⁵, Haiting Jiang⁶, Paola Masuzzo⁷, Cooper Smout⁷, Eric Billy⁸, Maxime Deforet⁹⁺ and Clémence Leyrat^{5,10+}



<https://doi.org/10.1101/2020.08.13.249847>

FAIR data through standardization - problem I/II/III - the goal



“Das Internet ist Neuland.”

(“The internet is uncharted territory.”)

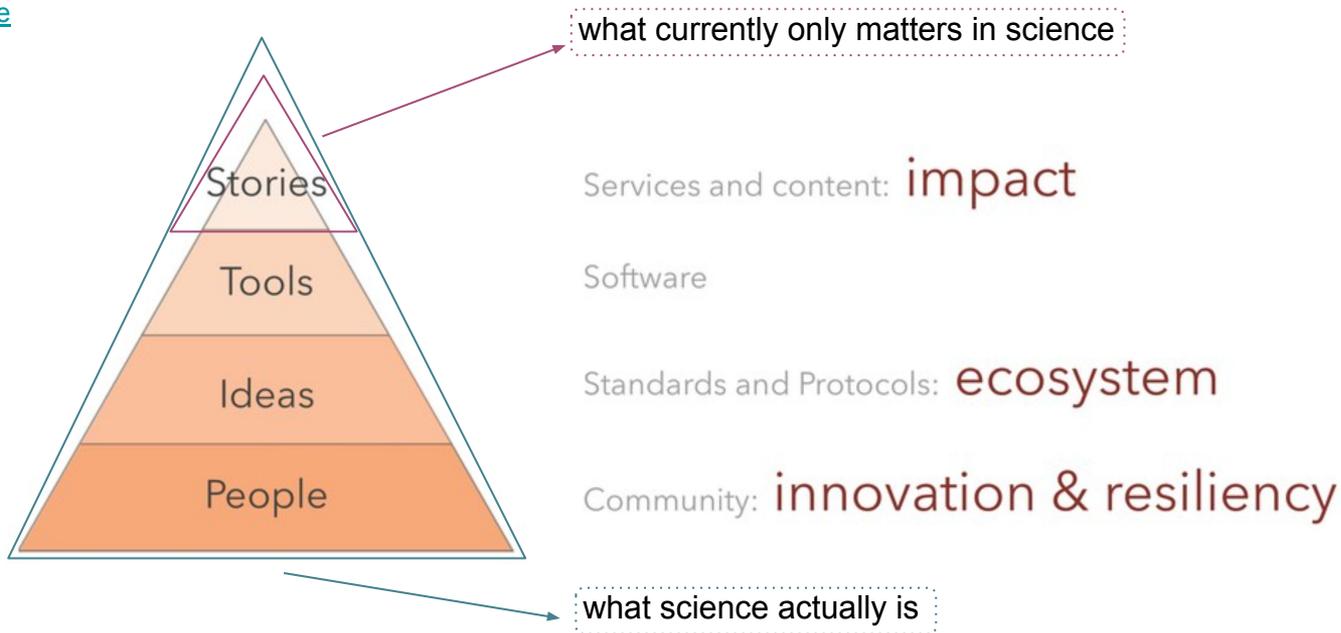
- * How do I manage projects and data?
- * Why do I have to conduct the x-th underpowered study with limited resources?
- * Why can't I access data and run analyses from everywhere?
- * Why do I have to learn everything on my own and am dependent on the good will of my supervisors?

“The future is now.”

- * I have a solid project and data management plan.
- * I can use open data from thousands of participants.
- * No matter where I am, I can work with my data.
- * There is a large international, open and supportive community that provides help no matter what.

FAIR data through standardization - problem IV

*adapted from [Fernando Perez](#)
and [Elizabeth DuPre](#)



*“An article about computational science in a scientific publication is not the scholarship itself, it is **merely advertising of the scholarship**. The **actual scholarship** is the complete software development environment and the complete set of instructions which generated the figures.”*

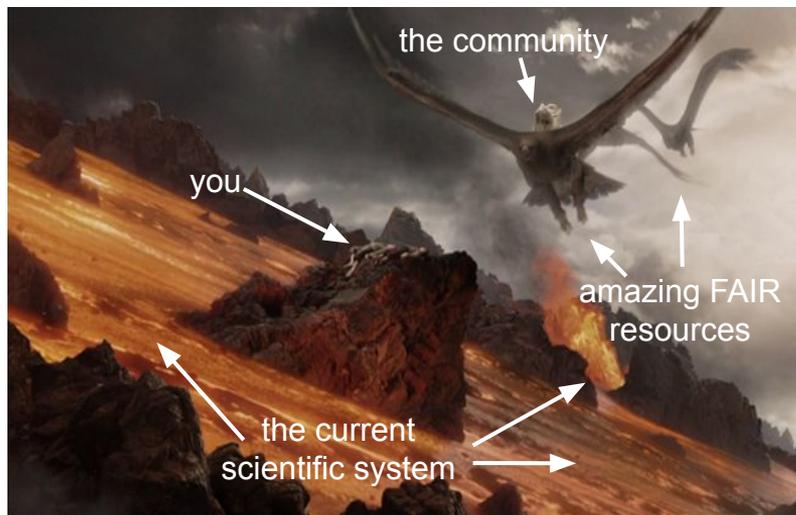
Buckhelt and Donoho
(paraphrasing John Claerbout)
WaveLab and Reproducible Research, 1995

FAIR data through standardization - problem IV

| | open communities | classic science/academia |
|--------------------------------|---|-------------------------------|
| Credit | Distributed | PI & hierarchy |
| Output/artifacts | Continuous & Project-specific | Discrete papers |
| Collaborators | Fluid: professionals, volunteers, ... | Structured, funding-dependent |
| Governance/ decision making | Open, community based | Top-down, PI |
| Authorship | Fluid, roles can evolve, no clear "first/ senior" author | Need to say more? |
| Peer review | Continuous, open, pervasive, friendly | The opposite |
| Value metric | Utility, need, impact | "Novel and transformative" |

*[adapted from Fernando Perez](#)

FAIR data through standardization - problem IV



VS



community based governance

- * The governance and decision making process is clearly structured, open to everyone from the community, actively seeking input from as many folks from as many backgrounds as possible, as well as allows to incorporate changes and updates in a continuous manner.

hierarchical top-down power structure

- * A few powerful chosen ones make decisions that affect a tremendous amount of those “below” them, even though they are not aware of latest developments and haven’t worked with data since ages.

Rhetorical question: did I chose an analogy depicting old white men on purpose?

FAIR data through community-driven development of standards and beyond

*adapted from [Elizabeth DuPre](#)

- **openly** develop standards, materials, resources
- **consensus** in decision-making
- **empower** and **equip** community members
- communities **grow from conscious investment** rather than the availability of resources
- communities and science in general need to be drastically more **open, welcoming** and **supportive** towards everyone, especially underrepresented and marginalized groups
- **every single individual** can and will bring something important to the scientific table, **if they get the chance to**



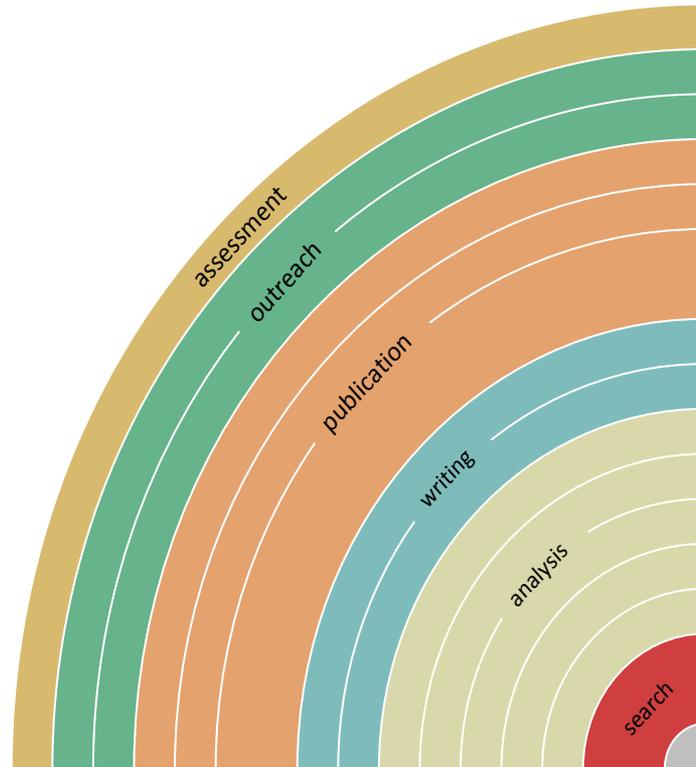
FAIR data through community-driven development of standards and beyond

*adapted from [Elizabeth DuPre](#)

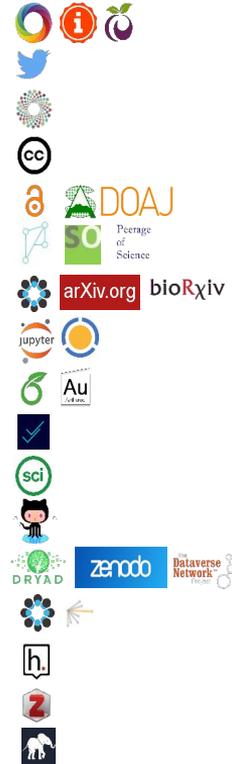
- **combine experience** across all levels (“big picture” folks and those that actually work with data)
- **community is required** as no one has all use cases
- **open discussion and development** necessary for acceptance and adaptation
- **community-driven standards** enable new kinds of science
- **everything needed for FAIR research outcomes exist** but individuals need to be empowered to use them and supported along the way



FAIR data through community-driven development of standards and beyond



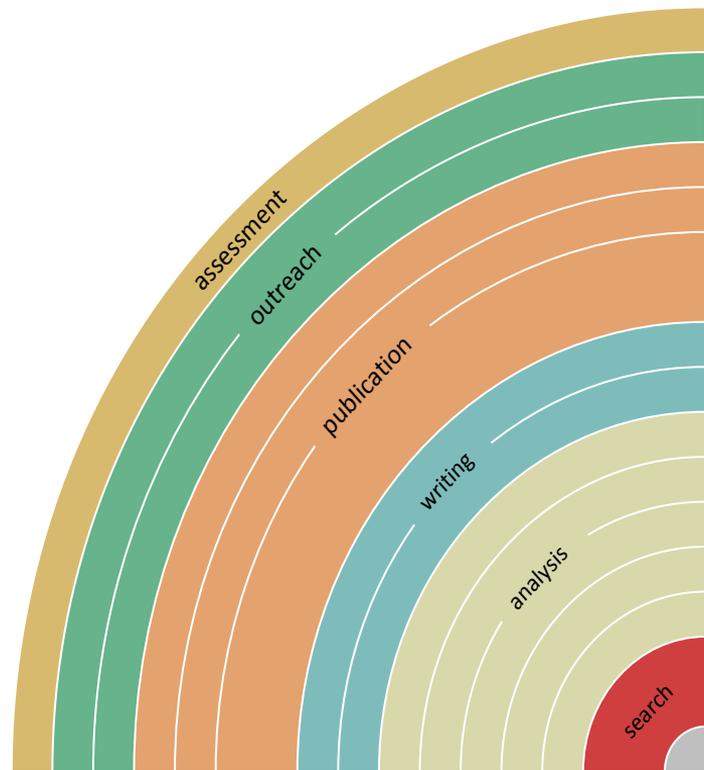
- adding alternative evaluation, e.g. with altmetrics
- communicating through social media, e.g. Twitter
- sharing posters & presentations, e.g. at FigShare
- using open licenses, e.g. CC0 or CC-BY
- publishing open access, 'green' or 'gold'
- using open peer review, e.g. at journals or PubPeer
- sharing preprints, e.g. at OSF, arXiv or bioRxiv
- using actionable formats, e.g. with Jupyter or CoCalc
- open XML-drafting, e.g. at Overleaf or Authorea
- sharing protocols & workfl., e.g. at Protocols.io
- sharing notebooks, e.g. at OpenNotebookScience
- sharing code, e.g. at GitHub with GNU/MIT license
- sharing data, e.g. at Dryad, Zenodo or Dataverse
- pre-registering, e.g. at OSF or AsPredicted
- commenting openly, e.g. with Hypothes.is
- using shared reference libraries, e.g. with Zotero
- sharing (grant) proposals, e.g. at RIO



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

[DOI: 10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR data through community-driven development of standards and beyond



- necessities - support
- prerequisites - initiatives
- necessities - repositories (standardized)
- necessities - more than a PDF
- necessities - reporting (standardized)
- necessities - documentation
- necessities - virtualization (standardized)
- necessities - workflows (standardized)
- necessities - validation (standardized)
- necessities - standardization
- prerequisites - data sharing (standardized)
- prerequisites - preregistration (standardized)



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

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FAIR data through community-driven development of standards and beyond



- necessities - standardization
- prerequisites - data sharing
- **prerequisites - preregistration**



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

[DOI: 10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR prerequisites - preregistration (standardized)

Future-proof your research.
Preregister your next study.

What is Preregistration?

When you preregister your research, you're simply specifying your research plan in advance of your study and submitting it to a registry.

Preregistration separates *hypothesis-generating* (exploratory) from *hypothesis-testing* (confirmatory) research. Both are important. But the same data cannot be used to generate and test a hypothesis, which can happen unintentionally and reduce the credibility of your results. Addressing this problem through planning improves the quality and transparency of your research. This helps you clearly report your study and helps others who may wish to build on it.

For additional insight and context, you can read [The Preregistration Revolution](#), (preprint)



[Get Started Now](#)

Confirmatory Research

- Hypothesis testing
- Results are held to the highest standards
- Data-independent
 - Minimizes false positives
 - P-values retain diagnostic value
- Inferences may be drawn to wider population

Exploratory Research

- Hypothesis generating
- Results deserve to be replicated and confirmed
- Data-dependent
 - Minimizes false negatives in order to find unexpected discoveries
 - P-values lose diagnostic value
- Not useful for making inferences to any wider population



Create a new pre-registration

[CREATE](#)

[Just trying it out; make this pre-registration self-destruct in 24 hours.](#)

See your pre-registrations

(e.g., to share with reviewers or make public)

[Sign in](#)

WHAT IS ASPREDICTED?

AsPredicted is a platform that makes it easy for researchers to pre-register their studies, and easy for others to read and evaluate those pre-registrations. To pre-register a study on AsPredicted, a researcher answers nine simple questions about their research design and analyses. The platform then generates a time-stamped, single page .pdf document that includes a unique URL for verification.

HOW DOES IT WORK?

- One author creates the pre-registration.
- Participating authors are emailed, requesting approval.
- If all approve, it is saved but remains private until an author makes it public; or remains private forever. ([Why?](#))
- Authors may share an anonymous version of the pre-registration with reviewers.
- If made public, the final .pdf (sample) is automatically stored in the [web-archive](#).

WHAT IF THINGS DON'T GO 'AS PREDICTED'?

You can just say so in the paper:

- 'Contrary to expectations, we found that...'
- 'Unexpectedly, we also found that...'
- 'In addition to the analyses we pre-registered we also ran...'
- 'We encountered an unexpected situation, and followed our Standard Operating Procedure ([pdf](#))'

FAIR prerequisites - preregistration (standardized)



MAX PLANCK INSTITUTE
FOR HUMAN COGNITIVE AND BRAIN SCIENCES

INSTITUTE | NEWS | RESEARCH | LIBRARY | CAREER | SCHOOLS

Home > Institute-wide cooperations > CBS Open Science > Open Science Events > Preregistration-Hackathon

Preregistration-Hackathon (June 18, 2020)

At this hackathon, we will be working on preregistration templates for (f)MRI and M/EEG.

JUNE 18, 2020

With these templates, we aim to promote and facilitate preregistrations for neuropsychological research in which a myriad of acquisition & analysis decisions have to be pre-specified. This work was started at the SIPS 2019 meeting and continued during one of our own events last fall. We are eager to move the templates forward with your help. If you have experience with (f)MRI or M/EEG and are interested in preregistrations, please join us! The hackathon will take place online via zoom (please contact Mariella Paul to receive the link (paulm@cbs.mpg.de)).

You can find the current versions of the templates here: [🔗 EEG](#), [🔗 fMRI](#)

<https://www.cbs.mpg.de/1543216/preregistration-hackathon>

fMRI preregistration template | Jessica Flannery, adapted by OSI MPI-CBS

The (f)MRI preregistration template

Usage Notes to the template

The goal of this template is to provide sufficient information in preregistration for (f)MRI data design to increase reproducible reporting practices. All prompts and tables can be used to fill in as it, or as a checklist of information you include for text for that section or checklist for attached files (e.g., json file of scan parameters). Tip: Using it as a checklist will allow you to write these sections as they will appear in your future paper. If a certain table/section does not apply, can just state "N/A."

Information on the evolution of the template:

The original document was written by Jessica Flannery. She adopted the text from the OSF preregistration challenge template to include the details important for fMRI psychology design. The OSF template was retained in some areas and edited in other areas to incorporate both prior published templates and guidelines regarding fMRI (Nichols et al., 2016; Poldrack et al., 2008; van't Veer & Giner-Sorolla, 2016).

The Open Science Initiative at the MPI-CBS worked on this template during two preregistration hackathons.

During the first hackathon in November 2019, we discussed the structure and rearranged the sections of the preregistration so that they matched the [OSF preregistration template](#) (Study Information, Design Plan, Sampling Plan, Variables, Analysis Plan, Other). We also agreed it would be best to first focus on task/resting state fMRI studies with this template (e.g. exclude sMRI and DWI for the moment).

During a second hackathon on June 18 2020, we worked on the details of the experimental design (regarding neuropsychological testing and design/implementation of the behavioral task). We also swapped the order of Design and Sampling Plan from the original OSF template. We restructured the Acquisition and Preprocessing section to improve the usability regarding the details of MRI sequences and preprocessing settings.

Remi Gau
16:52 23 Aug

We might want to cross check the MRI and the EEG templates for overlaps (especially in the "general" parts like sampling plan, study design). If some things can be fused between the 2 it would be better to have only one version.

Beyer Frauke
12:12 26 Aug

Add: "randomized controlled trial" = the study involves a manipulation of exposures on the subject-level (e.."



FAIR data through community-driven development of standards and beyond



- necessities - standardization
- **prerequisites - data sharing**
- prerequisites - preregistration



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

[DOI: 10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR prerequisites - data sharing (standardized)

<https://open-brain-consent.readthedocs.io/>



Docs • Make open data sharing a no-brainer for ethics committees. [Edit on GitHub](#)

Make open data sharing a no-brainer for ethics committees.

DOI: [10.5281/zenodo.3332807](https://doi.org/10.5281/zenodo.3332807)

Single access type version (all data shared publicly; recommended)

Version: OBC-ULT 1.0.0

The data and samples from this study might be used for other, future research projects in addition to the study you are currently participating in. Those future projects can focus on any topic that might be unrelated to the goals of this study. We will give access to the data we are collecting, including the imaging data, to the general public via the internet and a fully open database.

The data we share with the general public will not have your name on it, only a code number, so people will not know your name or which data are yours. In addition, we will not share any other information that we think might help people who know you guess which data are yours.

If you change your mind and withdraw your consent to participate in this study (you can call <PI name> at <phone number> to do this), we will not collect any additional data about you. We will delete your data if you withdraw before it was deposited in the database. However, any data and research results already shared with other investigators or the general public cannot be destroyed, withdrawn or recalled.

By agreeing to participate, you will be making a free and generous gift for research that might help others. It is possible that some of the research conducted using your information eventually could lead to the development of new methods for studying brain, new diagnostic tests, new drugs or other commercial products. Should this occur, there is no plan to provide you with any part of the profits generated from such products and you will not have any ownership rights in the products.

To the best of our knowledge, the data we release to the general public will not contain information that can directly identify you. The data will not have your name on it, only a code number, so people will not know your name or which data are yours. In addition, the data will not include data that we think might help people who know you guess which data are yours, such as your facial features or the date that you participated. If we write a report or article about this study or share the study data set with others, we will do so in such a way that you cannot be directly identified. However, by using additional data linked to your name (for example brain scans obtained from your medical records) one could potentially associate your imaging or other information in our database back to you. In addition a security breach (break in or cyber attack) might lead to someone being able to link you to your data. This risk is very low because your data are stored in a secure database, and the information about your identity is stored separately from the data themselves, linked only through a code.

We will keep the private portion (name, contact information etc.) of your data in a secure location for at least <x> years. This way if one of the researchers that obtained the data from us will find something in your brain scans that would have a diagnostic value we will be able to contact you. After this period of time we will destroy this information to protect your privacy.

Letting us use and share your data is voluntary. However, you must be willing to share your data in this way in order to participate in this study. If you are not willing, you cannot participate in this study.

By signing below, you agree to provide your data for future research. You agree that these may be shared with other investigators at other institutions from around the world. The details, results, and implications of these studies are unknown.

- data privacy & rights > FAIR/open science
- data privacy & rights need to be protected
- identification attacks & data breaches
- complicated legal issues and varying interpretations of laws
- researchers not trained



CONSIDER
NECESSARY
RESTRICTIONS
AROUND
SHARING
DIFFERENT
TYPES OF DATA



Sonbeia

<http://doi.org/10.5281/zenodo.3332807>



FAIR data through community-driven development of standards and beyond



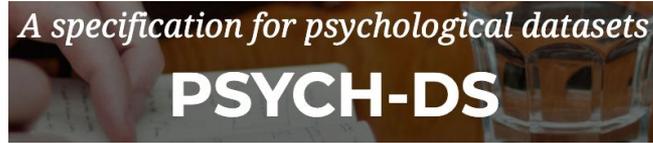
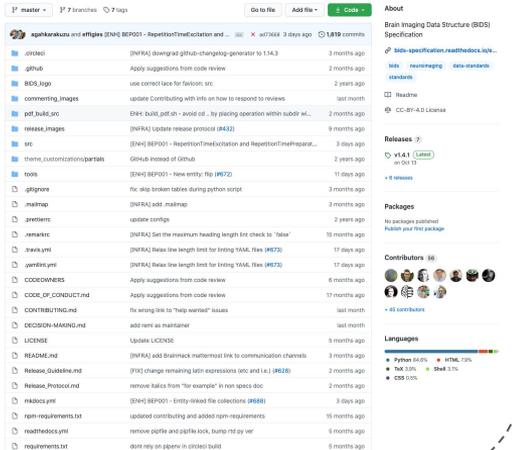
- **necessities - standardization**
- prerequisites - data sharing
- prerequisites - preregistration



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

[DOI: 10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR necessities - standardization - fantastic standards in the wild



Psych-DS

A technical specification for psychological datasets

Version 0.3.0 ("POST SIPS2020" DRAFT) - IN PROGRESS

Available under the CC-BY 4.0 International license
Based on the [BIDS specification](#) for fMRI data, with inspiration from the [BIDS Eye-tracking extension draft](#).

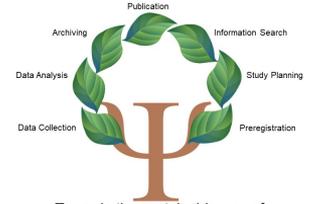
Welcome! If this is your first time checking out this project (and especially if this is your first time working with a technical specification document), please also check out [the project landing page](#) for an explanation of the goals of this project and notes on how you can contribute.

If you are working on creating a Psych-DS compliant dataset, please check out the [How-To page!](#)

The technical specification document should focus closely on the definition of Psych-DS. Background information on how or why to use Psych-DS (or comments for tool implementers) should be factored either into the [How-To page](#) or into the [position paper draft](#).

This document contains a draft of a data structure standard for psychological datasets. It is a community effort to define standards in data / metadata. This is a working document in draft stage and **comments are welcome**.

This specification is an extension of [BIDS](#), and general principles are shared. The Psych-DS specification aims to work for many different settings and facilitate integration with other types of data.



Towards the sustainable use of psychological research data



The Project PsyCurDat - Development of user-oriented curation criteria for psychological research data

Project description

In the context of the omniresent Open Science Movement, all stakeholders in psychological research (i.e. researchers, journals, and funders) increasingly strive for more transparency in order to foster scientific exchange and the quality of scientific work. However, to reach this overarching goal of the Open Science Movement, there is a need for discipline-specific documentation standards not only for research articles, but also for the data representing the basis of those articles. Against this background, the project [PsyCurDat](#): Development of user-oriented curation criteria for psychological research data, which is funded by the Federal Ministry of Education and Research (BMBWF), pursues two strategic objectives:

1. To meet researchers' needs in their role as data users, the project aims at developing a documentation standard for psychological research data considering the discipline specific methods relevant for reusing these data.
2. To meet researchers' needs in their role as contributors of research data, the project aims at developing a documentation standard considering determinants defined by the research process as well as by funders and journals.

Working towards these two strategic goals, [PsyCurDat](#) generally aims at promoting research economy and research integrity in Psychology by enabling a more effective and efficient documentation and reuse of psychological research data.

FAIR necessities - standardization - the idea of BIDS

How do you manage your data?

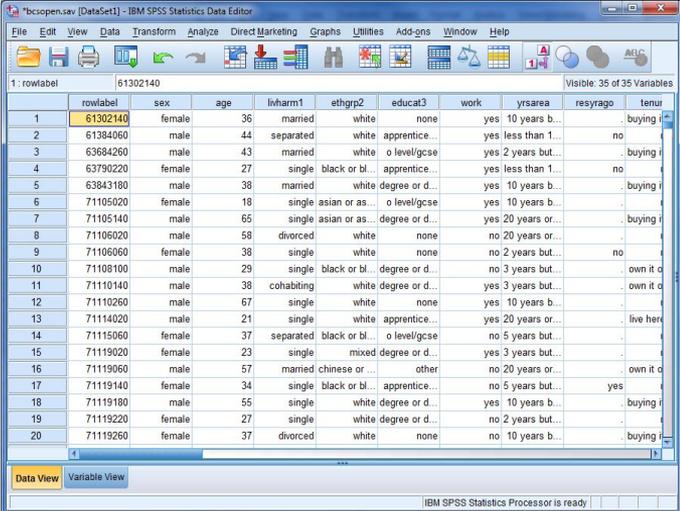
- storage, structure, metadata, version control?

How do you share your data?

- colleagues, students, other researchers?

The Problem with heterogeneity in data management is it is hard for others (and you) to understand your data and keep track of changes

- unnecessary metadata input
- codes / scripts have to be adapted
- huge effort to automate workflows and no way to automatically validate data sets
- sharing data becomes a hassle

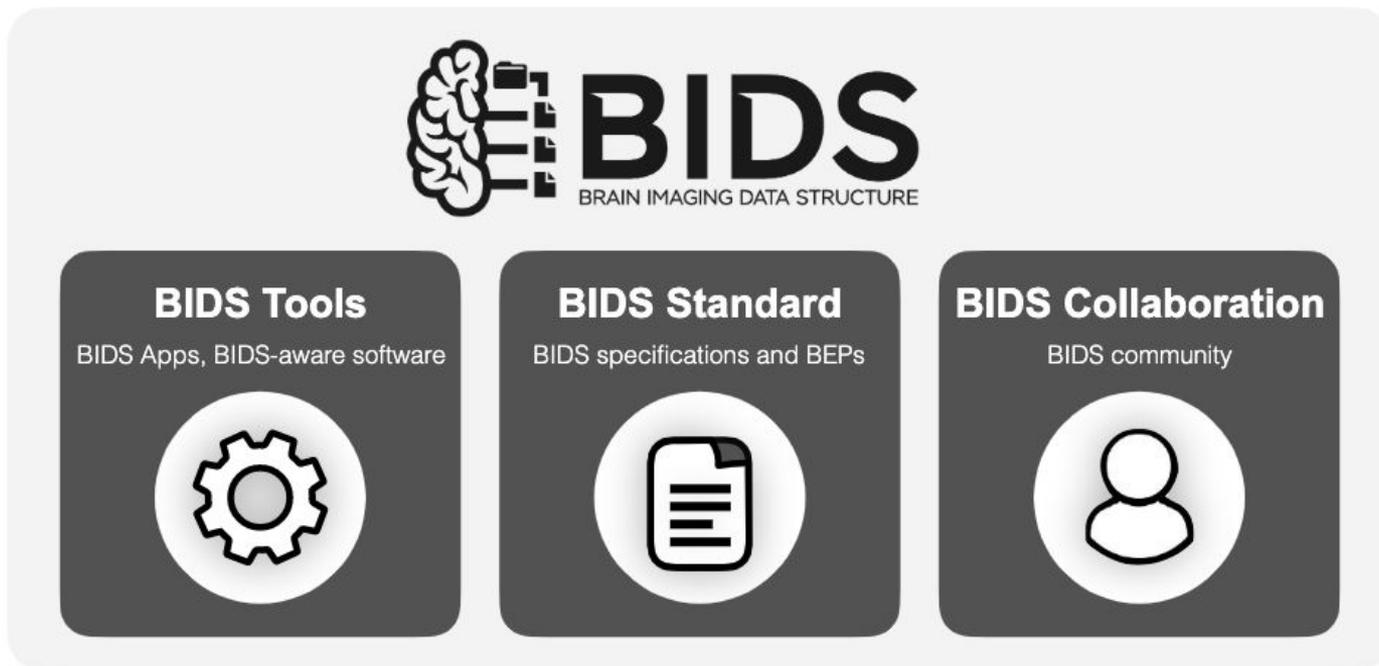


| rowlabel | sex | age | lvharm1 | ethgrp2 | educat3 | work | yrsarea | resyrago | tenu |
|----------|----------|--------|---------|------------|----------------|----------------|--------------------|----------|----------|
| 1 | 61302140 | female | 36 | married | white | none | yes 10 years b... | | buying r |
| 2 | 61384060 | male | 44 | separated | white | apprentice... | yes less than 1... | no | |
| 3 | 63684260 | male | 43 | married | white | o level/gcse | yes 2 years but... | | buying r |
| 4 | 63790220 | female | 27 | single | black or bl... | apprentice... | yes less than 1... | no | |
| 5 | 63843180 | male | 38 | married | white | degree or d... | yes 10 years b... | | buying r |
| 6 | 71105020 | female | 18 | single | asian or as... | o level/gcse | yes 10 years b... | | |
| 7 | 71105140 | male | 65 | single | asian or as... | degree or d... | yes 20 years or... | | buying r |
| 8 | 71106020 | male | 58 | divorced | white | none | no 20 years or... | | |
| 9 | 71106060 | female | 38 | single | white | none | no 2 years but... | no | |
| 10 | 71108100 | male | 29 | single | black or bl... | degree or d... | no 3 years but... | | own it d |
| 11 | 71110140 | male | 38 | cohabiting | white | degree or d... | yes 3 years but... | | own it d |
| 12 | 71110260 | male | 67 | single | white | none | yes 10 years b... | | |
| 13 | 71114020 | male | 21 | single | white | apprentice... | yes 20 years or... | | live her |
| 14 | 71115060 | female | 37 | separated | black or bl... | o level/gcse | no 5 years but... | | |
| 15 | 71119020 | female | 23 | single | mixed | degree or d... | yes 3 years but... | | |
| 16 | 71119060 | male | 57 | married | chinese or ... | other | no 20 years or... | | own it d |
| 17 | 71119140 | female | 34 | single | black or bl... | apprentice... | no 5 years but... | yes | |
| 18 | 71119180 | male | 55 | single | white | degree or d... | yes 10 years b... | | buying r |
| 19 | 71119220 | female | 27 | single | white | degree or d... | no 2 years but... | | |
| 20 | 71119260 | female | 37 | divorced | white | none | no 10 years b... | | buying r |

<https://bzlib247.files.wordpress.com/2014/04/data-editor-labels.png>

FAIR necessities - standardization - the idea of BIDS

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)



FAIR necessities - standardization - the idea of BIDS ... *with Marie Kondo*



```
dicomdir/  
  1208200617178_22/  
    1208200617178_22_8973.dcm  
    1208200617178_22_8943.dcm  
    1208200617178_22_2973.dcm  
    1208200617178_22_8923.dcm  
    1208200617178_22_4473.dcm  
    1208200617178_22_8783.dcm  
    1208200617178_22_7328.dcm  
    1208200617178_22_9264.dcm  
    1208200617178_22_9967.dcm  
    1208200617178_22_3894.dcm  
    1208200617178_22_3899.dcm  
  1208200617178_23/  
  1208200617178_24/  
  1208200617178_25/
```



```
my_dataset/  
  participants.tsv  
  sub-01/  
    anat/  
      sub-01_T1w.nii.gz  
    func/  
      sub-01_task-rest_bold.nii.gz  
      sub-01_task-rest_bold.json  
    dwi/  
      sub-01_dwi.nii.gz  
      sub-01_dwi.json  
      sub-01_dwi.bval  
      sub-01_dwi.bvec  
  sub-02/  
  sub-03/  
  sub-04/
```



FAIR necessities - standardization - the idea of BIDS

* content liberally borrowed from
Chris Markiewicz's [BIDS ecosystem presentation](https://bids-specification.readthedocs.io/en/stable/)

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

- **BIDS is a directory structure and naming convention**, based on common practices
- builds on **existing open standards** (NIfTI, JSON, TSV)
- intended for **human and machine legibility**
- **metadata** for all files and across different levels
- **maintained and extended** by a clearly structured and open community



```
sub-control01/  
  anat/  
    sub-control01_T1w.nii.gz  
    sub-control01_T1w.json  
    sub-control01_T2w.nii.gz  
    sub-control01_T2w.json  
  func/  
    sub-control01_task-nback_bold.nii.gz  
    sub-control01_task-nback_bold.json  
    sub-control01_task-nback_events.tsv  
    sub-control01_task-nback_physio.tsv.gz  
    sub-control01_task-nback_physio.json  
    sub-control01_task-nback_sbref.nii.gz  
  dwi/  
    sub-control01_dwi.nii.gz  
    sub-control01_dwi.bval  
    sub-control01_dwi.bvec  
  fmap/  
    sub-control01_phasediff.nii.gz  
    sub-control01_phasediff.json  
    sub-control01_magnitude1.nii.gz  
    sub-control01_scans.tsv  
  code/  
    deface.py  
  derivatives/  
  README  
  participants.tsv  
  dataset_description.json  
  CHANGES
```

```
{  
  "test": {  
    "LongName": "Education level",  
    "Description": "Education level, self-rated by participant",  
    "Levels": {  
      "1": "Finished primary school",  
      "2": "Finished secondary school",  
      "3": "Student at university",  
      "4": "Has degree from university"  
    }  
  },  
  "bmi": {  
    "LongName": "Body mass index",  
    "Units": "kg/m^2",  
    "TermURL": "http://purl.bioontology.org/ontology/SNOMEDCT/68621009"  
  }  
}
```

FAIR necessities - standardization - the idea of BIDS

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)

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

dataset_description.json

```
{  
  "Name": "The mother of all experiments",  
  "BIDSVersion": "1.4.0",  
  "DatasetType": "raw",  
  "License": "CC0",  
  "Authors": [ "Paul Broca", "Carl Wernicke" ],  
  "Acknowledgements": "Special thanks to KB for help in formatting  
this dataset in BIDS",  
  "HowToAcknowledge": "Please cite this paper: https://  
www.ncbi.nlm.nih.gov/pubmed/0928",  
  "Funding": [ "National Institute of Neuroscience Grant F37823MFH1" ],  
  "EthicsApprovals": [ "Human Research Protections Office (Protocol  
AR0928" ],  
  "ReferencesAndLinks": [ "https://www.ncbi.nlm.nih.gov/pubmed/  
0928", "Alzheimer et al (2015). Nature, 21. doi.org/0928" ],  
  "DatasetDOI": "10.0.2.3/dfjj.10"  
}
```



example1/

- |— CHANGES
- |— README.tsv
- |— **dataset_description.json**
- |— participants.json
- |— participants.tsv
- |— **sub-001**
 - |— ses-01
 - |— ses-02
 - |— sub-001_scans.tsv
- |— **sub-002**
- |— **sub-003**

FAIR necessities - standardization - the idea of BIDS

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)

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

participants.tsv

| participant_id | age | sex |
|----------------|-----|-----|
| sub-001 | 34 | M |
| sub-002 | 12 | F |
| sub-003 | 33 | F |

participants.json

```
"age": {  
  "Description": "age of the participant",  
  "Units": "years" },  
"sex": {  
  "Description": "sex as reported by the participant",  
  "Levels": {  
    "M": "male",  
    "F": "female" }
```



example1/

- |— CHANGES
- |— README.tsv
- |— dataset_description.json
- |— **participants.json**
- |— **participants.tsv**
- |— **sub-001**
 - |— ses-01
 - |— ses-02
 - |— sub-001_scans.tsv
- |— **sub-002**
- |— **sub-003**

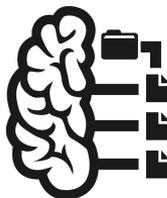
FAIR necessities - standardization - the idea of BIDS

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)

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

sub-001_ses-02_task-rest_eeg.json

```
{  
  "InstitutionName": "Montreal Neurological Institute",  
  "InstitutionAddress": "3801 University, Montreal QC",  
  "Manufacturer": "Brain Products",  
  "ManufacturersModelName": "BrainAmp DC",  
  "DeviceSerialNumber": "11035",  
  "PowerLineFrequency": 60,  
  "SamplingFrequency": 2400,  
  "EEGPlacementScheme": "10 percent system",  
  "EEGReference": "left mastoid",  
  "Softwarefilters": {"Anti-aliasing filter": {"half-amplitude cutoff  
(Hz)": 500, "Roll-off": "6dB/Octave"}},  
  "CapManufacturer": "EasyCap",  
  "CapManufacturersModelName": "M1-ext",  
  "EOGChannelCount": 2,  
  "ECGChannelCount": 1,  
  "EMGChannelCount": 0,  
  "RecordingDuration": 600, "RecordingType": "continuous",  
  "EpochLength": 0,  
  ...  
}
```



BIDS
BRAIN IMAGING DATA STRUCTURE

sub-001

```
├─ ses-02  
  │ └─ meg  
  │   │ └─ sub-001_ses-02_coordsystem.json  
  │   │ └─ sub-001_ses-02_headshape.pos  
  │   │ └─ sub-001_ses-02_task-rest_run-01_channels.tsv  
  │   │ └─ sub-001_ses-02_task-rest_run-01_meg.ds  
  │   └─ sub-001_ses-02_task-rest_run-01_meg.json  
├─ ses-03  
  │ └─ eeg  
  │   │ └─ sub-001_ses-03_task-rest_run-01_eeg.edf  
  │   └─ sub-001_ses-03_task-rest_run-01_eeg.json  
├─ beh  
  │ └─ sub-001_ses-03_task-oddball_run-01_events.tsv  
└─ sub-001_scans.tsv
```



FAIR necessities - standardization - the idea of BIDS

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)

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

- **adoption is crucial**

- minimize complexity and facilitate adoption
- don't reinvent the wheel
- reuse existing methods and technologies

- **80/20 rule**

- tackle 80% of most commonly used cases

- **engagement of a global community**

- everyone can provide input and work on maintenance and extension



```
sub-control01/  
  anat/  
    sub-control01_T1w.nii.gz  
    sub-control01_T1w.json  
    sub-control01_T2w.nii.gz  
    sub-control01_T2w.json  
  func/  
    sub-control01_task-nback_bold.nii.gz  
    sub-control01_task-nback_bold.json  
    sub-control01_task-nback_events.tsv  
    sub-control01_task-nback_physio.tsv.gz  
    sub-control01_task-nback_physio.json  
    sub-control01_task-nback_sbref.nii.gz  
  dwi/  
    sub-control01_dwi.nii.gz  
    sub-control01_dwi.bval  
    sub-control01_dwi.bvec  
  fmap/  
    sub-control01_phasediff.nii.gz  
    sub-control01_phasediff.json  
    sub-control01_magnitude1.nii.gz  
    sub-control01_scans.tsv  
code/  
  deface.py  
derivatives/  
README  
participants.tsv  
dataset_description.json  
CHANGES
```

```
{  
  "test": {  
    "LongName": "Education level",  
    "Description": "Education level, self-rated by participant",  
    "Levels": {  
      "1": "Finished primary school",  
      "2": "Finished secondary school",  
      "3": "Student at university",  
      "4": "Has degree from university"  
    }  
  },  
  "bmi": {  
    "LongName": "Body mass index",  
    "Units": "kg/m^2",  
    "TermURL": "http://purl.bioontology.org/ontology/SNOMEDCT/68621009"  
  }  
}
```



FAIR necessities - standardization - the idea of BIDS <https://bids-specification.readthedocs.io/en/stable/>

<https://bids.neuroimaging.io/>

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)

- BIDS is constantly expanded & updated through community driven development
 - Positron Emission Tomography (PET)
 - Common Derivatives
 - Models Specification
 - Magnetoencephalography (MEG)
 - Electroencephalography (EEG)
 - intracranial Electroencephalography (iEEG)
 - Eye Tracking including Gaze Position and Pupil Size
 - Susceptibility Weighted Imaging (SWI)
 - Genetic information
 - Microelectrode Recordings (MER)
 -

| BEP | Title | Moderators/Leads | BEP | Title | Moderators/Leads |
|--------|---|--------------------------------------|--------|--|--|
| BEP001 | Structural acquisitions with multiple contrasts (multi echo, flip angle, inversion time) seqs | G. de Hollander, K. Whitaker | BEP020 | Eye Tracking including Gaze Position and Pupil Size (ET) | B. Gagli, D. Draschlow |
| BEP002 | BIDS Models | T. Yarkoni | BEP021 | Common Electrophysiological Derivatives | M.Jas, Appelhoff, Pernet, G.Niso, R.Oostenveld |
| BEP004 | Susceptibility Weighted Imaging (SWI) | Available | BEP022 | Magnetic Resonance Spectroscopy (MRS) | M. Mikkelsen, W. Clarke |
| BEP005 | Arterial Spin Labeling (ASL) | Mutsaerts, Clement, Petr, Castellano | BEP023 | PET Preprocessing derivatives | M.Noergaard, Searle, Ganz |
| BEP009 | Positron Emission Tomography (PET) | M. Ganz | BEP024 | Computed Tomography scan (CT) | H. Boniface |
| BEP011 | Structural preprocessing derivatives | V. Siless | BEP025 | Medical Imaging Data structure (MIDS) | J.M.Saborit, M.de la Iglesia |
| BEP012 | Functional preprocessing derivatives | C.Maumet, Markiewicz | BEP026 | Microelectrode Recordings (MER) | G. Gilmore |
| BEP013 | Resting state fMRI derivatives | S. Giavasis | BEP027 | BIDS Execution | C. Markiewicz, G. Klar |
| BEP014 | Affine transformations and nonlinear field warps | O. Esteban | BEP028 | Provenance | S. Ghosh, C. Maumet |
| BEP016 | Diffusion weighted imaging derivatives | F. Pestilli, O. Esteban | BEP029 | Virtual and physical motion data | S. Jeung, J. Welzel |
| BEP017 | Generic BIDS connectivity data schema | E. Duff, P. McCarthy | BEP030 | Near Infrared Spectroscopy (NIRS) | R. Luke, L. Poltonini |
| BEP019 | DICOM Metadata | S. Ghosh | BEP031 | Microscopy | M.Bourget, J.Cohen-Adad |

| BEP | Title | Moderators/Leads |
|--------|--|--|
| BEP003 | Common Derivatives | C. Markiewicz |
| BEP006 | Electroencephalography (EEG) | C. Pernet, S. Appelhoff, R. Oostenveld |
| BEP007 | Hierarchical Event Descriptor (HED) Tags | C. Gorgolewski |
| BEP008 | Magnetoencephalography (MEG) | G. Niso |
| BEP010 | intracranial Electroencephalography (iEEG) | C. Holdgraf, D. Hermes |
| BEP018 | Genetic information | C. Pernet, C. Moreau, T. Nichols |



Brain Imaging Data Structure v1.4.1

The BIDS Specification >

The BIDS Starter Kit >

Filters - [x] Issue is open [x] Labels (2) [x] Milestones (2) [x] New Issue

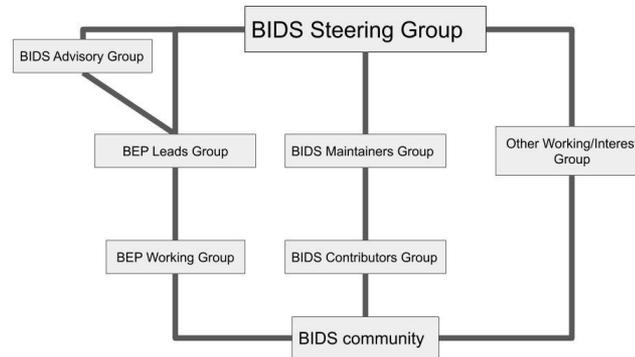
| Issue ID | Status | Author | Label | Projects | Milestones | Assignee | Sort |
|----------|--|------------------|---------------|----------|----------------|----------|------|
| 130 | Open | ✓ 186 | Closed | | | | |
| #588 | Fix example for IEEGElectrodeGroups field from object to string type | EEG | | | | | 5 |
| #585 | Migrate "Travis CI" service to "GitHub Actions" CI service | good first issue | documentation | | | | |
| #580 | Extend BIDS format to magnetic resonance spectroscopy | | | | | | 3 |
| #570 | Change requirement level "optional" -> "recommended" for three fields in dataset_description | options wanted | | | | | 2 |
| #567 | EEG - confusion between channels.tsv and electrodes.tsv | | | | | | 2 |
| #565 | [ENH] add realignment parameters in JSON for "MoCo" sequences | | | | | | |
| #564 | Add OPTIONAL SkullStripped metadata to any MR images | options wanted | | | | | 2 |
| #563 | Add consistent Units information in the json field tables | | | | | | 3 |
| #562 | Bare BIDS - can requirements for README be enhanced? | | | | | | 13 |
| #561 | "T1w" as keyword for <datatype>"CoordinateSystem" | MEG | MEG | new | consistency | EEG | 6 |
| #559 | Alternating row fills in tables within the PDF build | formatting | | | | | |
| #558 | Define supported coordinate systems in the BIDS specification instead of FieldTrip wiki | MEG | MEG | EEG | options wanted | | 8 |
| #554 | Add support for multi-echo T1w scans | | | | | | 11 |
| #547 | EEG channel type inconsistency | consistency | | | | | 2 |
| #544 | Harmonize .tsv examples (and their rendering) | enhancement | | | | | |
| #537 | unspecified status of extra content in the JSON files | | | | | | 3 |
| #527 | RFC: Maintainer/Steering credit on BEP papers | community | | | | | 6 |
| #525 | Is <source_entities> sufficient or should be extended to include "_mod_" when necessary? | | | | | | 2 |

The Brain Imaging Data Structure

This resource defines the Brain Imaging Data Structure (BIDS) specification, including the core specification as well as many modality-specific extensions.

To get started, [check out the introduction](#). If you'd like more information on how to adapt your own datasets to match the BIDS specification, we recommend exploring the [bids-specification starter kit](#).

For an overview of the BIDS ecosystem, visit the [BIDS homepage](#). The entire specification can also be [downloaded as PDF](#).



FAIR necessities - standardization - the idea of BIDS <https://bids-specification.readthedocs.io/en/stable/>

<https://bids.neuroimaging.io/>

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)

BIDS Steering Group 2019-2020



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The Netherlands
M/EEG



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Kirstie Whitaker
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BIDS Maintainers



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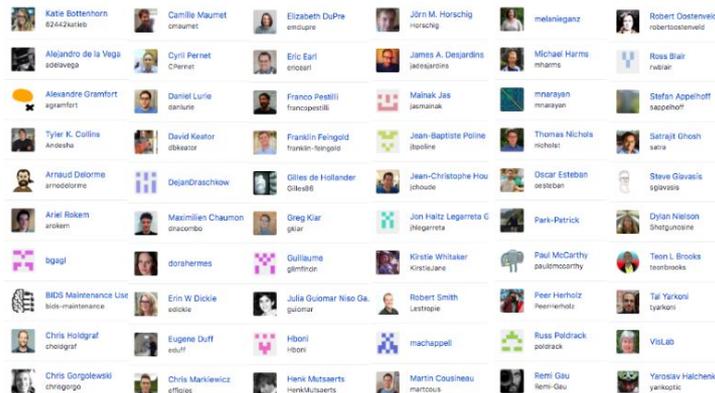
Chris Markiewicz
USA



Taylor Sato
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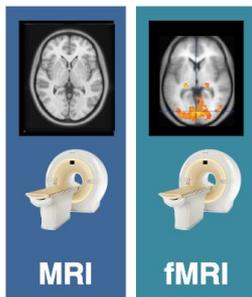


Franklin Feingold
USA



*only some contributors are shown

* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)



SCIENTIFIC DATA 

OPEN

SUBJECT CATEGORIES

- Data publication and archiving
- Research data

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

Received: 18 December 2015
Accepted: 19 May 2016
Published: 21 June 2016

Krzysztof J. Gorgolewski¹, Tibor Auern², Vince D. Calhoun^{3,4}, R. Cameron Craddock^{5,6}, Samir Das⁷, Eugene P. Duff⁸, Guillaume Flandin⁹, Sarajit S. Ghosh^{10,11}, Tristan Glatard¹², Yaroslav O. Halchenko¹³, Daniel A. Handwerker¹⁴, Michael Hanke^{15,16}, David Keator¹⁷, Xiangru Li¹⁸, Zachary Michael¹⁹, Camille Maurice²⁰, B. Nolan Michael^{21,22}, Thomas E. Nichols^{23,24}, John Poline²⁵, Jean-Baptiste Poline²⁶, Ariel Rokem²⁷, Guntmar Schaefer²⁸, Vanessa Sochat²⁷, William Toplett²⁷, Jessica A. Turner²⁸, Gaël Varoquaux²⁸ & Russell A. Poldrack²

(Gorgolewski et al. 2016)



SCIENTIFIC DATA 

OPEN

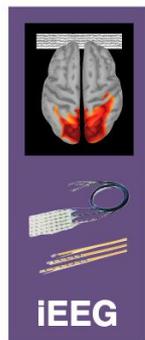
COMMENT

EEG-BIDS, an extension to the brain imaging data structure for electroencephalography

Received: 16 January 2019
Accepted: 7 May 2019
Published online: 25 June 2019

Cyril R. Pernet¹, Stefan Appelhoff², Krzysztof J. Gorgolewski³, Guillaume Flandin⁴, Christophe Phillips⁵, Arnaud Delorme^{6,7} & Robert Oostenveld^{8,9}

(Pernet et al., 2019)



SCIENTIFIC DATA 

OPEN

COMMENT

iEEG-BIDS, extending the Brain Imaging Data Structure specification to human intracranial electrophysiology

Received: 29 January 2019
Accepted: 24 May 2019
Published online: 25 June 2019

Christopher Holdgraf^{1,2}, Stefan Appelhoff³, Stephan Bickel⁴, Kristofer Bouchard⁵, Sasha D'Ambrosio⁶, Olivier David⁷, Orrin Devinsky⁸, Benjamin Dichter⁹, Adeen Flinker¹⁰, Brett L. Foster¹¹, Krzysztof J. Gorgolewski¹², Iris Groen¹³, David Groppa¹⁴, Aysegül Gunduz¹⁵, Liberty Hamilton¹⁶, Christopher J. Honey¹⁷, Mainak Jas¹⁸, Robert Knight¹⁹, Jean-Philippe Lachaux²⁰, Jonathan C. Lau²¹, Christopher Lee-Messer²², Brian N. Lundstrom²³, Kai J. Miller²⁴, Jeffrey G. Ojemann²⁵, Robert Oostenveld²⁶, Natalia Petridou²⁷, Gio Piantoni²⁸, Andrea Pigorini²⁹, Nader Pouratian³⁰, Nick F. Ramsey³¹, Arjen Stolk³², Nicole C. Swann³³, François Tadel^{34,35}, Bradley Voytek³⁶, Brian A. Wandell³⁷, Jonathan Winawer³⁸, Kirstie Whitaker^{39,40}, Lyuba Zehl⁴¹ & Dora Hermes^{42,43}

(Holdgraf et al., 2019)



SCIENTIFIC DATA 

OPEN

COMMENT: MEG-BIDS, the brain imaging data structure extended to magnetoencephalography

Received: 14 November 2017
Accepted: 3 May 2018
Published: 19 June 2018

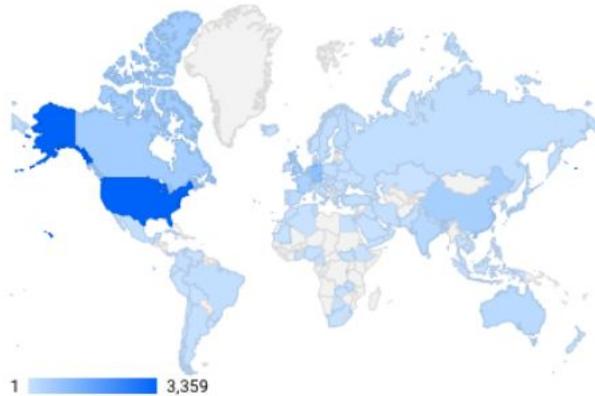
Guiomar Niso^{1,2}, Krzysztof J. Gorgolewski³, Elizabeth Bock⁴, Teon L. Brooks⁵, Guillaume Flandin⁶, Alexandre Gramfort^{7,8}, Richard N. Henson⁹, Mainak Jas¹⁰, Vladimir Litvak¹¹, Jeremy T. Moore¹², Robert Oostenveld¹³, Jan-Mathijs Schoffelen¹⁴, François Tadel^{15,16}, Joseph Wexler¹⁷ & Sylvain Baillet¹⁸

(Niso et al., 2018)

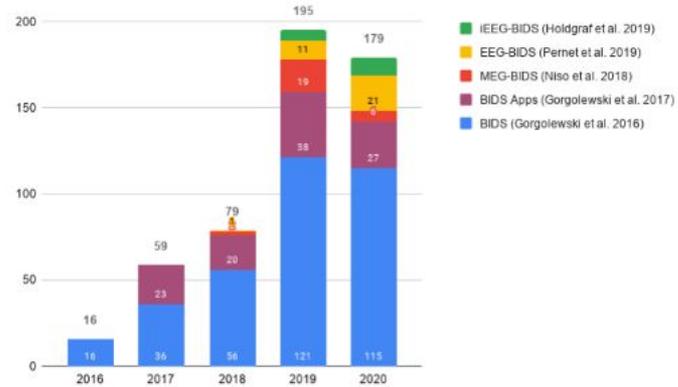
FAIR necessities - standardization - the idea of BIDS <https://bids-specification.readthedocs.io/en/stable/>

<https://bids.neuroimaging.io/>

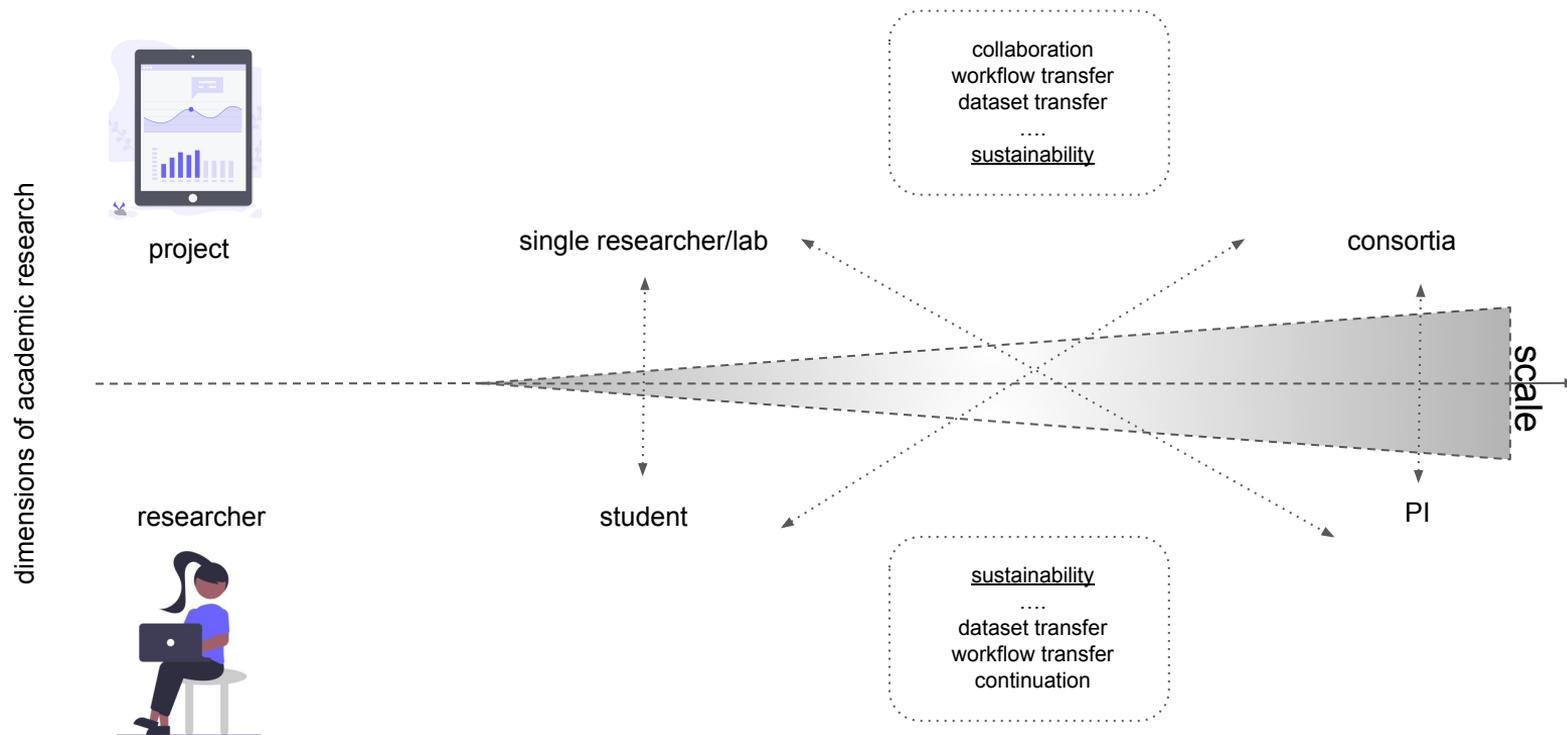
* content liberally borrowed from
Guiomar Niso's [BIDS ecosystem presentation](#)



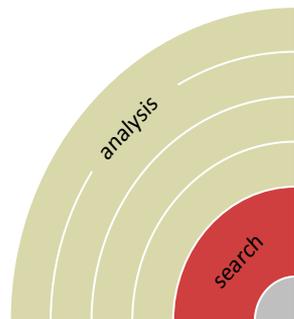
BIDS new citations per year



FAIR necessities - standardization - the benefits of BIDS



FAIR data through community-driven development of standards and beyond



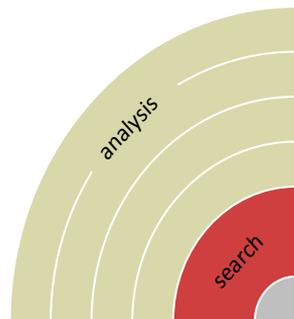
- necessities - documentation
- **necessities - virtualization (standardized)**
- **necessities - workflows (standardized)**
- **necessities - validation (standardized)**
- necessities - standardization
- prerequisites - data sharing (standardized)
- prerequisites - preregistration (standardized)



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR data through community-driven development of standards and beyond



- necessities - documentation
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Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

[DOI: 10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR necessities - standardization - the benefits of BIDS - data conversion

* content liberally borrowed from
Chris Markiewicz's [BIDS ecosystem presentation](https://bids-specification.readthedocs.io/en/stable/)

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



- vast amount of converters to convert raw data to BIDS in a (semi) automated and reproducible manner
- support for various modalities and input data types
- AFNI BIDS-tools, BIDS2ISATab, BIDS2NDA, bidsfy, bidskit, dac2bids, Data2Bids, Dcm2bids, heudiconv, DCM2NIfx, ReproIn, XNAT2BIDS, bids2xar, BIDS2NIDM, BIDScoin, MNE-BIDS...

```
dicomdir/  
├── 1208200617178_22/  
│   ├── 1208200617178_22_8973.dcm  
│   ├── 1208200617178_22_8943.dcm  
│   ├── 1208200617178_22_2973.dcm  
│   ├── 1208200617178_22_8923.dcm  
│   ├── 1208200617178_22_4473.dcm  
│   ├── 1208200617178_22_8783.dcm  
│   ├── 1208200617178_22_7328.dcm  
│   ├── 1208200617178_22_9264.dcm  
│   ├── 1208200617178_22_9967.dcm  
│   ├── 1208200617178_22_3894.dcm  
│   └── 1208200617178_22_3899.dcm  
├── 1208200617178_23/  
├── 1208200617178_24/  
└── 1208200617178_25/
```



```
my_dataset/  
├── participants.tsv  
├── sub-01/  
│   ├── anat/  
│   │   └── sub-01_T1w.nii.gz  
│   ├── func/  
│   │   ├── sub-01_task-rest_bold.nii.gz  
│   │   └── sub-01_task-rest_bold.json  
│   └── dwi/  
│       ├── sub-01_dwi.nii.gz  
│       ├── sub-01_dwi.json  
│       ├── sub-01_dwi.bval  
│       └── sub-01_dwi.bvec  
├── sub-02/  
├── sub-03/  
└── sub-04/
```

FAIR necessities - standardization - the benefits of BIDS - data validation

* content liberally borrowed from
Chris Markiewicz's [BIDS ecosystem presentation](#)

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

<https://bids.neuroimaging.io/>

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

- BIDS' machine readable structure makes automated validation possible and easy
- BIDS-validator does a comprehensive check
- for every single file and aspect in dataset:
 - warnings
 - errors
- fast and reliable integrity check

Select a [BIDS dataset](#) to validate

No file chosen

Note: Selecting a dataset only performs validation. Files are never uploaded.

ds00123

| Summary | Available Tasks | Available Modalities |
|---|--|--|
| <ul style="list-style-type: none">• 39 Files, 757.25MB• 4 - Subjects• 1 - Session | <ul style="list-style-type: none">• memory | <ul style="list-style-type: none">• bold• T1w |

Your dataset is not a valid BIDS dataset.

[view 1 error in 1 file](#)

[view 2 warnings in 3 files](#)

[Download error log for ds00123](#)

[Click to view details on BIDS specification](#)

FAIR necessities - standardization - the benefits of BIDS - data validation

* content liberally borrowed from
Chris Markiewicz's [BIDS ecosystem presentation](https://bids-specification.readthedocs.io/en/stable/)

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

<https://bids.neuroimaging.io/>

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

ds00123

| Summary | Available Tasks | Available Modalities |
|---|--|--|
| <ul style="list-style-type: none">39 Files, 757.25MB4 - Subjects1 - Session | <ul style="list-style-type: none">memory | <ul style="list-style-type: none">boldT1w |

Your dataset is not a valid BIDS dataset.

view 1 error in 1 file

Error: 1 1 file

Repetition time did not match between the scan's header and the associated JSON metadata file.

sub-04_ses-01_task-memory_run-03_bold.nii.gz 55354.461 KB | application/gzip

Location:

ds00123/sub-04/ses-01/func/sub-04_ses-01_task-memory_run-03_bold.nii.gz

Reason:

Repetition time defined in the JSON (2.007 sec.) did not match the one defined in the NIFTI header (1.99 sec.)

view 2 warnings in 3 files

ds00123

| Summary | Available Tasks | Available Modalities |
|---|--|--|
| <ul style="list-style-type: none">39 Files, 757.25MB4 - Subjects1 - Session | <ul style="list-style-type: none">memory | <ul style="list-style-type: none">boldT1w |

Your dataset is not a valid BIDS dataset.

view 1 error in 1 file

view 2 warnings in 3 files

Warning: 1 1 file

Not all subjects contain the same files. Each subject should contain the same number of files with the same naming unless some files are known to be missing.

sub-02_ses-01_T1w.nii.gz NaN KB |

Location:

/sub-02/ses-01/ana/sub-02_ses-01_T1w.nii.gz

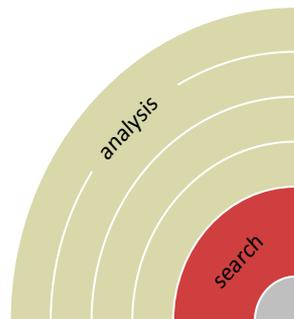
Reason:

This file is missing for subject sub-02, but is present for at least one other subject.

Warning: 2 2 files

Not all subjects/sessions/runs have the same scanning parameters.

FAIR data through community-driven development of standards and beyond



- necessities - documentation
- **necessities - virtualization (standardized)**
- **necessities - workflows (standardized)**
- necessities - validation (standardized)
- necessities - standardization
- prerequisites - data sharing (standardized)
- prerequisites - preregistration (standardized)



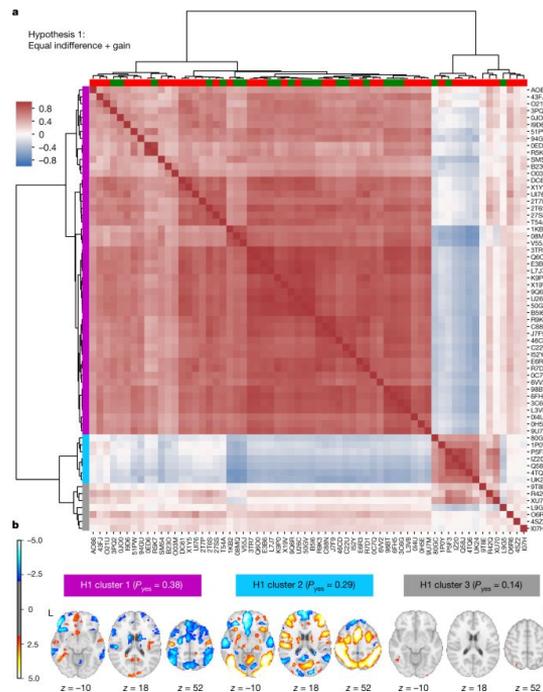
Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR necessities - standardization - the benefits of BIDS - data processing

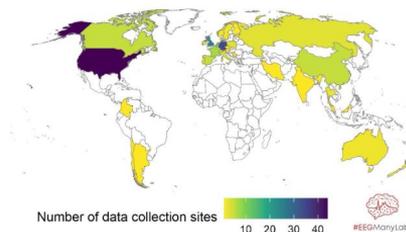
- a lot of sources of analyses/results variability
- every aspect of scientific workflow matters and therefore should be **FAIR**
- acquisition, datasets and metadata
- computing environment (OS, software package, version number, etc.)
- mega & multiverse analyses demonstrated the need for **FAIR** and **standardization**

Botvinik-Nezer et al. (2020)
The NARPS project



<https://www.nature.com/articles/s41586-020-2314-9>

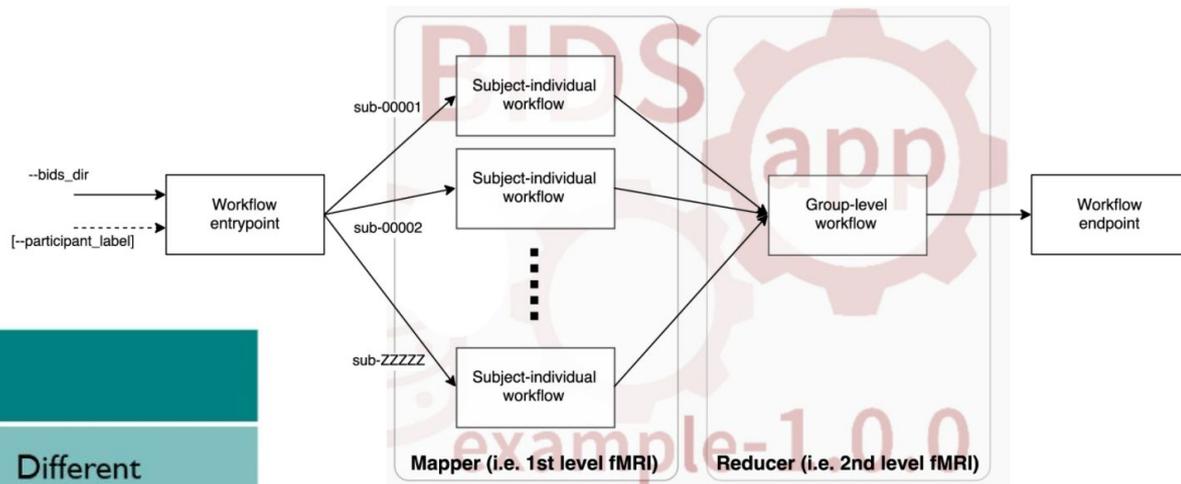
Pavlov et al. (2020)
#EEGManyLabs



<https://psyarxiv.com/528nr/>

FAIR necessities - standardization - the benefits of BIDS - data processing

* content liberally borrowed from
Chris Markiewicz's [BIDS ecosystem presentation](#)



From doi:10.1371/journal.pcbi.1005209.g002

| | | Data | |
|----------|-----------|--------------|---------------|
| | | Same | Different |
| Analysis | Same | Reproducible | Replicable |
| | Different | Robust | Generalisable |

From [The Turing Way](#), Ch. 2; doi:10.5281/zenodo.3233853

- BIDS Apps aims to address aspects of this problem via a common interface, container technologies and standardization
- open & reproducible analyses on (open) data
- each container/app includes a dedicated pipeline/workflow
- input: BIDS dataset
- pipeline/workflow: virtualization using containers
- output: BIDS conform standardized derivatives
- standardization + flexibility + reproducibility

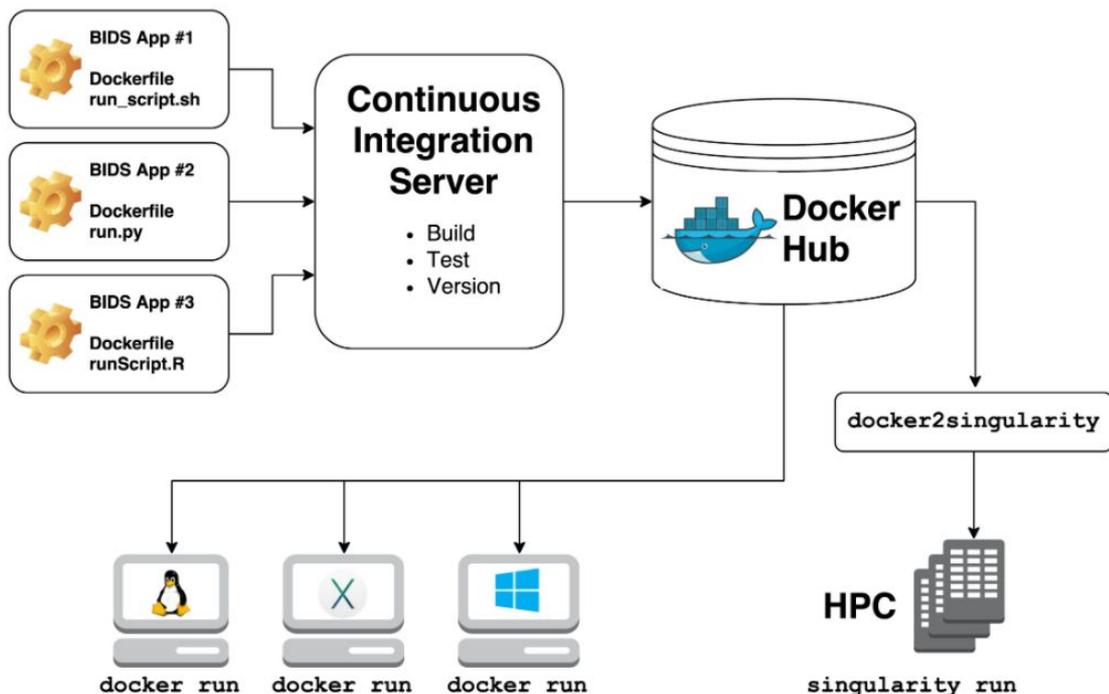
<https://bids-apps.neuroimaging.io/>

FAIR necessities - standardization - the benefits of BIDS - data processing

* content liberally borrowed from
Chris Markiewicz's [BIDS ecosystem presentation](https://bids-ecosystem.neuroimaging.io/)

<https://bids-ecosystem.neuroimaging.io/>

- containerized pipelines/workflow that understand BIDS datasets
- set of core command line arguments
- should not depend on software outside the container
- openly hosted on docker/singularity hub and include version tags
- targeting automated and standardized analyses that are reproducible



From doi:10.1371/journal.pcbi.1005209.g001

FAIR necessities - standardization - the benefits of BIDS - data processing



BIDS Apps

portable neuroimaging pipelines that understand BIDS datasets

[About](#) [Tutorials](#) [Apps](#)

Available BIDS Apps

<https://bids-apps.neuroimaging.io/>

BIDS-Apps/example
BIDS-Apps/freesurfer
BIDS-Apps/ndmg
BIDS-Apps/BROCCOLI
BIDS-Apps/FibreDensityAndCrosssection
BIDS-Apps/SPM
poldracklab/mriqc
BIDS-Apps/QAP
BIDS-Apps/CPAC
BIDS-Apps/hyperalignment
BIDS-Apps/mindboggle
BIDS-Apps/MRtrix3_connectome
BIDS-Apps/rs_signal_extract
BIDS-Apps/aa
BIDS-Apps/niak
BIDS-Apps/oppni
poldracklab/fmriprep
BIDS-Apps/brainiak-srm
BIDS-Apps/nipypelines
BIDS-Apps/HCPPIpipelines
BIDS-Apps/MAGeTbrain
BIDS-Apps/tracula
BIDS-Apps/baracus
BIDS-Apps/antsCorticalThickness

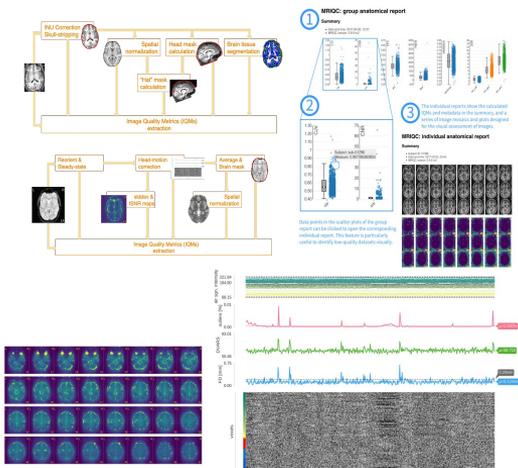
| | | | | | | |
|--------------------|-----------------|-----------------|--------------------------|-------------------|---------|------------|
| version v0.0.7 | open issues 5 | circled passing | open bug pull requests 0 | docker pulls 14x | 439.5MB | 23 layers |
| version v6.0.1-6.1 | open issues 17 | circled passing | open bug pull requests 0 | docker pulls 11x | 2.6GB | 52 layers |
| version v0.1.0 | open issues 1 | circled passing | open bug pull requests 0 | docker pulls 8x | 920.9MB | 31 layers |
| version v1.0.1 | open issues 8 | circled passing | open bug pull requests 0 | docker pulls 522 | 3GB | 21 layers |
| version v0.0.1 | open issues 1 | circled passing | open bug pull requests 0 | docker pulls 250 | 576.8MB | 31 layers |
| version v0.0.20 | open issues 3 | circled passing | open bug pull requests 0 | docker pulls 1.5k | 7GB | 24 layers |
| version v0.19.1 | open issues 125 | circled passing | open bug pull requests 0 | docker pulls 69x | 3GB | 41 layers |
| version v0.0.1 | open issues 5 | circled passing | open bug pull requests 0 | docker pulls 7 | Image | not found |
| version v1.0.1a_22 | open issues 10 | circled passing | open bug pull requests 0 | docker pulls 3.2k | 1.6GB | 45 layers |
| version v0.0.5 | open issues 0 | circled passing | open bug pull requests 0 | docker pulls 169 | Image | not found |
| version v0.0.4-1 | open issues 8 | circled passing | open bug pull requests 0 | docker pulls 752 | 1.9GB | 81 layers |
| version v0.4.2 | open issues 3x | circled passing | open bug pull requests 0 | docker pulls 1.4k | 7.3GB | 56 layers |
| version v0.1 | open issues 0 | circled passing | open bug pull requests 0 | docker pulls 238 | 240MB | 17 layers |
| version v0.2.0 | open issues 2 | circled passing | open bug pull requests 0 | docker pulls 272 | 6.3GB | 30 layers |
| version v1.0 | open issues 3 | circled passing | open bug pull requests 0 | docker pulls 279 | 2.7GB | 103 layers |
| version v0.7.0-1 | open issues 5 | circled passing | open bug pull requests 0 | docker pulls 315 | 2.9GB | 41 layers |
| version v20.2.0 | open issues 192 | circled passing | open bug pull requests 0 | docker pulls 376x | 5GB | 49 layers |
| version initial | open issues 0 | circled failing | open bug pull requests 0 | docker pulls 207 | 559.3MB | 13 layers |
| version v0.3.0 | open issues 0 | circled passing | open bug pull requests 0 | docker pulls 381 | 478.1MB | 20 layers |
| version v4.1.3 | open issues 7 | circled passing | open bug pull requests 0 | docker pulls 1.9k | 4GB | 31 layers |
| version v0.3.1 | open issues 2 | circled passing | open bug pull requests 0 | docker pulls 466 | Image | not found |
| version v6.0.0-5 | open issues 0 | circled passing | open bug pull requests 0 | docker pulls 689 | 3.4GB | 57 layers |
| version v1.1.4 | open issues 2 | circled passing | open bug pull requests 0 | docker pulls 1.2k | Image | not found |
| version v2.2.0-1 | open issues 0 | circled passing | open bug pull requests 0 | docker pulls 207 | 351.9MB | 21 layers |



FAIR necessities - standardization - the benefits of BIDS - data processing

quality control (MRIQC)

mriqc data/ output/ --specifications

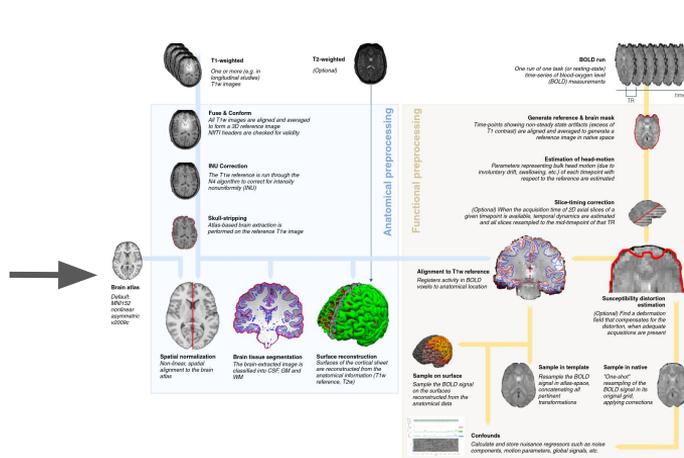


[Esteban et al. \(2017\),](https://mriqc.readthedocs.io/en/stable/index.html)

<https://mriqc.readthedocs.io/en/stable/index.html>

preprocessing (fmriprep)

fmriprep data/ output/ --specifications

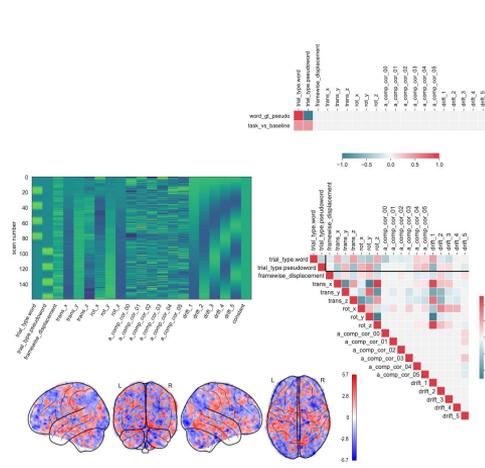


[Esteban et al. \(2019\),](https://fmriprep.org/en/stable/)

<https://fmriprep.org/en/stable/>

statistical analyses (fitlins)

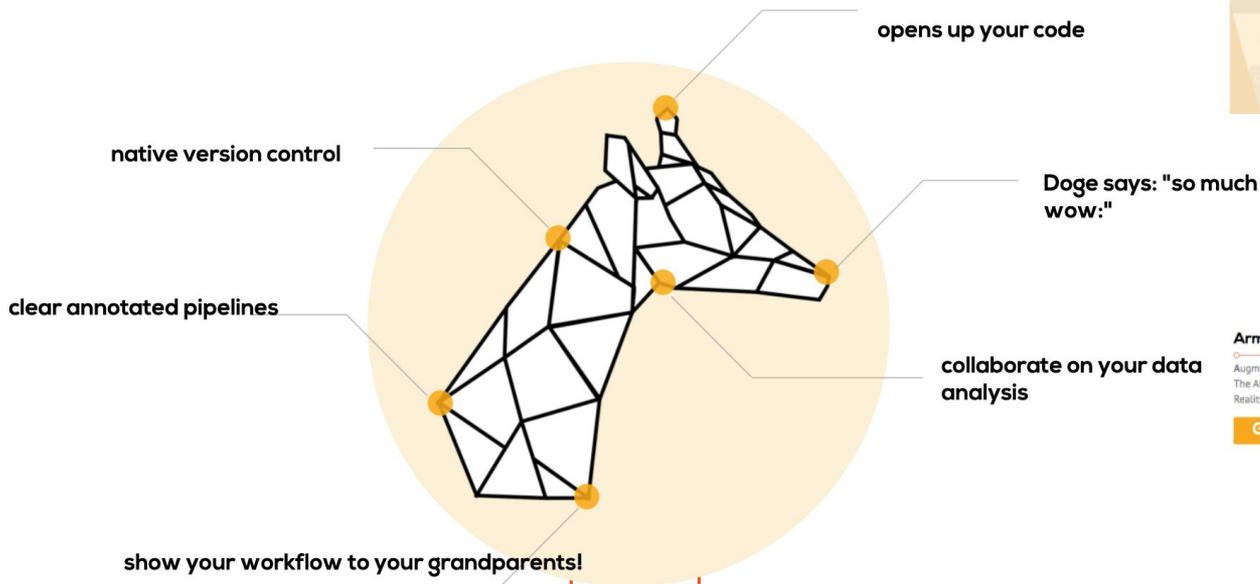
fitlins data/ output/ --specifications



<https://fitlins.readthedocs.io/en/latest/index.html>

FAIR necessities - standardization - the idea of data processing

<https://giraffe.tools/>



Porcupine

Porcupine Creates U- Pipeline
With the Porcupine workflow editor, you can visually build your pipeline!

[Get started!](#)

Armadillo

Augmented Reality [madillo]
The ARmadillo web app creates 3D Augmented Reality images from the Neurovault database

[Get started!](#)



A platform for fast and flexible re-analysis of (naturalistic) fMRI studies

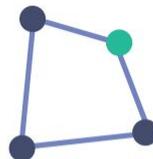
Browse public analyses

Re-use public data



Select from openly available naturalistic fMRI datasets, from sources such as [OpenNeuro](#) and [DataLad](#).

Design your analysis



Browse hundreds of annotations automatically extracted from stimuli using **state-of-the-art machine learning** algorithms, such as Google Cloud Vision, IBM Watson, and more.

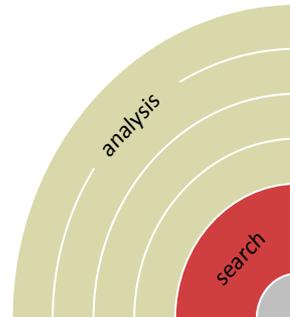
Execute and share



Portable BIDS pipelines enable execution with no configuration. Results are automatically uploaded to [NeuroVault](#) for easy sharing.

Learn more

FAIR data through community-driven development of standards and beyond



- **necessities - documentation**
- necessities - virtualization (standardized)
- necessities - workflows (standardized)

- necessities - standardization
- prerequisites - data sharing (standardized)
- prerequisites - preregistration (standardized)



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR necessities - analyses documentation - Jupyter

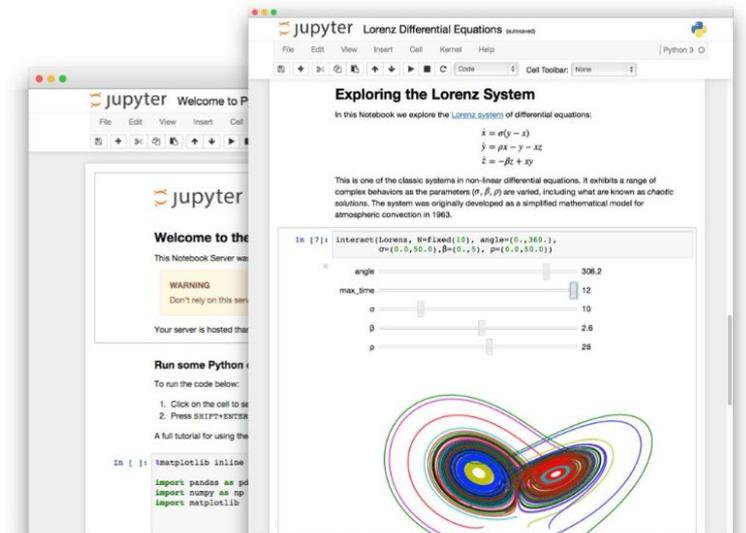
*adapted from [Fernando Perez](#)
and [Elizabeth DuPre](#)



- a **community** of people
- an **ecosystem** of **open tools** and **standards** for **interactive computing**
- **language-agnostic** and **modular**
- **empower people** to use other **open tools**

FAIR necessities - analyses documentation - Jupyter notebooks

*adapted from [Fernando Perez](#)
and [Elizabeth DuPre](#)

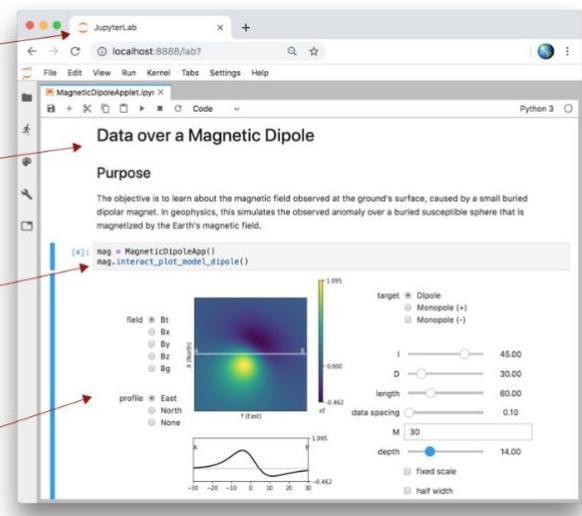


Runs in web browser

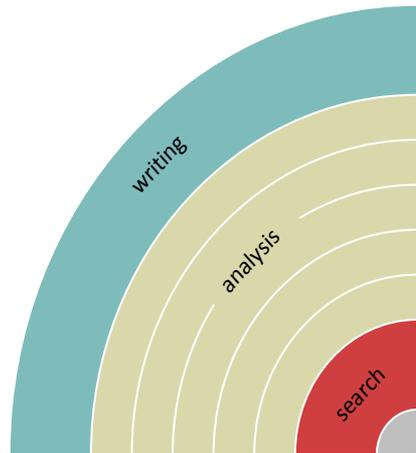
Text, comments, equations

Lines of code
(Python, Julia, R, Matlab, ...)

Output: values, images, plots,
tables, interactive widgets



FAIR data through community-driven development of standards and beyond



- **necessities - reporting (standardized)**
- necessities - documentation
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- necessities - workflows (standardized)
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FAIR data through community-driven development of standards and beyond



- necessities - repositories (standardized)
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FAIR necessities - more than a PDF - binder



Turn a Git repo into a collection of interactive notebooks

Have a repository full of Jupyter notebooks? With Binder, open those notebooks in an executable environment, making your code immediately reproducible by anyone, anywhere.

New to Binder? Get started with a Zero-to-Binder tutorial in Julia, Python or R.

Build and launch a repository

GitHub repository name or URL

GitHub

Git ref (branch, tag, or commit) Path to a notebook file (optional)

HEAD File

Copy the URL below and share your Binder with others:

Copy the text below, then paste into your README to show a binder badge:  [launch binder](#)

<https://mybinder.org/>



Explicit Dependencies



*[adapted from Fernando Perez](#)

FAIR necessities - more than a PDF - Jupyter Books

jupyter {book}

Search this book...

GET STARTED

Overview

Build your book

Publish your book online

Configure book settings

Table of contents structure

Types of content source files

WRITE BOOK CONTENT

MyST Markdown overview

Special content blocks

References and citations

Math and equations

Images and figures

Control the page layout

Execute and cache your pages

Formatting code outputs

MAKE YOUR BOOK INTERACTIVE

Launch buttons for interactivity

Hide or remove content

Interactive data visualizations

Commenting and annotating

ADVANCED AND MISCELLANEOUS

PDFs for your book

Custom Sphinx configuration

How-to and FAQ

Contribute to Jupyter Book

REFERENCE

Gallery of Jupyter Books

MyST cheat sheet

The command-line interface

Glossary

Change log

Powered by Jupyter Book



Contents

Books with Jupyter

Jupyter Book is an open source project for building beautiful, publication-quality books and documents from computational material.

Here are some of the features of Jupyter Book:

- ✓ [Write publication-quality content in Markdown](#)
You can write in either Jupyter Markdown, or an extended flavor of Markdown with [publishing features](#). This includes support for rich syntax such as [citations](#) and [cross-references](#), [math](#) and [equations](#), and [figures](#).
- ✓ [Write content in Jupyter Notebook](#)
This allows you to include your code and outputs in your book. You can also write notebooks entirely in Markdown that get executed when you build your book.
- ✓ [Execute and cache your book's content](#)
For `.jupyter` and Markdown notebooks, execute code and insert the latest outputs into your book. In addition, [cache](#) and [re-use](#) outputs to be used later.
- ✓ [Insert notebook outputs into your content](#)
Generate outputs as you build your documentation, and insert them in-line with your content across pages.
- ✓ [Add interactivity to your book](#)
You can [toggle cell visibility](#), include [interactive outputs](#) from Jupyter, and [connect with online services](#) like Binder.
- ✓ [Generate a variety of outputs](#)
This includes single- and multi-page websites, as well as [PDF outputs](#).
- ✓ [Build books with a simple command-line interface](#)
You can quickly generate your books with one command, like so: `jupyter-book build mybook/`

This website is built with Jupyter Book! You can browse its contents to the left to see what is possible.

Get involved with Jupyter Book!

Jupyter Book is an open community that welcomes your feedback, input, and contributions!

- [Open an issue](#)
to provide feedback and new ideas, and to help others.
- [Vote for new features](#)
by adding a 🗳️ to issues you'd like to see completed.
- [Contribute to Jupyter Book](#)
by following our contributing guidelines and finding an issue to work on. See the [feature voting leaderboard](#) for inspiration.

Install Jupyter Book

You can install Jupyter Book [via pip](#):

```
pip install -U jupyter-book
```

This will install everything you need to build a Jupyter Book locally.

<https://jupyterbook.org/>

FAIR necessities - more than a PDF - Jupyter + binder



Supported by the Canadian Open Neuroscience Platform (CONP).

NeuroLibre is a curated repository of interactive neuroscience notebooks.

Committed to publishing curated Jupyter notebooks with zero article processing charges or subscription fees.

[Submit a paper to NeuroLibre](#)

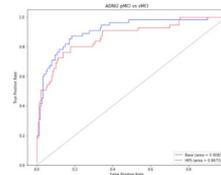
- [Explore Papers](#)
- [Documentation](#)
- [Learn More](#)

<https://www.neurolibre.com/>

A highly predictive signature (HPS) of Alzheimer's disease dementia from cognitive and structural brain features

A jupyter notebook containing analyses that give a highly predictive signature (HPS) of Alzheimer's disease dementia from cognitive and structural features using simulated data.

[NeuroLibre Book](#) [GitHub Code](#)



Introduction to machine learning with Nilearn

An introductory tutorial for using the popular Nilearn software package to perform machine learning analyses with neuroimaging data. This material is adapted from the Montreal AI and Neuroscience (MAIN) 2018 workshops.

[NeuroLibre Book](#) [GitHub Code](#)



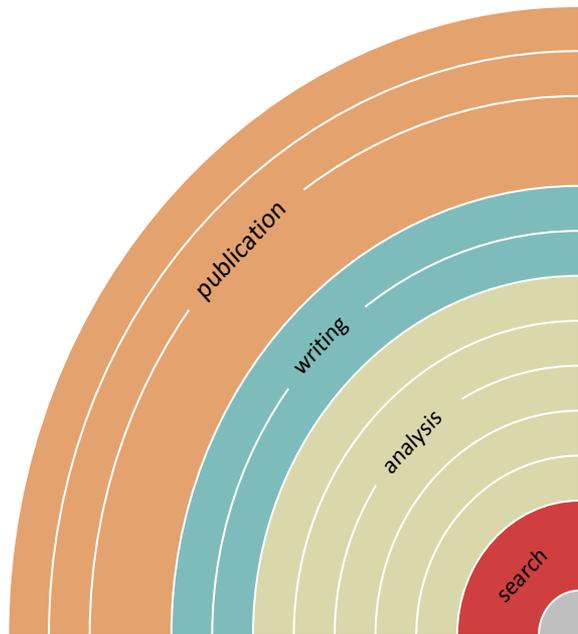
Image processing with Spinal Cord Toolbox (SCT)

This notebook presents an example analysis pipeline using the Spinal Cord Toolbox (SCT), a suite of tools specialized for analysis of spinal cord MRI images of the spinal. Topics covered include: segmentation, masking, registration, warping, and quantitative metric computation. This tutorial was generated in a Jupyter Notebook and coded in Python.

[NeuroLibre Book](#) [GitHub Code](#)



FAIR data through community-driven development of standards and beyond



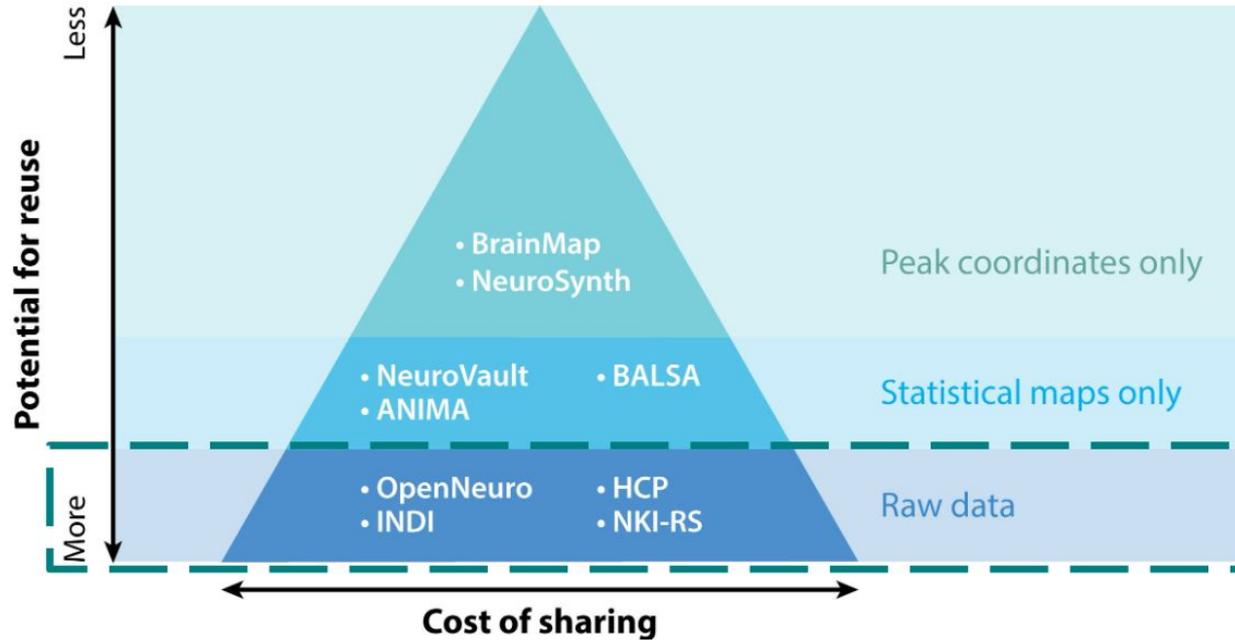
- **necessities - repositories (standardized)**
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FAIR data through community-driven development of standards and beyond



*[adapted from Russ Poldrack](#)

FAIR necessities - repositories (standardized)



OpenNEURO

A free and open platform for sharing MRI,
MEG, EEG, iEEG, and ECoG data



[Browse All Public Datasets](#)

436

Public Datasets

14277

Participants



Get Data

Browse and download datasets from contributors all over the world.



Share Data

Upload your data to an NIH Brain Initiative approved repository.



Use Data

Use our affiliated website to process applicable data.

<https://openneuro.org/>



NEUROVAULT

A public repository of unthresholded statistical maps, parcellations, and atlases of the brain.

What is it?

A place where researchers can publicly store and share unthresholded statistical maps, parcellations, and atlases produced by MRI and PET studies.

Why use it?

- Interactive visualization
- A permanent URL
- Publicly shareable
- Improves meta-analyses

Supported by



[Get started and upload an image!](#)

NeuroVault Collections ▾ Metaanalyses ▾ About ▾ herholz.peer ▾ Search Search

Improvising at rest: Differentiating jazz and classical music training with resting state functional connectivity

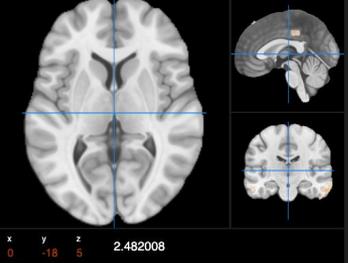
Description: Improvisationally trained musicians, Classically trained musicians, and the Minimally Musically Trained (MMT) in seed-based functional connectivity and network analyses in resting state functional MRI.

Related article: Belden, A., Zeng, T., Przynsinda, E., Anteraper, S. A., Whitfield-Gabrieli, S., & Loui, P. (2020). Improvising at rest: Differentiating jazz and classical music training with resting state functional connectivity. *NeuroImage*, 207, 116384. doi:10.1016/j.neuroimage.2019.116384

[Source data:](#)

[3D View](#) [Download](#)

File View Settings Help 



x y z 2.482008
0 -18 5

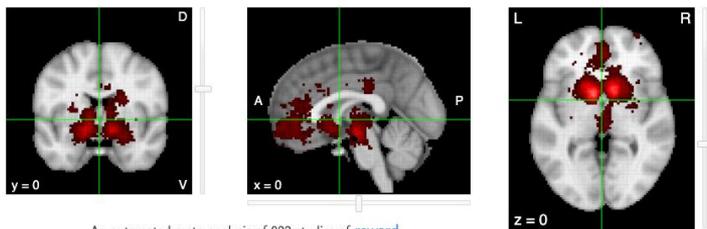
| Group | Metadata | | |
|----------------|------------------------------|---------------------------|-------|
| Show 7 entries | Search: <input type="text"/> | | |
| View | ID | Name | Type |
| | 133730 | Class LECN spmT 0001 | T map |
| | 133731 | Class Precun spmT 0001 | T map |
| | 133732 | Class RECN spmT 0001 | T map |
| | 133733 | Class Vis spmT 0001 | T map |
| | 133734 | Class pSalience spmT 0001 | T map |
| | 133735 | Class vDMN spmT 0001 | T map |
| | 133736 | Imp LECN spmT 0001 | T map |

Showing 1 to 7 of 76 entries [First](#) [Previous](#) [Next](#) [Last](#)

neurosynth.org

Neurosynth is a platform for large-scale, automated synthesis of functional magnetic resonance imaging (fMRI) data.

It takes thousands of published articles reporting the results of fMRI studies, chews on them for a bit, and then spits out images that look like this:



An automated meta-analysis of 922 studies of [reward](#)

Database Status

507891 activations reported in [14371 studies](#)

Interactive, downloadable meta-analyses of [1335 terms](#)

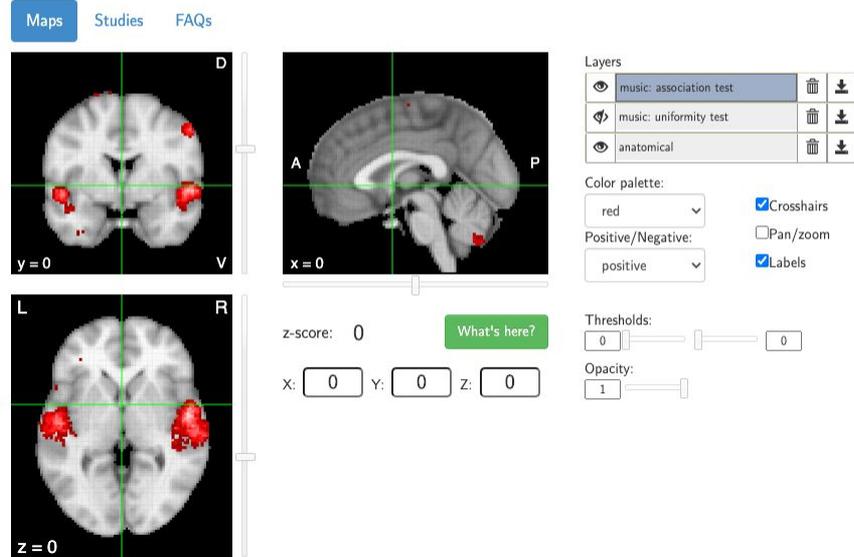
Functional connectivity and coactivation maps for over [150,000 brain locations](#)

music

An automated meta-analysis of 163 studies

Search for another term:

Maps Studies FAQs



Layers

| | | | |
|-------------------------------------|-------------------------|--------------------------|--------------------------|
| <input checked="" type="checkbox"/> | music: association test | <input type="checkbox"/> | <input type="checkbox"/> |
| <input checked="" type="checkbox"/> | music: uniformity test | <input type="checkbox"/> | <input type="checkbox"/> |
| <input checked="" type="checkbox"/> | anatomical | <input type="checkbox"/> | <input type="checkbox"/> |

Color palette: red

Positive/Negative: positive

Thresholds: 0

Opacity: 1

z-score: 0 [What's here?](#)

X: 0 Y: 0 Z: 0

Crosshairs
 Pan/zoom
 Labels

FAIR necessities - repositories (standardized)

<https://neuroquery.org/>

NeuroQuery [Dockès et al 2020] Brain maps by querying the neuroscience literature

A query on neuroscience, cognition, or brain pathologies

music

Click to edit. [Edit query](#)

Terms related to the query

| In query | Term | Similarity | Weight in brain map | N |
|--------------|------------------------|------------|---------------------|-------|
| | music | | | 623 |
| In expansion | auditory | | | 4009 |
| | temporal | | | 11897 |
| | auditory cortex | | | 846 |
| | cerebellum | | | 5578 |
| | premotor | | | 3200 |
| | motor | | | 7928 |
| | sound | | | 2261 |
| | parahippocampal | | | 3512 |
| | superior | | | 9978 |
| | matter | | | 6324 |
| | right | | | 13076 |

Show 12 entries Showing 1 to 12 of 76 entries



Publications related to the query

[Separate cortical networks involved in music perception: preliminary functional MRI evidence for modularity of music processing](#)

[Music and Emotions in the Brain: Familiarity Matters](#)

[It don't mean a thing...](#)

[A Functional MRI Study of Happy and Sad Emotions in Music with and without Lyrics](#)

[The rewards of music listening: Response and physiological connectivity of the mesolimbic system](#)

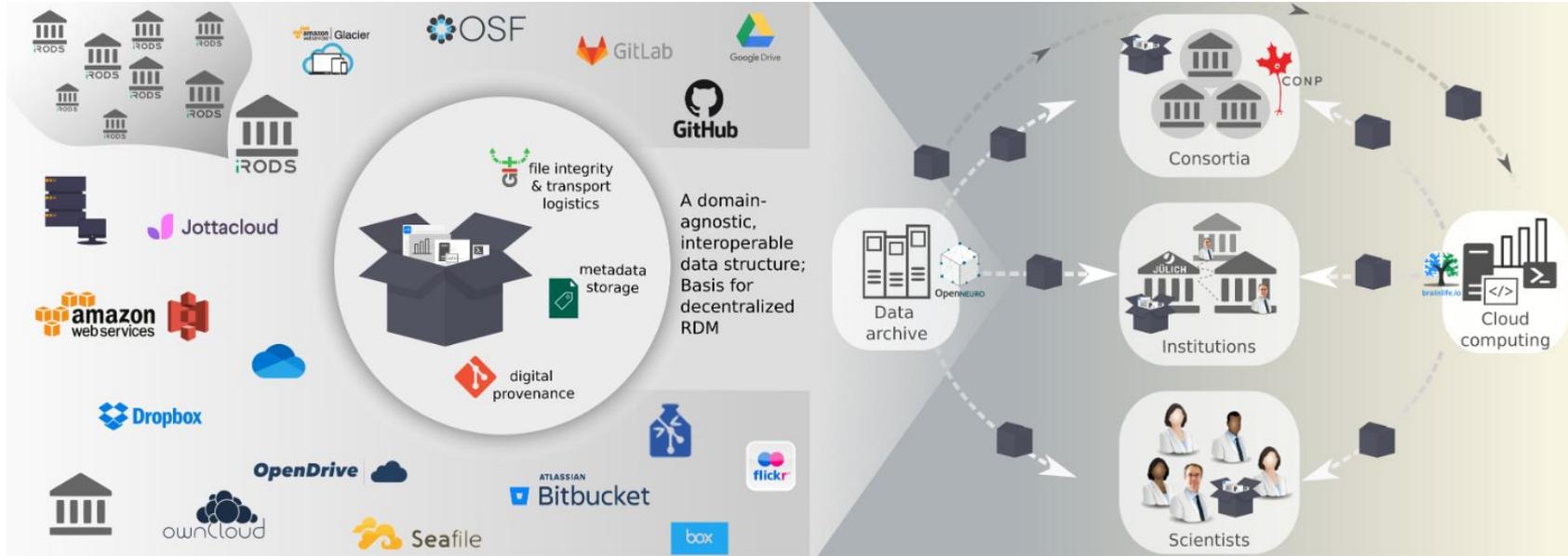
[Moving to Music: Effects of Heard and Imagined Musical Cues on Movement-Related Brain Activity](#)

FAIR data management

*adapted from [Adina Wagner](#)



<https://www.datalad.org>
<http://handbook.datalad.org>



Discover Data

DataLad has built-in support for **metadata** extraction and **search**. With only a few steps, you can search through a large collection of readily available datasets and immediately download them. [See more...](#)

Consume Data

DataLad offers direct **access to individual files** — great when you only need a few files from some large datasets for an analysis. Files in a dataset can be distributed across multiple download sources with tailored permissions to match your **data privacy** needs. [See more...](#)

Publish Data

DataLad supports sharing datasets with the **public or just some colleagues** on platforms that you are using already — **no need for a central service**. You have complete freedom to share your work in multiple platforms simultaneously (your own server, DropBox, GitHub, etc.) without losing track. [See more...](#)

Reproducibility

DataLad provides **joint management of analysis code and data**. This enables you to comprehensively track the exact state of any analysis inputs that produced your results — across the entire lifetime of a project, and across multiple datasets. [See more...](#)

Data Portal

The DataLad project operates a crawler that regularly indexes datasets from scientific data portals such as [OpenfMRI](#) and [CRCNS](#), making them trivial to acquire and work with using DataLad. Take a look at the [available datasets](#).

Community

Have a question? The primary support forum for [DataLad is on Neurostars](#). If you're having a problem with DataLad, you can use this [pre-filled form](#) to help you report your problem. Just wanna chat? We are hanging out in our [matrix chatroom](#). Come join us.



 Star 53

The DataLad handbook will supply you with everything you need to get started and break new grounds with DataLad.



DataLad

The Handbook

Welcome!

This handbook is a living resource about why and — more importantly — *how* to use DataLad. It aims to provide novices and advanced users of all backgrounds with both the basics of DataLad and start-to-end use cases of specific applications. If you want to get hands-on experience and learn DataLad, the *Basics* part of this book will teach you. If you want to know what is possible, the *use cases* will show you. And if you want to help others to get started with DataLad, the [companion repository](#) provides [free and open source teaching material](#) tailored to the handbook.

Before you read on, please note that the handbook is based on **DataLad version 0.12**, but the section [Installation and configuration](#) will set you up with what you need if you currently do not have DataLad 0.12 or higher installed.

If you're new here, please start the handbook [here](#). Alternatively, try to identify with one of several user-types in this [user specific guide to the handbook](#).

Important:

The handbook is currently in beta stage. If you would be willing to provide feedback on its contents, please [get in touch](#).

Contributors

This guide is the result of the collaboration of many people, and your contributions are welcome!

Useful Links

DataLad Website
Developer Docs
DataLad@GitHub
Handbook@GitHub
Frequently Asked Questions
Handbook Index
DataLad cheat sheet

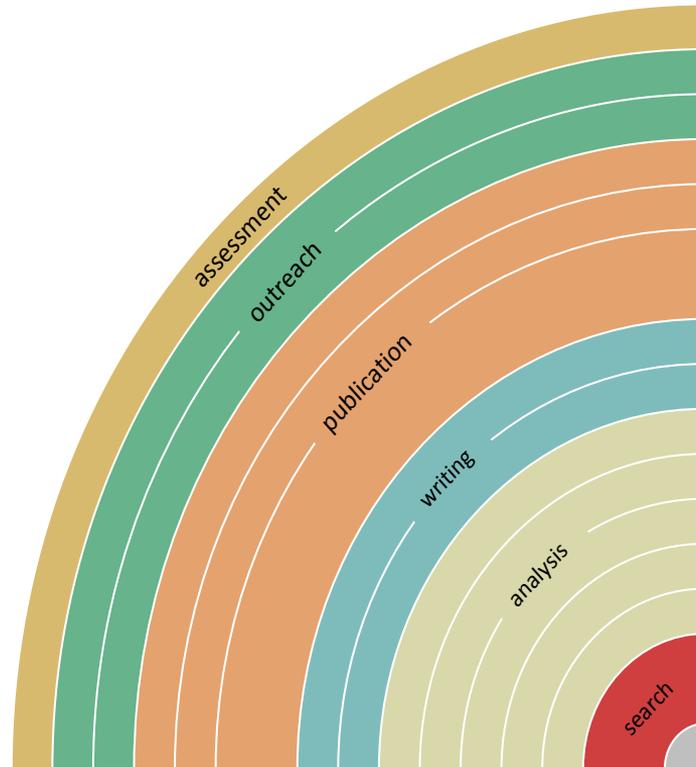


Feedback

We highly appreciate your feedback on the handbook and on DataLad



FAIR data through community-driven development of standards and beyond



- necessities - support
- prerequisites - initiatives
- necessities - repositories (standardized)
- necessities - more than a PDF
- necessities - reporting (standardized)
- necessities - documentation
- necessities - virtualization (standardized)
- necessities - workflows (standardized)
- necessities - validation (standardized)
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Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)

FAIR initiatives & support

Project T(eaching) I(ntegrity in) E(mpirical) R(eseach)

Raw Data

A copy of **every original data file** from which you extract any of data used in your study.

Your original data files **serve as a record** of the data you began the project with.

Analysis Code + Analysis Data

One or more files containing code used for the study... should **execute all the data processing and analysis necessary to replicate the study** and reproduce the reported results

Documents

Documentation to understand the study

- A copy of your final paper
- Your Data Appendix
- Your README file

<https://www.projecttier.org/>

The Turing Way



https://zenodo.org/record/3695300#_X8q1_6pkJLJ

<https://the-turing-way.netlify.app/>



ReproNim: A Center for Reproducible Neuroimaging Computation

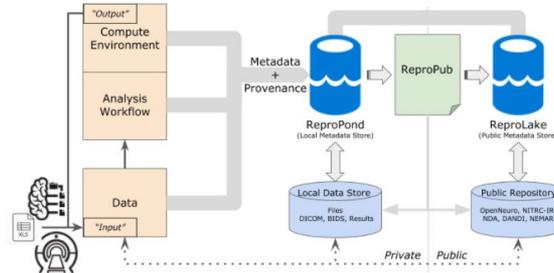
Welcome to ReproNim!

ReproNim's goal is to improve the reproducibility of neuroimaging science and extend the value of our national investment in neuroimaging research, while making the process easier and more efficient for investigators.

ReproNim delivers a reproducible analysis framework comprised of components that include:

- data and software discovery
- implementation of standardized description of data, results and workflows
- development of execution options that facilitates operation in all computational environments
- provide training and education to the community

All components of the framework are intended to foster continued use and development of the reproducible and generalizable framework in neuroimaging research. See our [Blog Post](#) for more discussion of ReproLake and ReproPond.



Welcome to INCF Neurostars.

A question and answer forum for neuroscience researchers, infrastructure providers and software developers.

How to start

If you are here for the first time, say hello and [introduce yourself!](#) Please visit the ["Welcome"](#) post to learn how to use the forum.

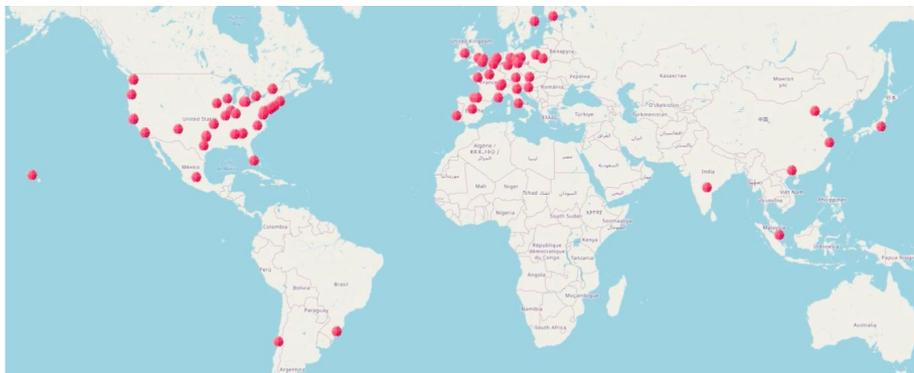
Important links

- View free online neuroscience courses and tutorials, visit [TrainingSpace](#)
- Looking for sample datasets, visit [KnowledgeSpace](#)
- Want to join the open neuroscience revolution, visit [INCF.org](#)

| all categories ▾ | Categories | Latest | Top | My Posts | Bookmarks | + New Topic |
|--|------------|----------|-----|----------|--|-------------|
| Category | | Topics | | | Latest | |
| Neuro Questions | | 2898 | | | | |
| This is a place to ask any and all questions about Neuroscience. | | 3 new | | |  Datalad on github + cluster storage • datalad | 2 42m |
| Announcements | | 162 | | |  BIDS and cookiecutter? • bids | 0 1h |
| Announcements is the place to post some general information that is of interest to the community. It can be a new project, a software release, a job posting, or any information you want to share that's not a "question". | | 2 new | | | OCNS | |
| Community Chat | | 10 | | |  Infrastructure/software/tools SIG: meet and greet, initial discussions | 64 4h |
| A place for friendly conversation, ideas, discussion, and otherwise un-categorizable things in the NeuroStars community. | | 1 unread | | |  OCNS cns2020 | |
| Institutions | | 37 | | |  Seed map - HELP from a newbie | 0 6h |
| Category for Institutions that are partnered with INCF to discuss the institution-specific topics. | | | | | ■ Neuromatch Academy projects | |
| ABCD ReproNim | | 13 | | | Deadline extended to Nov. 15th | |
| The ABCD-ReproNim Course provides training for reproducible analyses of the Adolescent Brain Cognitive Development (ABCD) Study® data. Course starts October 16, 2020. Course website: https://www.abcd-repronim.org | | 3 new | | |  Postdoctoral position for the development of acquisition, storage and processing pipelines for reproducible science building on the BIDS standard. • | 0 20h |
| | | | | | ■ Announcements jobs | |



<http://brainhack.org/>

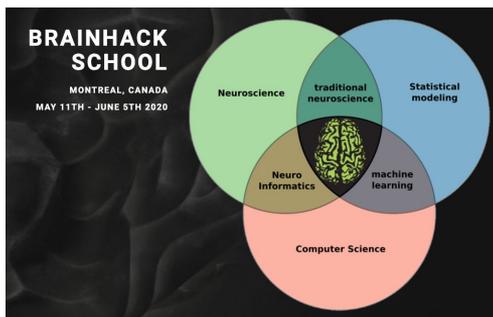


- started in 2012, spread across 25 countries, over 140 events, over 4000 members
- over 1000 attendees annually, focus on ECRs-collaboration-community-hacking-teaching
- model for various large scale teaching initiatives

The image shows a screenshot of the Brainhack Global 2020 event page. The page has a dark green header with navigation links: Home, Events, Projects, Code of Conduct, Team, Contact, and FAQ. The main content area features the text "Brainhack Global 2020 is happening!" and "November 30th - December 13th". There is a button that says "Host your own local Brainhack" with a location pin icon. Below this is a section titled "What is Brainhack?" with a description of the event format and a link to "2020 being 2020, this year's Brainhack Global will unite several virtual regional Brainhack events throughout the world." The page also features a stylized brain graphic and a footer with social media icons.

FAIR initiatives & support

- **open** to everyone interested
- mostly **free** (if not, dedicated funding to support individuals from underrepresented and marginalized groups exists)
- **all materials publicly available** (including video recordings of lectures)
- **comprehensive, application oriented training**
- **decentralized and digital**
- **individual support and supervision**



<https://school.brainhackmtl.org/>

A screenshot of the NeuroHackAcademy website. The header includes the title 'NEUROHACKADEMY' and an 'APPLY' button. The navigation menu contains 'Home', 'Courses', 'Materials', 'Apply', and 'Contact'. The main content area shows brain scan images with neural network overlays. Text on the page describes the course as a summer school in neuroimaging and data science, held at the University of Washington eScience Institute, and provides details about the course schedule and application process.

<https://neurohackademy.org/>

A screenshot of the Neuromatch Academy website. The title is 'Neuromatch Academy'. Below the title, it states 'An online school for Computational Neuroscience'. Further down, it mentions 'Started in response to the COVID-19 pandemic, Neuromatch Academy is a non-profit course in computational neuroscience offered online from July 13-31, 2020.' The bottom of the screenshot features the Neuromatch Academy logo, which consists of a stylized brain icon and the text 'neuromatch academy'.

<https://www.neuromatchacademy.org/>

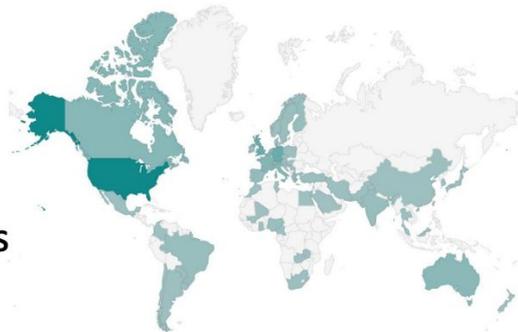
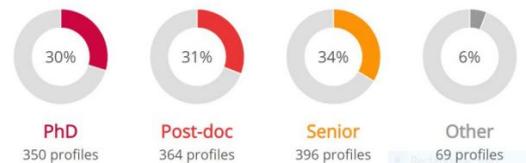


Repository for Women in Neuroscience

- www.winrepo.org
- over 1,100 profiles
- easy search
- recommendations

Support the project:

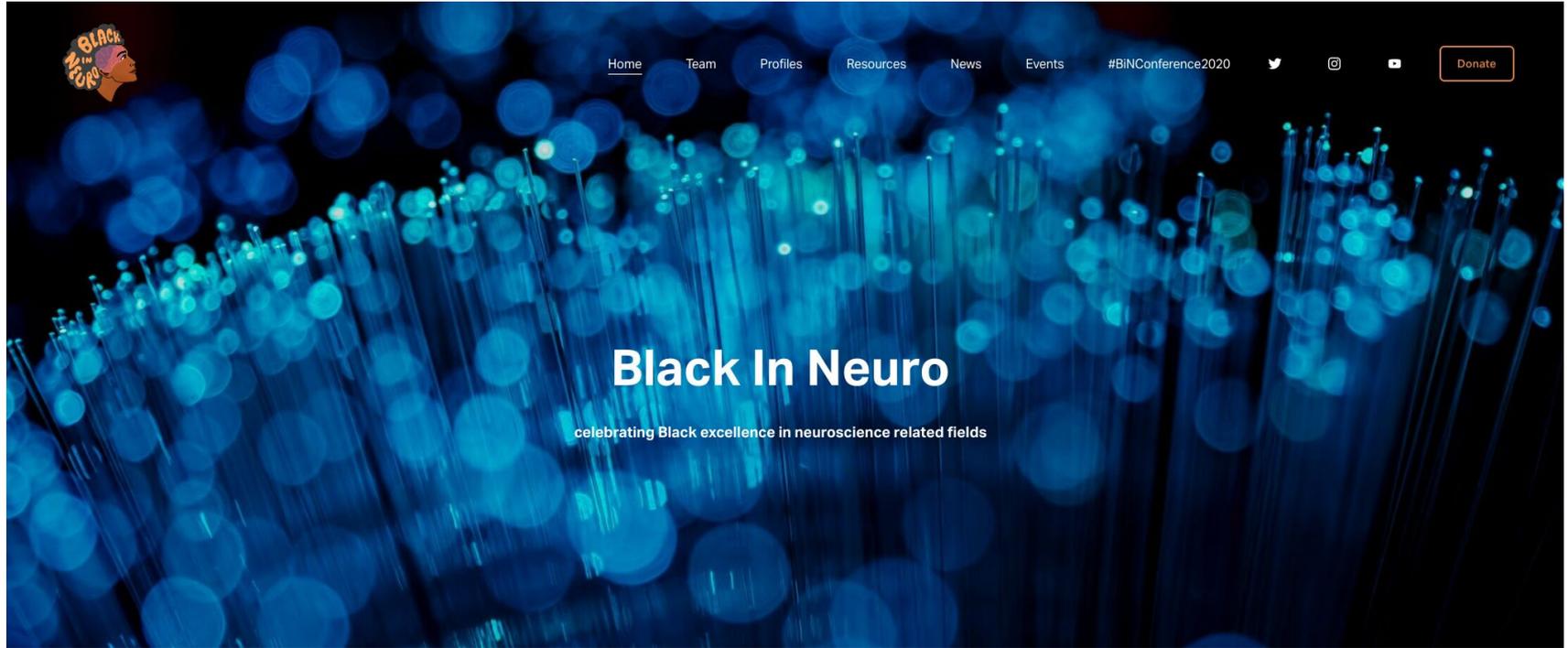
- sign up
- spread the word
- submit recommendations



 WINRePo1
 www.facebook.com/WiNRepository/

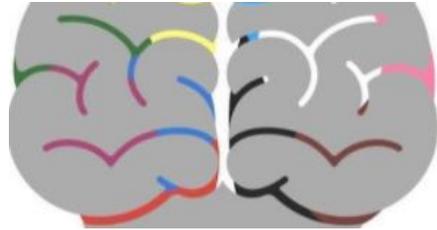


amazing things - super important



<https://www.blackinneuro.com/>

FAIR initiatives & support



Queer in Neuro

[Google form for signing up](#)



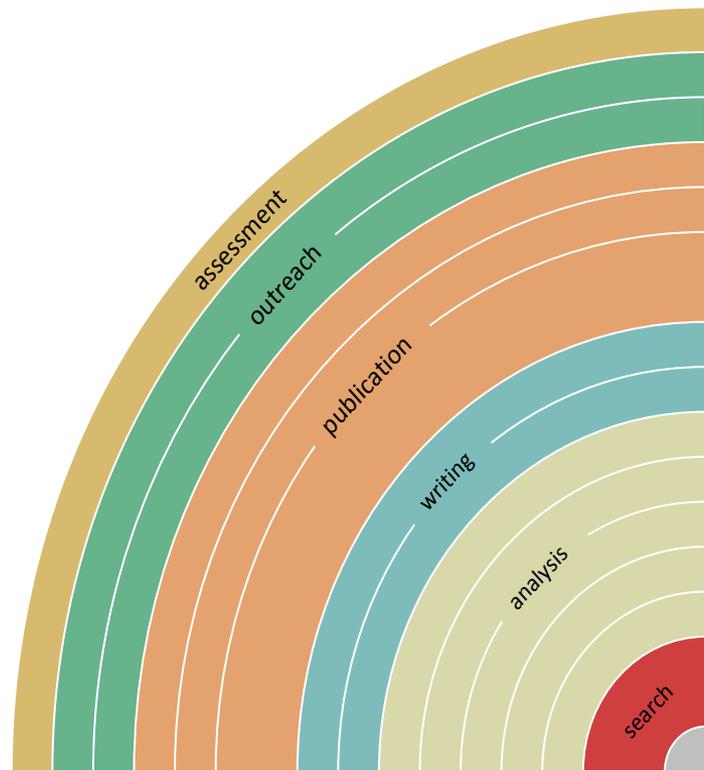
OHBM Student
& Postdoc SIG

[https://www.ohbmtrainees.com/
mentoring-programme/](https://www.ohbmtrainees.com/mentoring-programme/)



<https://ossig.netlify.app/>

FAIR data through community-driven development of standards and beyond



- necessities - support
- prerequisites - initiatives
- necessities - repositories (standardized)
- necessities - more than a PDF
- necessities - reporting (standardized)
- necessities - documentation
- necessities - virtualization (standardized)
- necessities - workflows (standardized)
- necessities - validation (standardized)
- necessities - standardization
- prerequisites - data sharing (standardized)
- prerequisites - preregistration (standardized)



Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)



FAIR data through community-driven development of standards and beyond

- **community-driven FAIR standards** enable new kinds of science:
 - open and collaborative development that is fast-paced, yet highly structured, understandable and traceable
- **community-driven FAIR standards** introduce changes on a global scale:
 - highly adapted across all levels of individuals and projects
- **community-driven FAIR standards** empower individuals:
 - support and engagement for as many folks with as many backgrounds as possible



FAIR data through community-driven development of standards and beyond

Standardization driven by an **open and structured community** helps **continuous integration, development, testing, adaption, extension, control, broadens science, includes everyone, teaching/training sustainability, getting rid of toxic structures and science islands** and if done early in workflow can **drastically benefit entire research workflow** as every other step can build upon standardization and can itself be standardized.



FAIR

(literally and figuratively)



Big kudos and thank you go out to ...

CONP
PCNO

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HEALTHY LIVES

ReproNim

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NIH
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of Mental Health
R01MH096906

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www.wmde.org/opensciencefellows

funding

Brain Hack

BIDS
BRAIN IMAGING DATA STRUCTURE

brainhack

community

Thank you very much for your attention.

I'm happy to answer questions during the chat and/or via twitter
(@peerherholz) or email (herholz dot peer at gmail dot com).



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