



McGill  
UNIVERSITY



neuro

Brms ·)))  
International Laboratory for  
Brain, Music, and Sound Research



brainhack.org DataLad

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

CSPD 2020

Peer Herholz (he/him)  
Postdoctoral researcher - NeuroDataScience-ORIGAMI lab at MNI, McGill  
- BRAMS

Member - BIDS, ReproNim, Brainhack, UNIQUE



BIDS  
BRAIN IMAGING DATA STRUCTURE



docker



Singularity



BIDS Apps



@peerherholz



Neuroscout



OpenNEURO



git

December 07, 2020



## A few things before we start ...

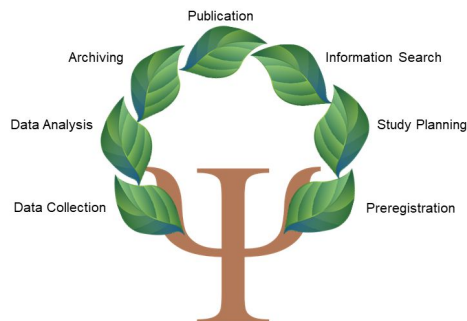
- to the folks with hearing impairments:
  - There will be live captions at the bottom of the screen. If something doesn't work, please let me/us know in the chat.
- to the folks with vision impairments:
  - All graphics (except those taken from publications) should be high in contrast and color-blind 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 ...

# Sharing Psychological Research Data: Best Practices and New Developments

07.12.2020 – 08.12.2020



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

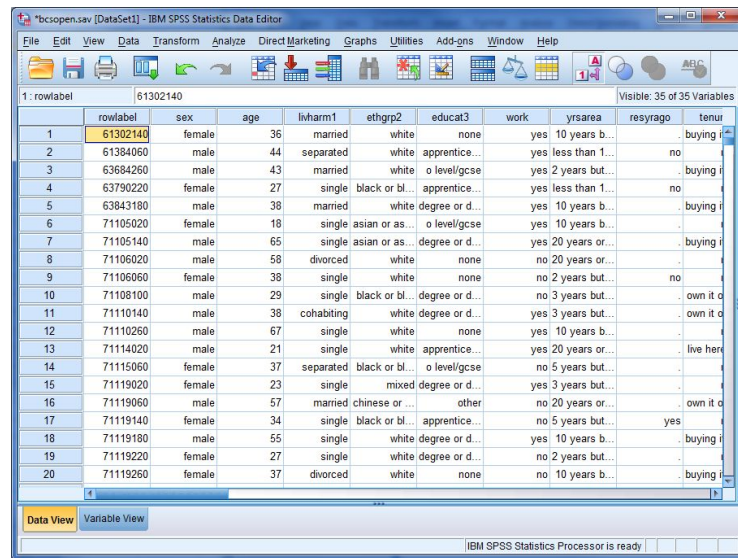
The data that support the findings of this study are available from the corresponding author upon reasonable request.

### Data Availability Statement

[Go to: ☐](#)

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

Code used to generate the analyses are available from the corresponding



The screenshot shows the IBM SPSS Statistics Data Editor window. 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 35 variables. The first variable is 'rowlabel', and the second is '61302140'. The data grid shows various demographic and behavioral variables such as 'sex', 'age', 'livharm1', 'ethgrp2', 'educat3', 'work', 'ysarea', 'resyrago', and 'tenur'. The status bar at the bottom indicates "IBM SPSS Statistics Processor is ready".

<https://bitb247.files.wordpress.com/2014/04/data-editor-labels.png>

“upon (reasonable) request” =  
you might get it if I (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

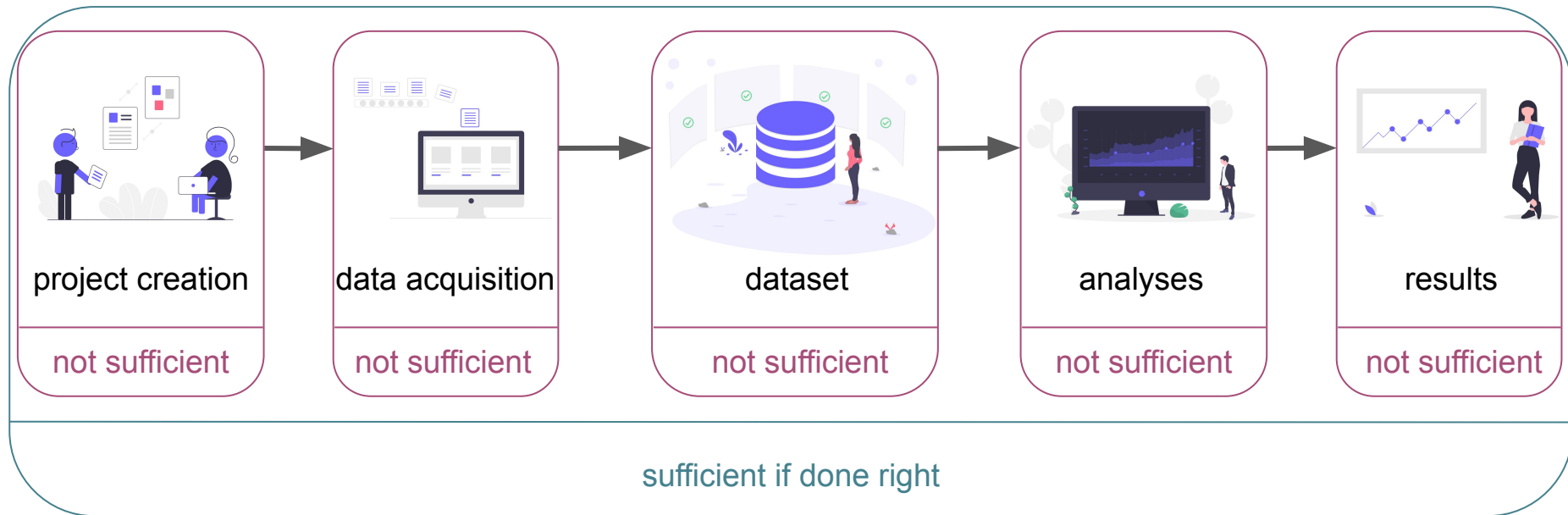


data/results in  
fancy journal





# 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://ogsci.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

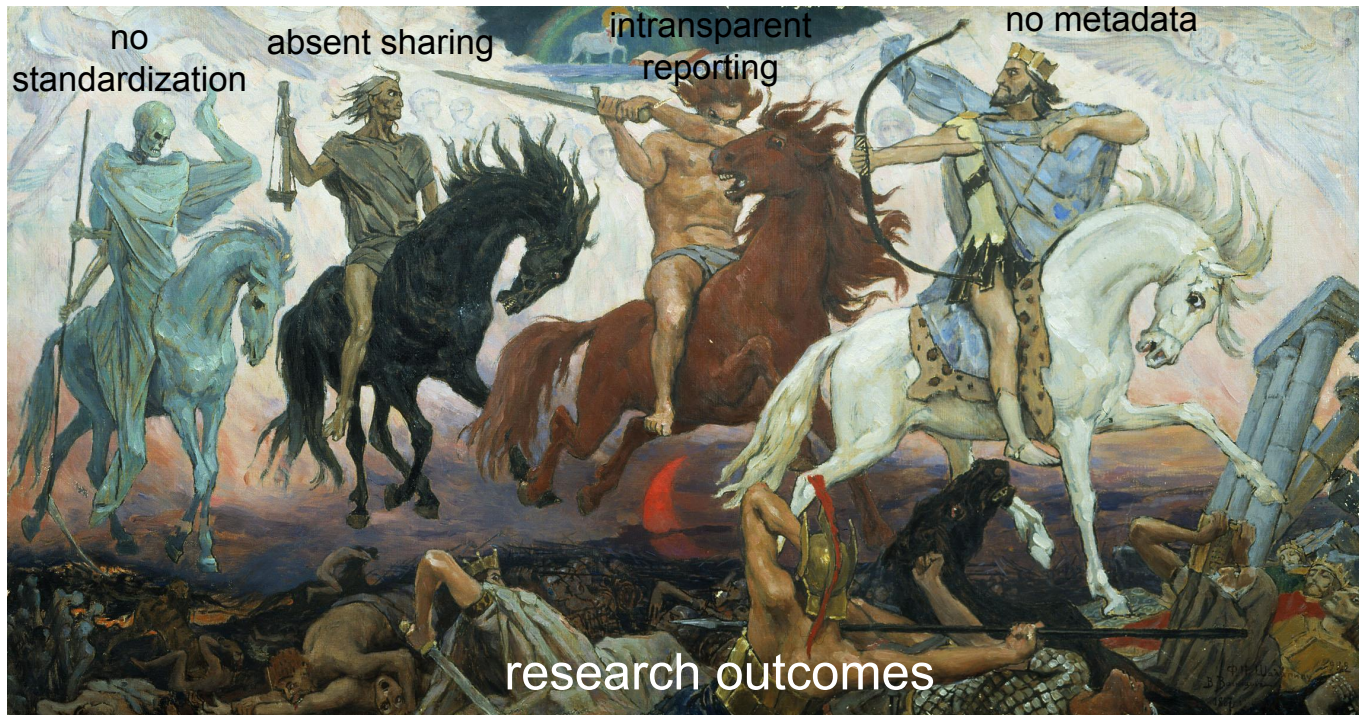
- FAIR is *fuzzy per design*
- a *general set of principles* that should *guide* research *across domains*
- FAIR has *natural variation across domains* (e.g. metadata in Psychology or Neuroscience)
- each domain *needs* to develop its own FAIR *standards and terms*, as no solution can fit all



- <https://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>



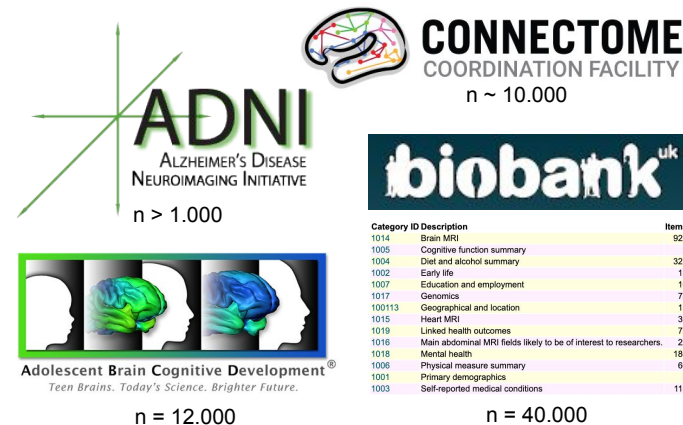
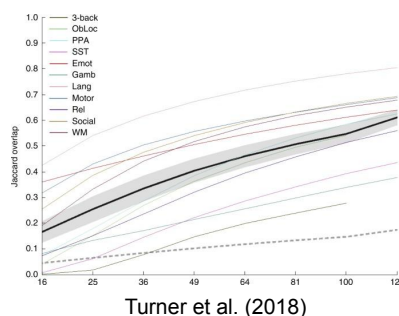
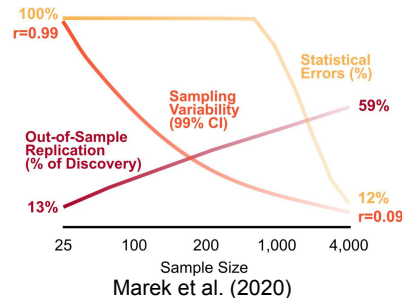
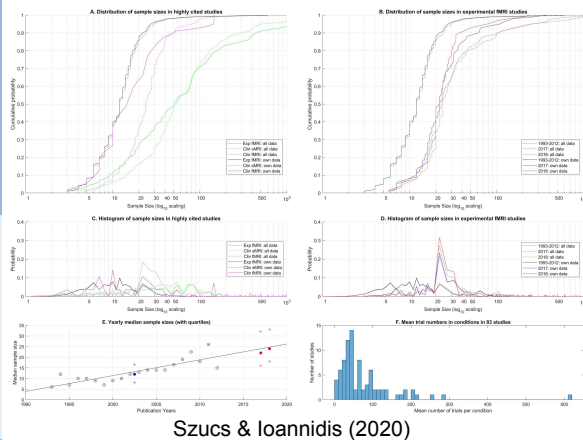
# The four horsemen of ... unFAIR research outcomes



Source: Public domain



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





# FAIR data through standardization ... with Marie Kondo



IBM SPSS Statistics Data Editor

File Edit View Data Transform Analyze Direct Marketing Graphs Utilities Add-ons Window Help

1: rowlabel 61302140 Visible: 35 of 35 Variables

	rowlabel	sex	age	ltham1	ethgr2	educat3	work	years	resygr	tenur
1	61302140	female	36	married	white	none	yes	10 years b.	buying	
2	61384060	male	44	separated	white	apprentice	yes	less than 1.	no	
3	63040400	male	43	married	white	o level/grade	yes	2 years but.	buying	
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	
6	71105020	female	18	single	asian or as.	o level/grade	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	71106060	female	38	single	white	none	no	3 years but.	own it c.	
10	71108100	male	29	single	black or bl.	degree or d.	no	3 years but.	own it c.	
11	71110140	male	38	cohabiting	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	apprentice	yes	20 years or.	line her	
14	71115060	female	37	separated	black or bl.	o level/grade	no	5 years but.	own it c.	
15	71119020	female	23	single	mixed	degree or d.	yes	3 years but.	own it c.	
16	71119060	male	57	married	chinese or ...	other	no	20 years or.	own it c.	
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	
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	

Data View Variable View

IBM SPSS Statistics Processor is ready

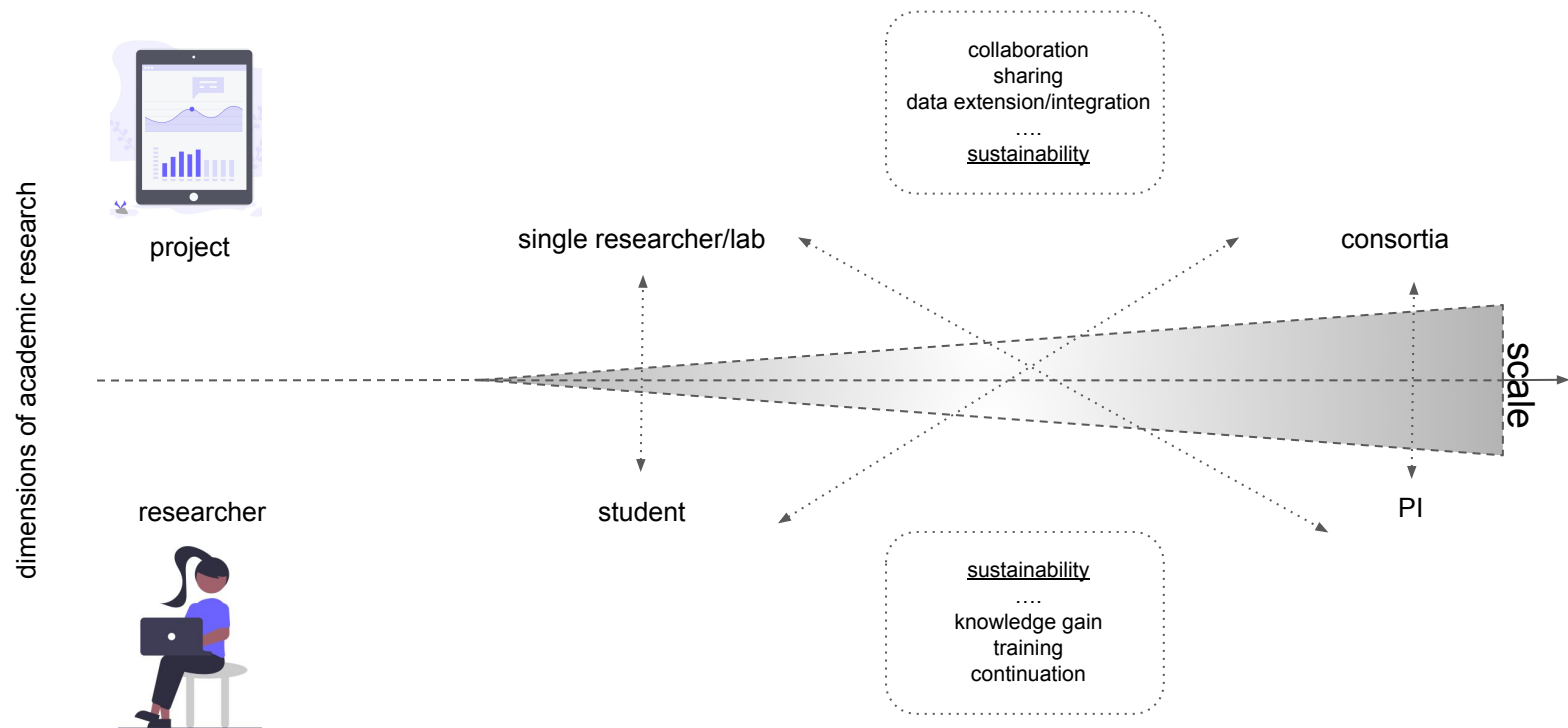
<https://bizlib247.files.wordpress.com/2014/04/data-editor-labels.png>

```
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_sref.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_magnitudel.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!

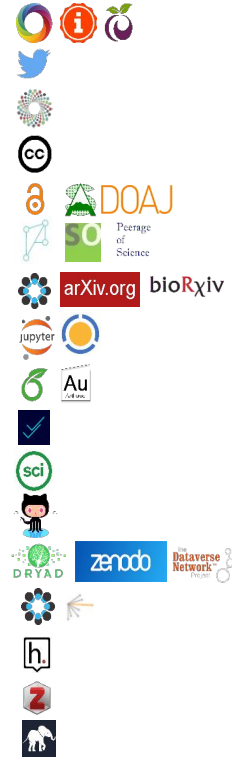




# FAIR by default, from the beginning to the end and beyond



- 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



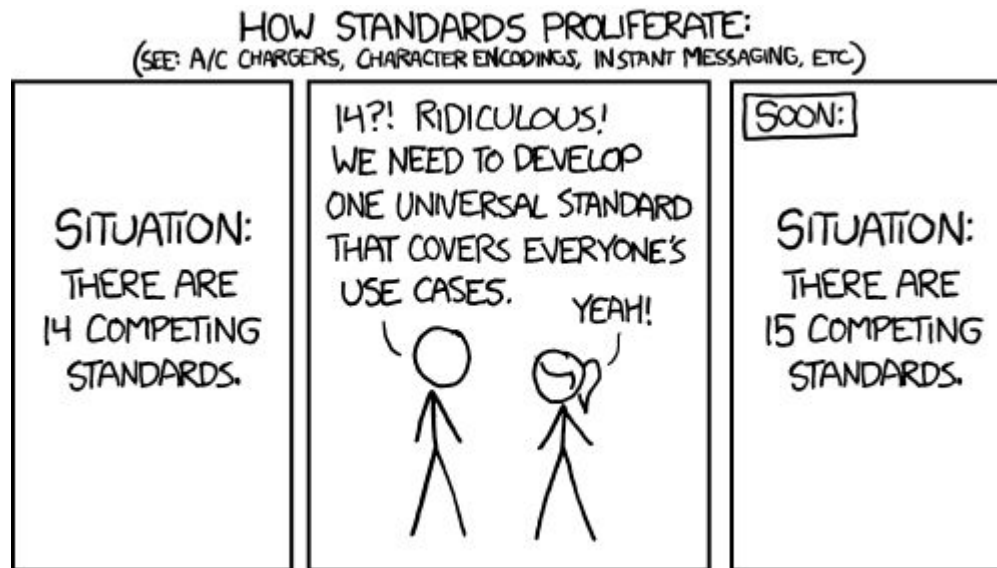
Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

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





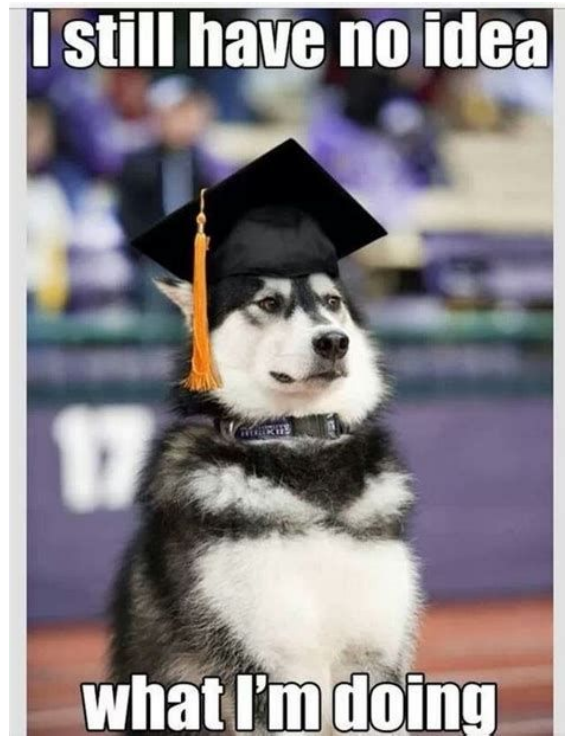
# FAIR data through standardization - problem I



<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/1H5ABVS>

\* 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



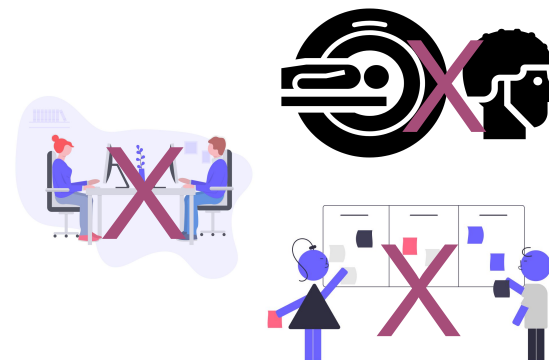
# FAIR data through standardization - problem III

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 Beitner<sup>1</sup>, Garvin Brod<sup>2,3</sup>, Benjamin Gagl<sup>1</sup>, Dominik Kraft<sup>4</sup>, und Martin Schultze<sup>5</sup>

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

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

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

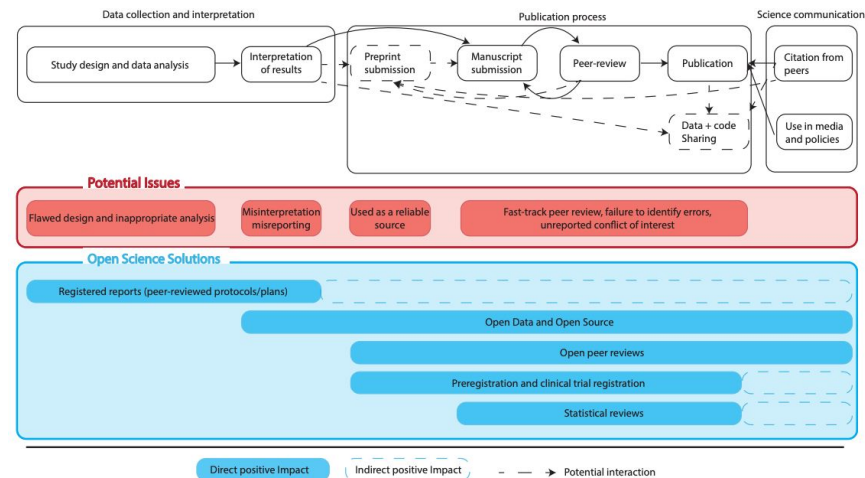
<sup>4</sup> Neurokognitive Psychologie, Goethe-Universität Frankfurt am Main. Theodor-W.-Adorno-Platz 6, D-60323 Frankfurt am Main.

<sup>5</sup> Psychologische 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. Autorenenreihenfolge alphabetisch.

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

Lonni Besançon<sup>1,2\*</sup>, Nathan Peiffer-Smadja<sup>3,4</sup>, Corentin Segalas<sup>5</sup>, Haiting Jiang<sup>6</sup>, Paola Masuzzo<sup>7</sup>, Cooper Smout<sup>7</sup>, Eric Billy<sup>8</sup>, Maxime Deforet<sup>9+</sup> and Clémence Leyrat<sup>5,10+</sup>



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

<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?



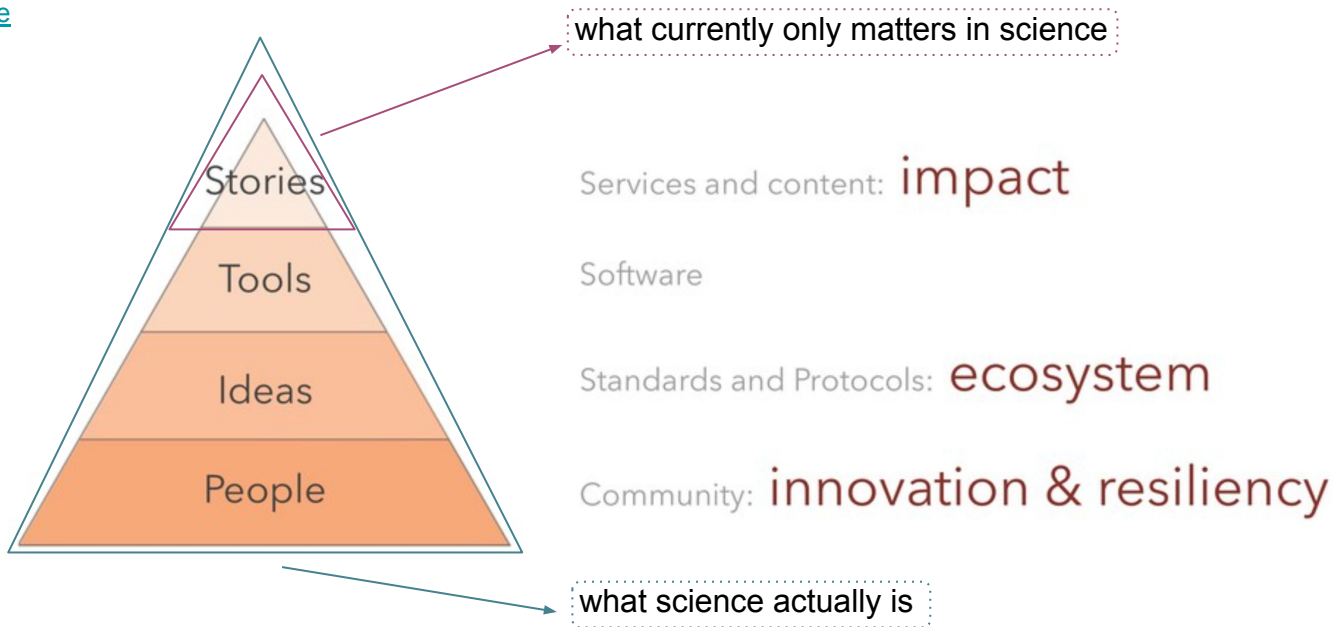
## “The future is now.”

- \* 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.”*

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



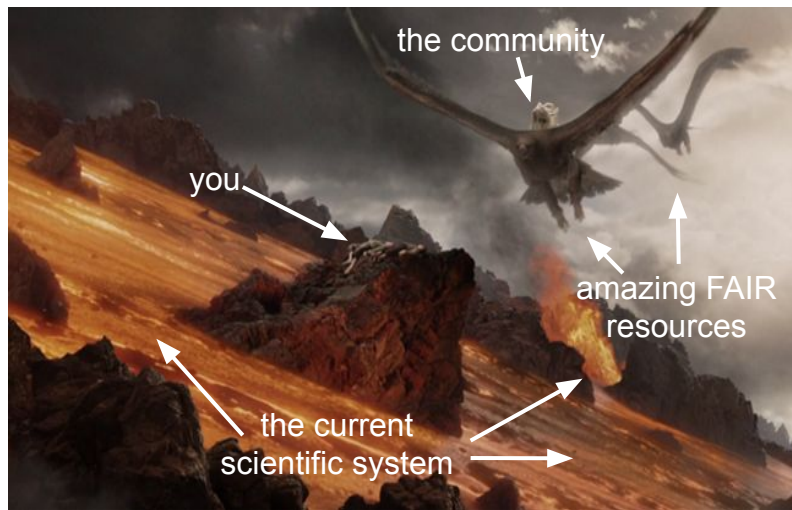
## FAIR data through standardization - problem IV

	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

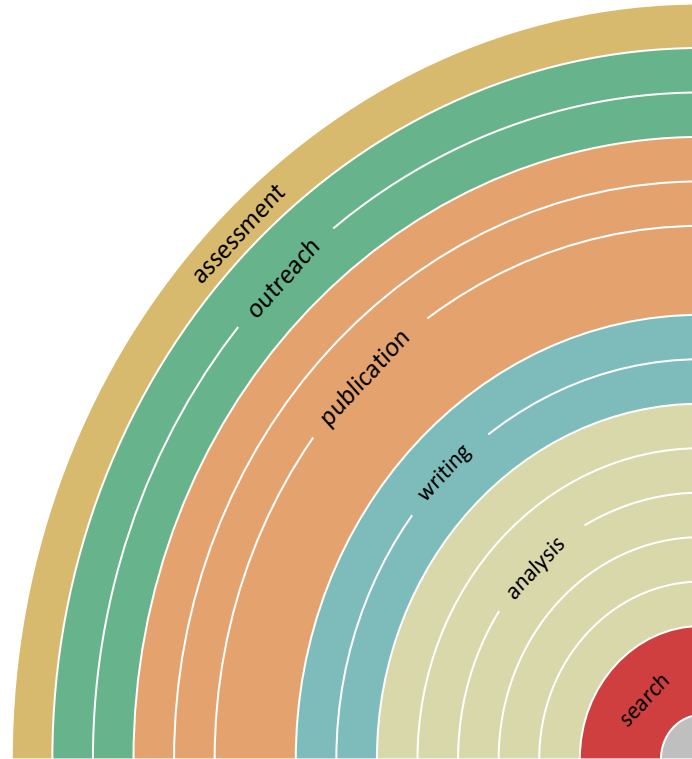
\*adapted from [Elizabeth DuPre](#)

- **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

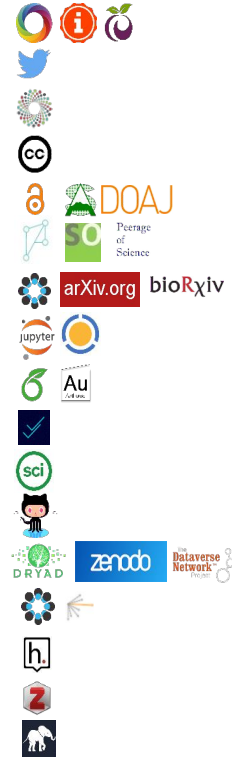




# FAIR data through community-driven development of standards and beyond



- 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



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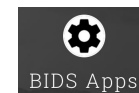
DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)



# 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)



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# FAIR data through community-driven development of standards and beyond



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



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# 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)

email address you have used in AsPredicted  
I cannot access my AsPredicted email account anymore

[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

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## 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](mailto: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/testing 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.

Bayer 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.1411525](https://doi.org/10.5281/zenodo.1411525)

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 information 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.

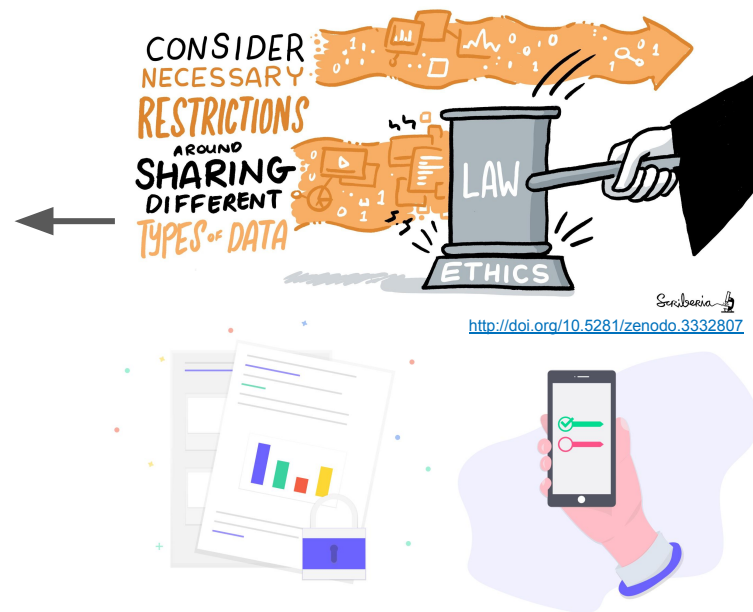
To the best of our knowledge, the data we release to the general public will not contain information that can directly identify you. The data will not have your name on it, only a code number, so people will not know your name or which data are yours. In addition, the data will not include data that we think might help people who know you guess which data are yours, such as your facial features or the data 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





## 2018



↓

2019

A group of nine people, seven men and two women, are standing in a row in an office setting. They are dressed in business casual attire. Behind them is a grey wall featuring the 'Cosmo' logo, which consists of a stylized 'C' made of three interlocking shapes followed by the word 'cosmo' in lowercase. The floor is made of light-colored wood. To the right, a portion of a desk and a window with a view of a city skyline are visible.



# 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)

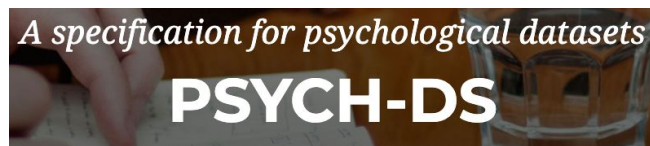


# FAIR necessities - standardization - fantastic standards in the wild



120+ contributors\*

master	7 branches	3 tags	Go to file	Add file	3	Code	About
Brain Imaging Data Structure (BIDS) Specification							
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.							



20+ contributors\*

## 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.

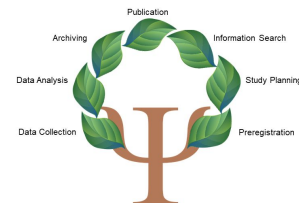
Welcome! If this is your first time checking out this project (and especially if this is your first time working with a technical specification document), please also check out the project landing page for an explanation of the goals of this project and notes on how you can contribute.

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 PsychCurat - Development of user-oriented curation criteria for psychological research data

### Project description

In the context of the omni-present 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 PsychCurat: Development of user-oriented curation criteria for psychological research data, which is funded by the Federal Ministry of Education and Research (BMBF), 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, PsychCurat 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?

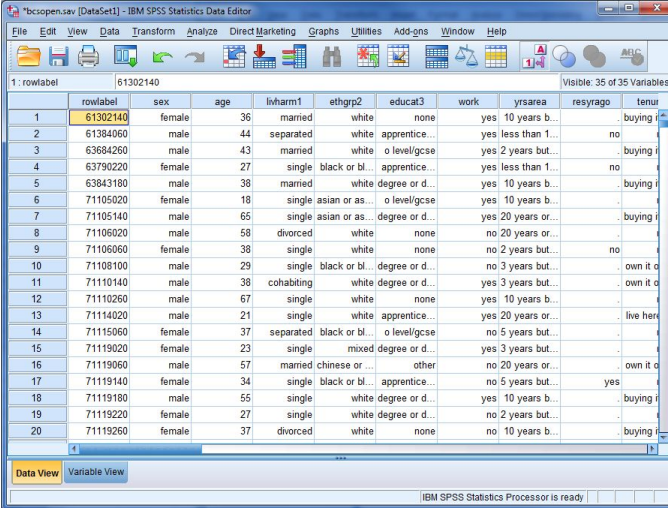
- storage, structure, metadata, version control?

## 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



The screenshot shows the IBM SPSS Statistics Data Editor window. The menu bar includes File, Edit, View, Data, Transform, Analyze, Direct Marketing, Graphs, Utilities, Add-ons, Window, and Help. The toolbar contains icons for file operations, data manipulation, and analysis. The main window displays a data grid with 20 rows and 12 visible columns. The first column is labeled 'rowlabel' and contains row numbers 1 through 20. The second column is labeled '61302140' and contains a list of IDs. The third column is labeled 'sex' and contains 'female' or 'male'. The fourth column is labeled 'age' and contains ages from 36 to 27. The fifth column is labeled 'lvharm1' and contains marital status terms like 'married', 'separated', 'single', 'divorced', 'cohabiting'. The sixth column is labeled 'ethgrp2' and contains racial/ethnic categories like 'white', 'black or bl...', 'asian or as...', 'mixed', 'chinese or ...', 'black or bl...'. The seventh column is labeled 'educat3' and contains education levels like 'none', 'apprentice...', 'o level/gcse', 'degree or d...', 'other'. The eighth column is labeled 'work' and contains employment status like 'yes', 'no'. The ninth column is labeled 'yrsare' and contains years of experience like '10 years b...', 'less than 1...', '2 years but...', 'less than 1...', '10 years b...', '20 years or...', '2 years but...', '3 years but...', '10 years b...', '5 years but...', '3 years but...', '20 years or...', '5 years but...', '10 years b...', '2 years but...', '10 years b...'. The tenth column is labeled 'resyago' and contains 'no' or 'yes'. The eleventh column is labeled 'tenur' and contains 'buying r...', 'no', 'buying r...', 'no', 'buying r...', 'no', 'own it o...', 'own it o...', 'live her...', 'own it o...', 'own it o...', 'yes', 'buying r...', 'buying r...', 'buying r...'. The status bar at the bottom indicates 'IBM SPSS Statistics Processor is ready'.

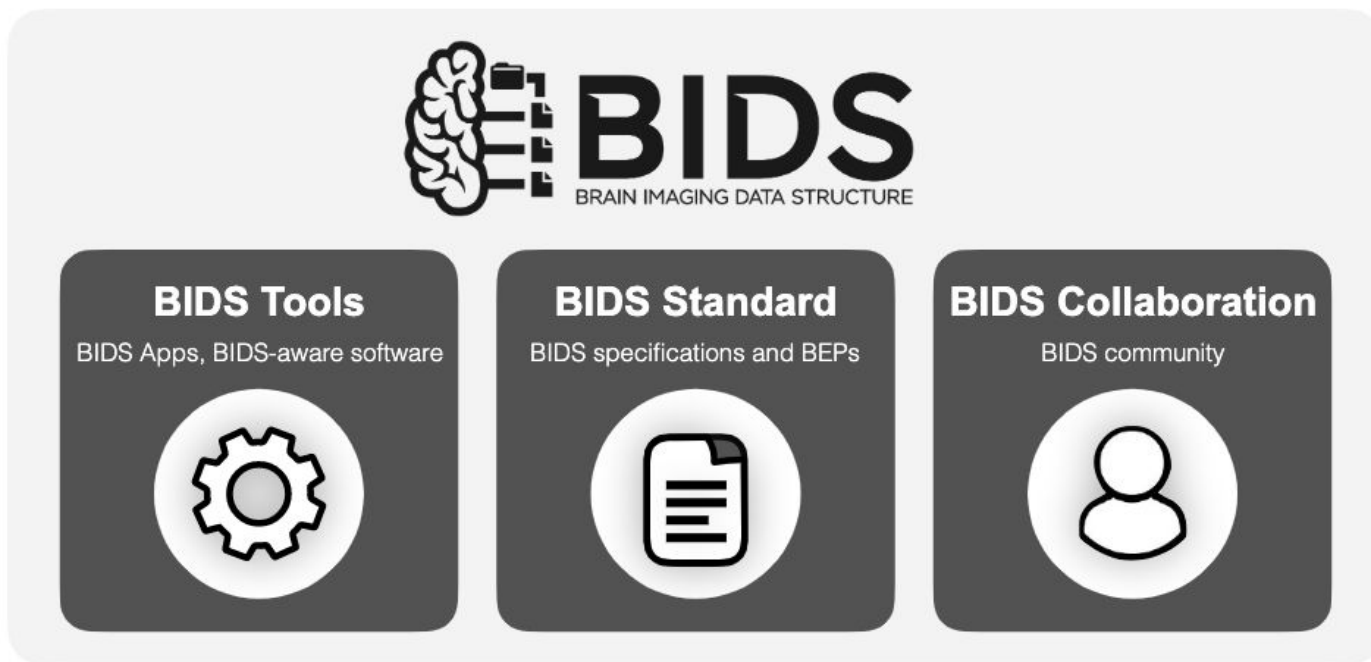
rowlabel	61302140	sex	age	lvharm1	ethgrp2	educat3	work	yrsare	resyago	tenur
1	61302140	female	36	married	white	none	yes	10 years b...		buying r...
2	61384060	male	44	separated	white	apprentice...	yes	less than 1...	no	
3	63684260	male	43	married	white	o level/gcse	yes	2 years but...		buying r...
4	63790220	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 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 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/60621009"  
  }  
}
```



# 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/dfj.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

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<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**



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<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,  
  ...  
}
```



## 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

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<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/60621009"  
  }  
}
```



# 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. Draschkow
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. Duft, 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



# FAIR necessities - standardization - the idea of BIDS

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

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



Brain Imaging Data Structure v1.4.1

Brain Imaging Data Structure  
v1.4.1

The BIDS Specification

The BIDS Starter Kit

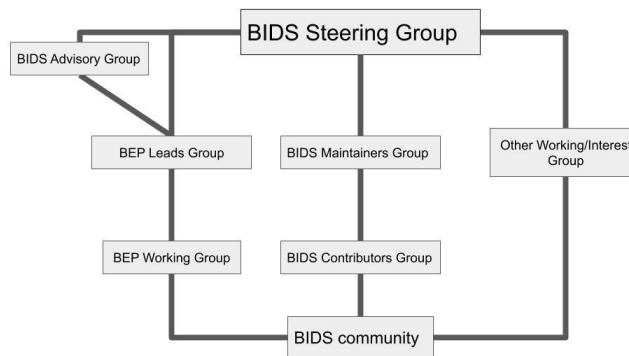
Filters	Labels (2)	Milestones (2)	New Issue
130 Open	✓ 186 Closed		
Fix example for IEEGElectrodeGroups field from object to string type	ESD		5
Migrate "Travis CI" service to "GitHub Actions" CI service	good first issue	enhancement	
Extend BIDS format to magnetic resonance spectroscopy			3
Change requirement level "optional" -> "recommended" for three fields in dataset_description	options wanted		2
ESD - confusion between channels.tsv and electrodes.tsv			2
[ENH] add realignment parameters in JSON for "McCo" sequences			
Add OPTIONAL SkullStripped metadata to any MR images	options wanted		2
Add consistent Units information in the json field tables			3
Bare BIDS - can requirements for README be enhanced?			13
"T1w" as keyword for <datatype>"CoordinateSystem"	MEG	MEG	ESD
Alternating row fills in tables within the PDF build	formatting		
Define nested coordinate systems in the BIDS specification instead of FieldTrip wiki	MEG	MEG	ESD
Add support for multi-echo T1w scans			11
EEG channel type inconsistency	consistency		2
Harmonize .tsv examples (and their rendering)	enhancement		
Unspecified status of extra content in the JSON files			3
RFC: Maintainer/Steering credit on BEP papers	community		6
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



Guiomar Niso  
Spain  
MEG



Melanie Ganz  
Denmark  
PET



Robert Oostenveld  
The Netherlands  
M/EEG



Russ Poldrack  
USA  
fMRI



Kirstie Whitaker  
UK  
MRI

## BIDS Maintainers



Stefan Appelhoff  
Germany



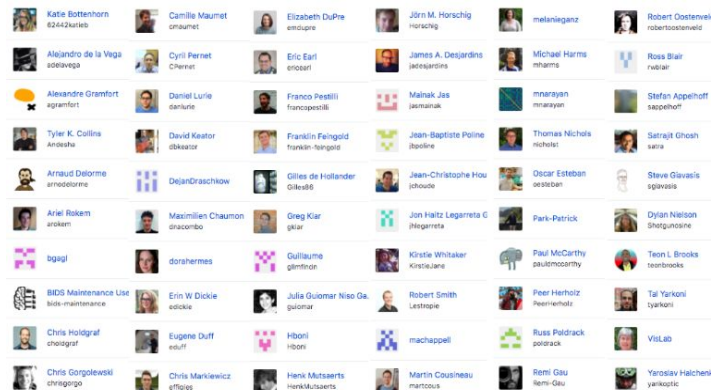
Chris Markiewicz  
USA



Taylor Salo  
USA



Franklin Feingold  
USA



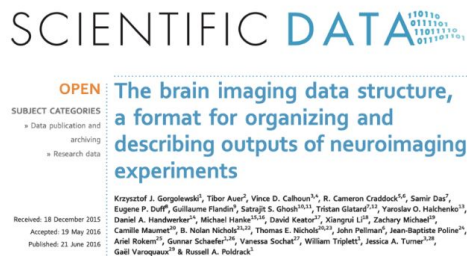
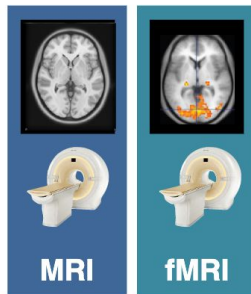
\*only some contributors are shown



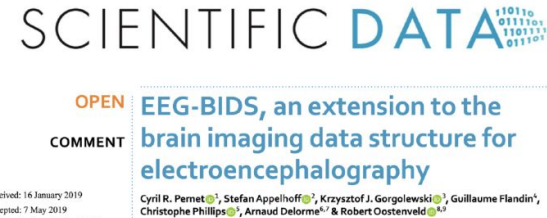
# FAIR necessities - standardization - the idea of BIDS <https://bids-specification.readthedocs.io/en/stable/>

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

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

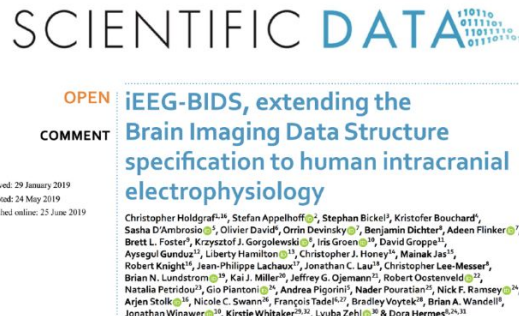
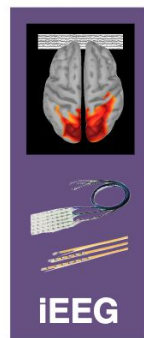


(Gorgolewski et al. 2016)



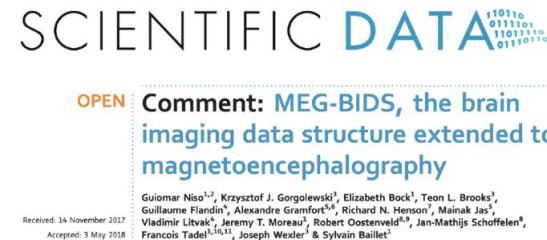
Received: 16 January 2019  
Accepted: 7 May 2019  
Published online: 25 June 2019

(Pernet et al., 2019)



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

(Holdgraf et al., 2019)



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

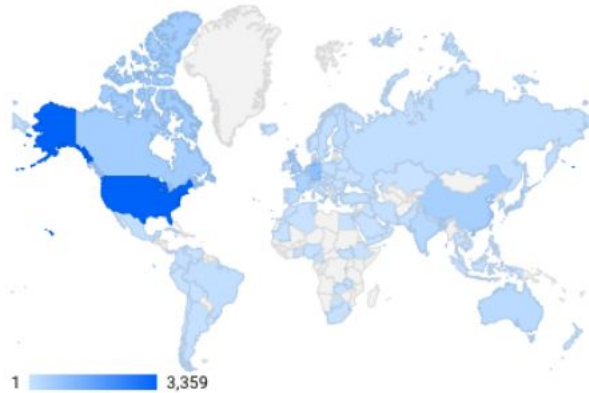
(Niso et al., 2018)



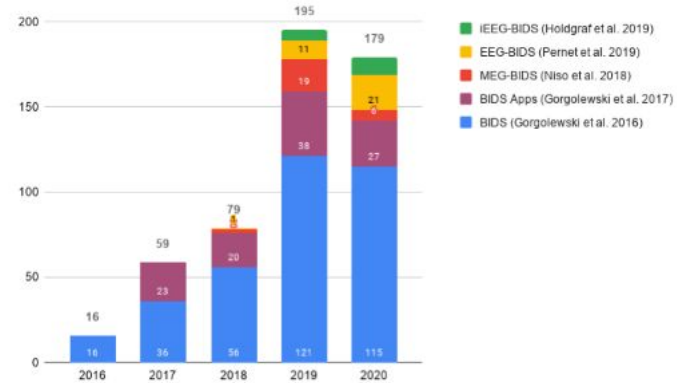
# FAIR necessities - standardization - the idea of BIDS <https://bids-specification.readthedocs.io/en/stable/>

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

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

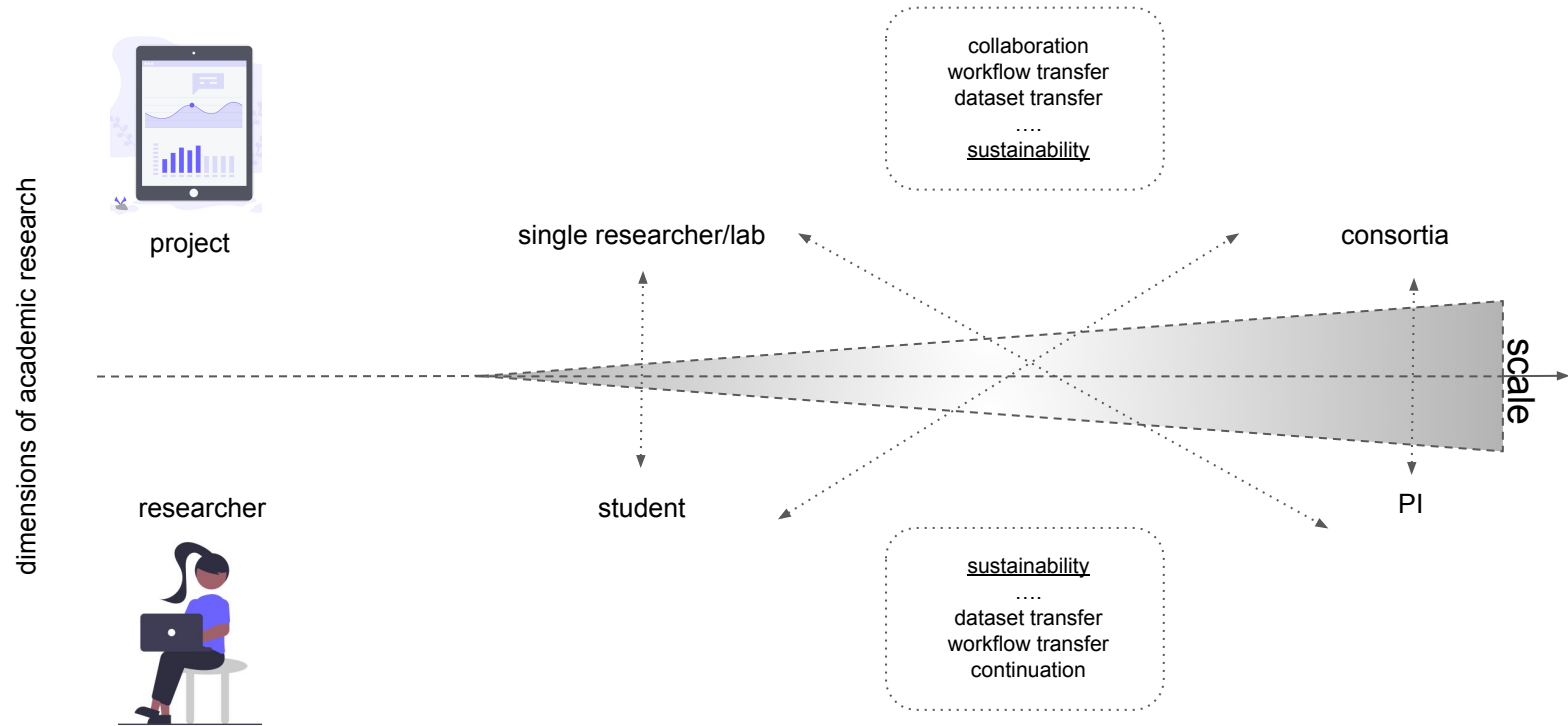


**BIDS new citations per year**



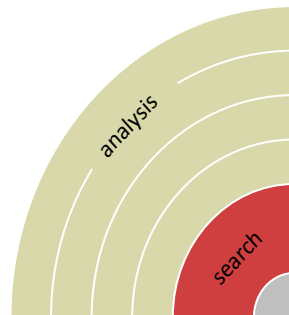


# FAIR necessities - standardization - the benefits of BIDS

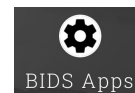




# FAIR data through community-driven development of standards and beyond



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

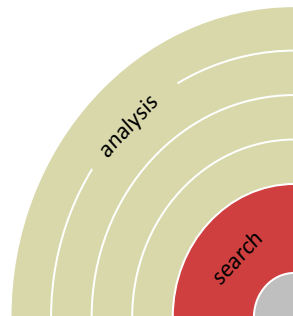


Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

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



# FAIR data through community-driven development of standards and beyond



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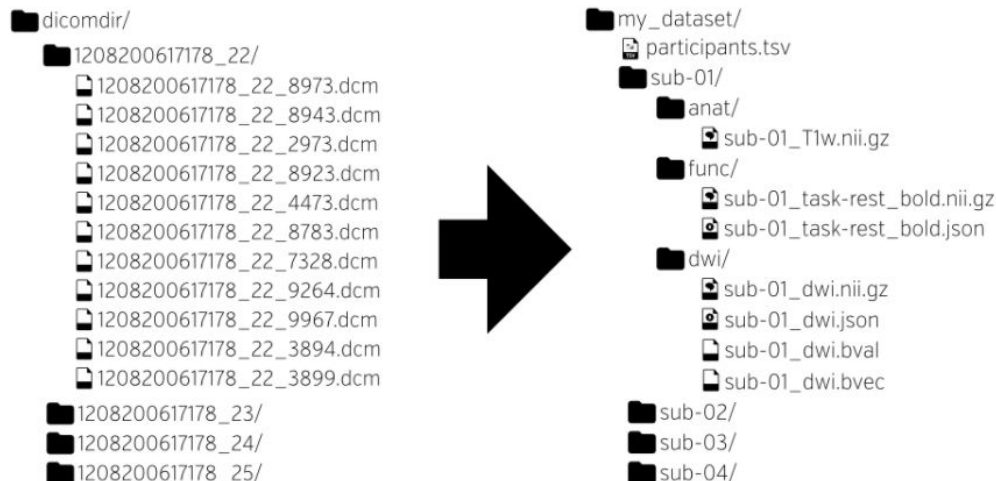
# FAIR necessities - standardization - the benefits of BIDS - data conversion

\* 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, BIDS23col, BIDS2NDA, bidsfy, bidskit, dac2bids, Data2Bids, Dcm2bids, heudiconv, DCM2NIfx, ReproIn, XNAT2BIDS, bids2xar, BIDS2NIDM, BIDScoin, MNE-BIDS...





# 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"><li>• 39 Files, 757.25MB</li><li>• 4 - Subjects</li><li>• 1 - Session</li></ul>	<ul style="list-style-type: none"><li>• memory</li></ul>	<ul style="list-style-type: none"><li>• bold</li><li>• T1w</li></ul>

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"><li>• 39 Files, 757.25MB</li><li>• 4 - Subjects</li><li>• 1 - Session</li></ul>	<ul style="list-style-type: none"><li>• memory</li></ul>	<ul style="list-style-type: none"><li>• bold</li><li>• T1w</li></ul>

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"><li>• 39 Files, 757.25MB</li><li>• 4 - Subjects</li><li>• 1 - Session</li></ul>	<ul style="list-style-type: none"><li>• memory</li></ul>	<ul style="list-style-type: none"><li>• bold</li><li>• T1w</li></ul>

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/anat/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.

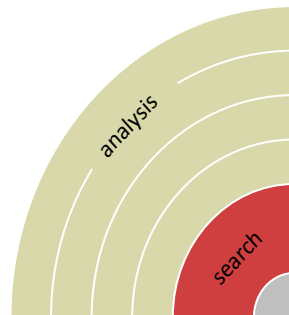
## Warning: 2

2 files

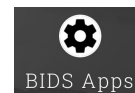
Not all subjects/sessions/runs have the same scanning parameters.



# FAIR data through community-driven development of standards and beyond



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

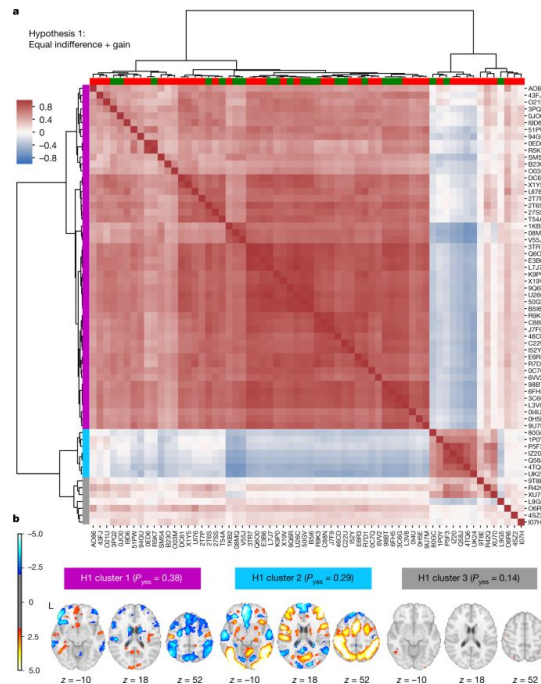
DOI: [10.5281/zenodo.1147025](https://doi.org/10.5281/zenodo.1147025)



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

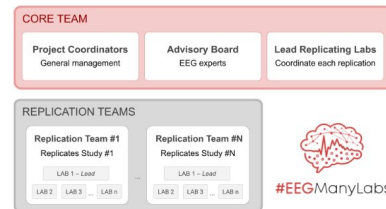
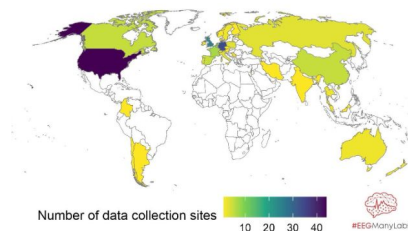
- a lot of sources of analyses/results variability
- every aspect of scientific workflow matters and therefore should be **FAIR**
- acquisition, datasets and metadata
- computing environment (OS, software package, version number, etc.)
- mega & multiverse analyses demonstrated the need for **FAIR** and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

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



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

Pavlov et al. (2020)  
#EEGManyLabs

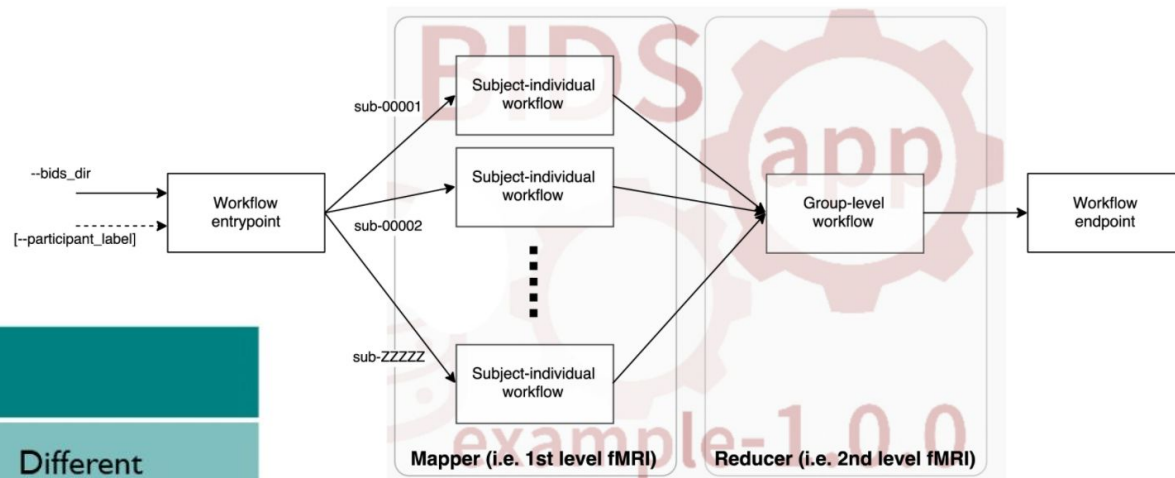


<https://psyarxiv.com/528nr/>



# 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/>



McGill  
UNIVERSITY



neuro

Brams  
International Laboratory for  
Brain, Music, and Sound Research



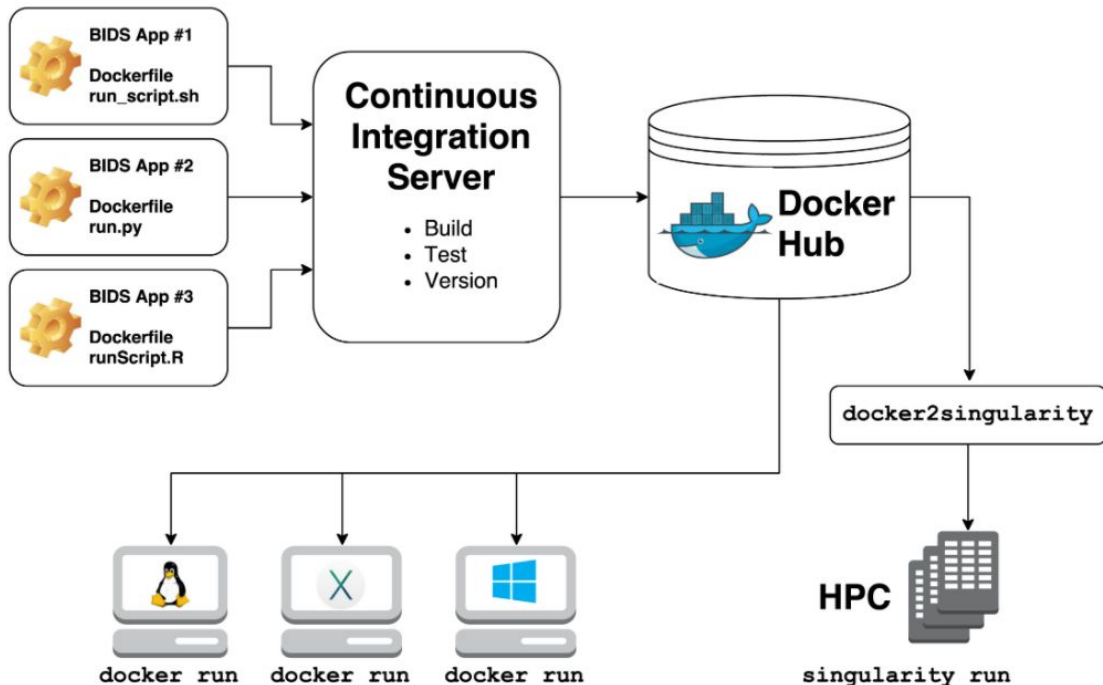


# FAIR necessities - standardization - the benefits of BIDS - data processing

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

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

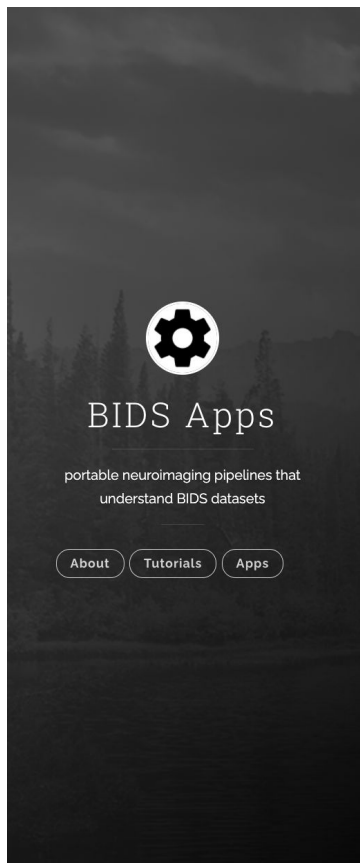
- containerized pipelines/workflow that understand BIDS datasets
- set of core command line arguments
- should not depend on software outside the container
- openly hosted on docker/singularity hub and include version tags
- targeting automated and standardized analyses that are reproducible



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



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



## Available BIDS Apps

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/HCPipelines  
BIDS-Apps/MAGeTbrain  
BIDS-Apps/tracula  
BIDS-Apps/baracus  
BIDS-Apps/antsCorticalThickness

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

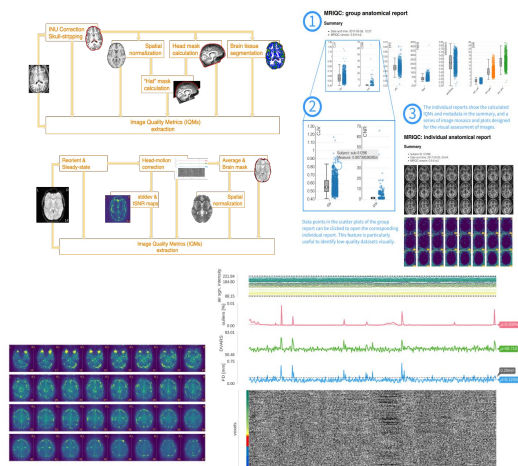
version v0.0.7	open issues 5	circled passing	open bug pull requests 0	docker pulls 14k	439.5MB	23 layers
version v0.9.1-5.1	open issues 11	circled passing	open bug pull requests 0	docker pulls 1.1k	2.6GB	52 layers
version v0.1.0	open issues 11	circled passing	open bug pull requests 0	docker pulls 8k	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 11	circled passing	open bug pull requests 0	docker pulls 250	576.8MB	31 layers
version v0.9.20	open issues 3	circled passing	open bug pull requests 0	docker pulls 1.5k	2GB	24 layers
version v0.15.1	open issues 125	circled passing	open bug pull requests 0	docker pulls 69k	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 3k	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 27k	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 376k	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 v2.7.0-1	open issues 0	circled passing	open bug pull requests 0	docker pulls 207	391.9MB	21 layers



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

## quality control (MRIQC)

mriqc data/ output/ --specifications

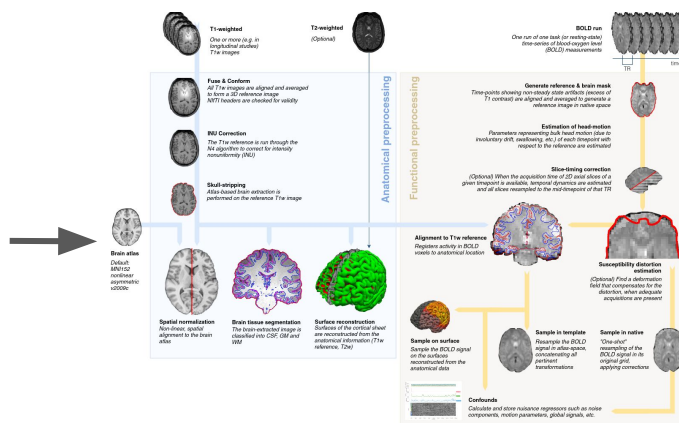


[Esteban et al. \(2017\),](https://mriqc.readthedocs.io/en/stable/index.html)

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

## preprocessing (fMRIPREP)

fmriprep data/ output/ --specifications

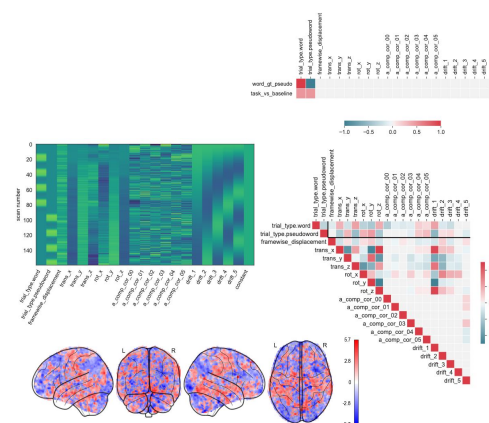


[Esteban et al. \(2019\),](https://fmriprep.org/en/stable/)

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

## statistical analyses (fitlins)

fitlins data/ output/ --specifications

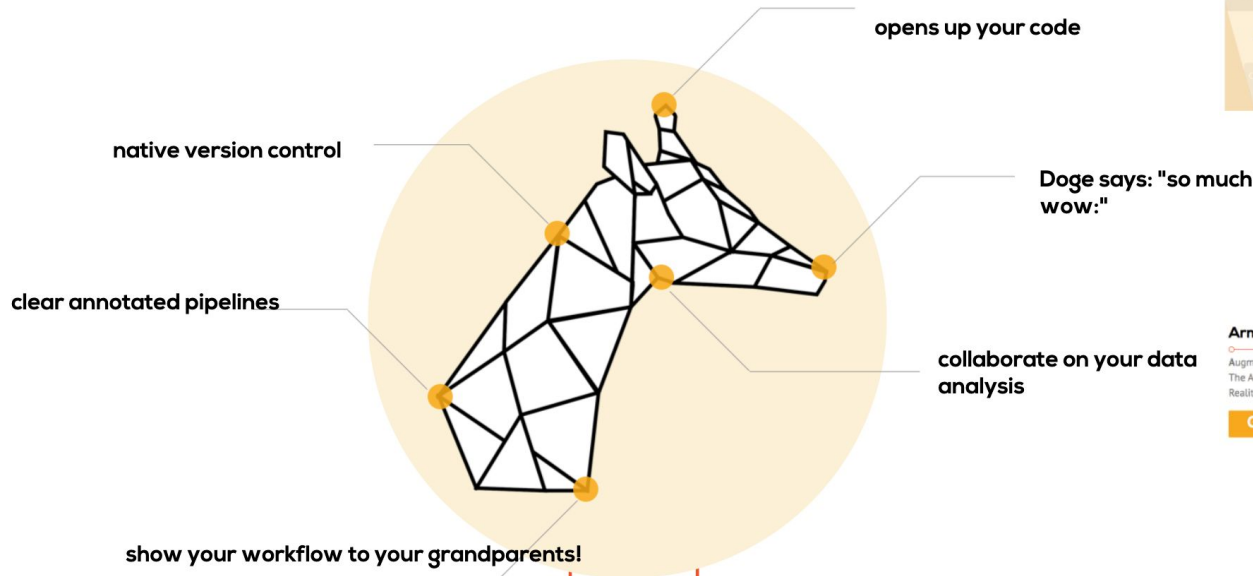


<https://fitlins.readthedocs.io/en/latest/index.html>



# FAIR necessities - standardization - the idea of data processing

<https://giraffe.tools/>



## Porcupine

**Porcupine Creates Ur 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!**





# FAIR necessities - standardization - combining data, processing, outcomes

<https://neuroscout.org/>



# Neuroscout

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

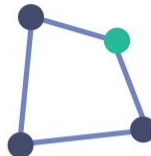
[Browse public analyses](#)

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

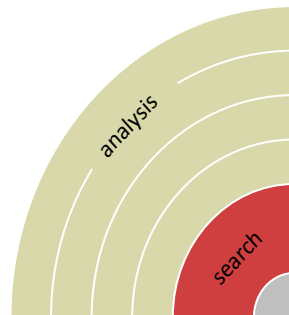


Portable BIDS pipelines enable execution with no configuration. Results are automatically uploaded to [NeuroVault](#) for easy sharing.

[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](#)

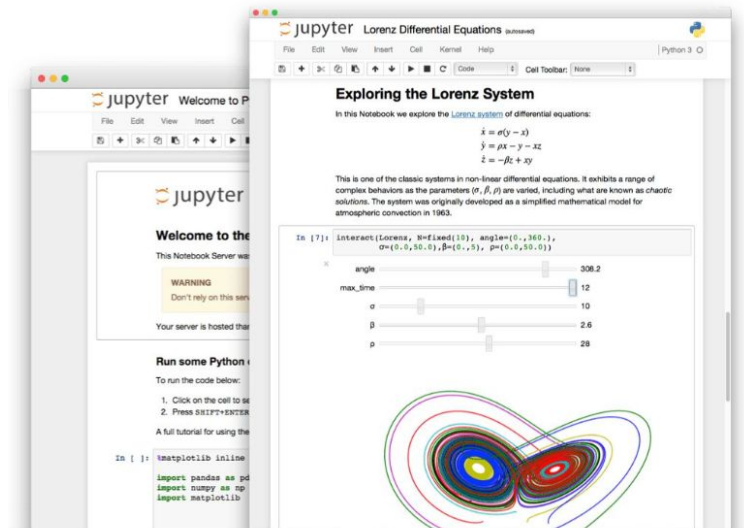


- a **community** of people
- an **ecosystem** of **open tools** and **standards** for **interactive computing**
- **language-agnostic** and **modular**
- **empower people** to use other **open tools**



# FAIR necessities - analyses documentation - Jupyter notebooks

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

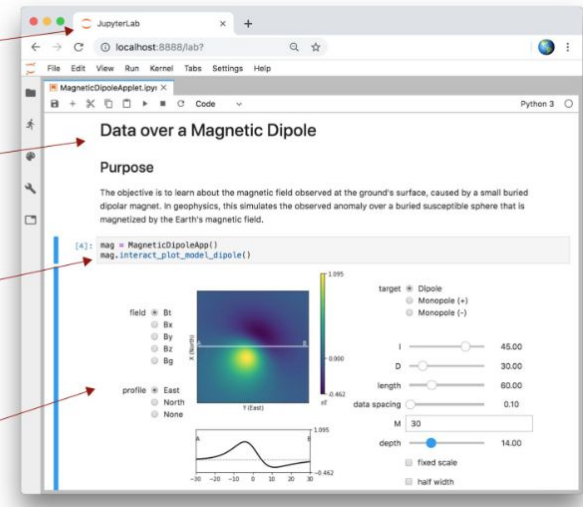


Runs in web browser

Text, comments, equations

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

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

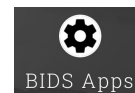




# FAIR data through community-driven development of standards and beyond



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



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







# FAIR necessities - reporting (standardized)



Link to this doc too  
<https://docs.google.com/document/d/19C6uJhV3JXenQNW5y4t8dMTEde888LKK7UxmUxdt>

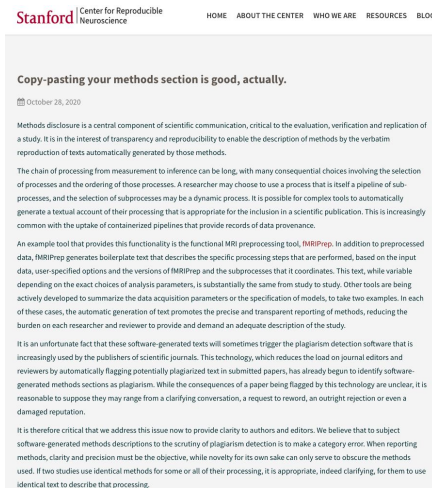
Methods disclosure is a central component of scientific communication, critical to the evaluation, verification and replication of a study. It is in the interest of transparency and reproducibility to enable the precise description of methods by the verbatim reproduction of texts generated by automatic methods.

The chain of processing from measurement to inference can be long, with many consequential choices involving the selection of processes and the ordering of those processes. A researcher may choose to use a process that is itself a pipeline of sub-processes, and the selection of sub-processes may be a dynamic process. It is possible for complex tools to automatically generate a textual account of their processing that is appropriate for the inclusion in a scientific publication and whose descriptions will tend to be more accurate than those generated by individual researchers. This is increasingly common with the uptake of containerised pipelines that provide records of data provenance.

An example tool that provides this functionality is the functional MRI preprocessing tool, fMRIPrep. In addition to preprocessed data, fMRIPrep generates boilerplate text that describes the specific processing steps that are performed, based on the input data, user-specified options and the versions of fMRIPrep and the subprocesses that it coordinates. This text, while variable depending on the exact choices of analysis parameters, is structured to summarise study-relevant information in as standardized of a format as possible. Other tools are being actively developed to summarise the data acquisition parameters or the specification of models, to take two examples. In each of these cases, the automatic generation of text promotes the precise and transparent reporting of methods, reducing the burden on each researcher and reviewer to provide and demand an adequate description of the study.

As more and more researchers use these software-generated texts, they will increasingly trigger the plagiarism detection software used by the publishers of scientific journals as well as graduate and postgraduate student exam administrators. This technology, which reduces the load on journal editors and reviewers by automatically flagging potentially plagiarized text in submitted papers, has already begun to identify software-generated methods sections as plagiarism. While the consequences of a paper being flagged by this technology are dependent on local university regulations and publishers' internal rules, it is reasonable to suppose they may range from a clarifying conversation, a request to reword, a downgrading of marks, article rejection or even a damaged reputation.

Therefore, we must address this issue now to provide clarity to authors, students' examiner committees, and editors. We believe that to subject software-generated methods descriptions to the scrutiny of plagiarism detection is to make a category error. When reporting methods, clarity and precision must be the objective, while novelty for its own sake can only serve to obscure the



- plagiarism software flags automatically generated method sections
- community gets together to gather information and draft statement to address the issue
- statement posted, continuing working together with journals





# FAIR data through community-driven development of standards and beyond



- necessities - repositories (standardized)
- **necessities - more than a PDF**
- necessities - reporting (standardized)
- necessities - documentation
- necessities - virtualization (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 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/>



\*[adapted from Fernando Perez](#)



# FAIR necessities - more than a PDF - Jupyter Books

jupyter {book}

Q 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



## 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 [.ipynb](#) 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.

Contents

Install Jupyter Book

Get started

A small example project

Under the hood - the components

of Jupyter Book

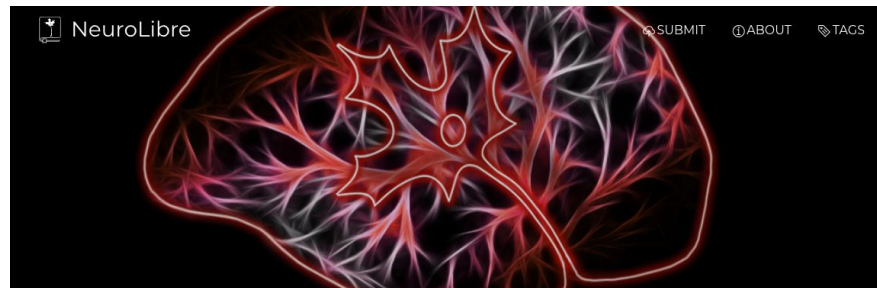
Contribute to Jupyter Book

Acknowledgements


<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

ABOUT

PAPERS

TOOLS

QUESTIONS

Log in with ORCID

Explore Papers

Documentation

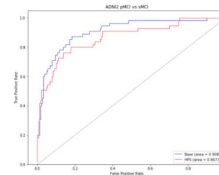
Learn More

<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.

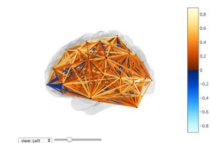
NeuroLibre Book GitHub Code



## 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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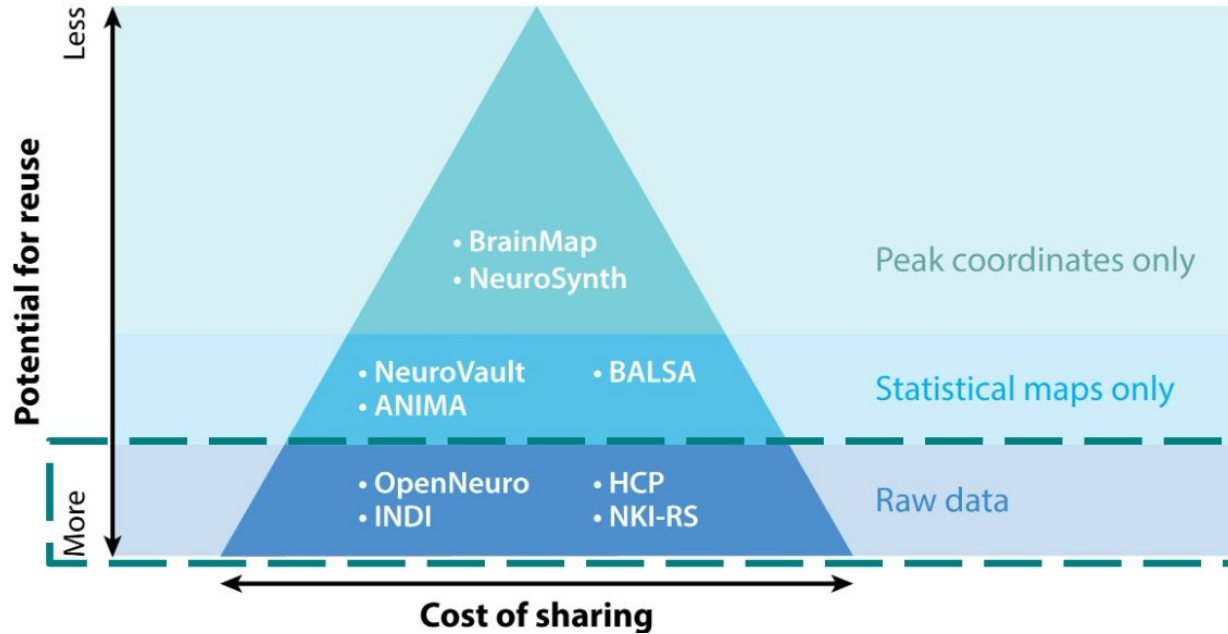


Bianca Kramer & Jeroen Bosman <https://101innovations.wordpress.com>

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



# FAIR data through community-driven development of standards and beyond



\*[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** Collections ▾ Metaanalyses ▾ About ▾ herholz.peer ▾ Search Search



# 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., Przysinda, 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:

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](#)

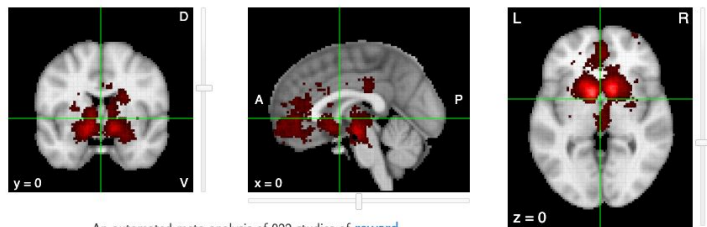


Neurosynth.org Home Meta-analyses Studies Locations Genes Decoder Code FAQs

## 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](#)

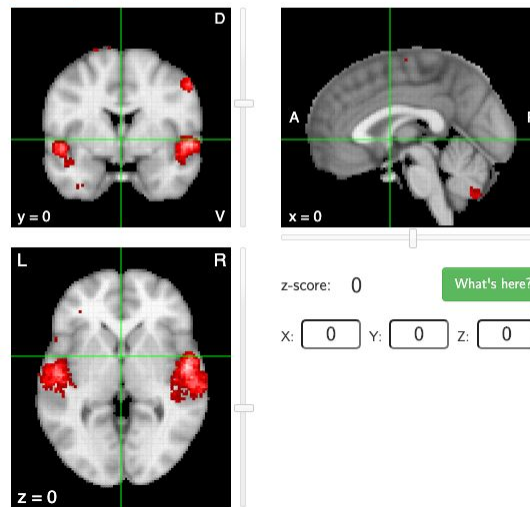
Neurosynth.org Home Meta-analyses Studies Locations Genes Decoder Code FAQs

## 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 ☐ Crosshairs  
Positive/Negative: ☐ Pan/zoom  
positive ☒ Labels

Thresholds:

0  0

Opacity:

1

z-score: 0

What's here?

X: 0 Y: 0 Z: 0



# 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

	Term	Similarity	Weight in brain map	N
In query	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



Predicted distribution of activations in the literature

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](#)

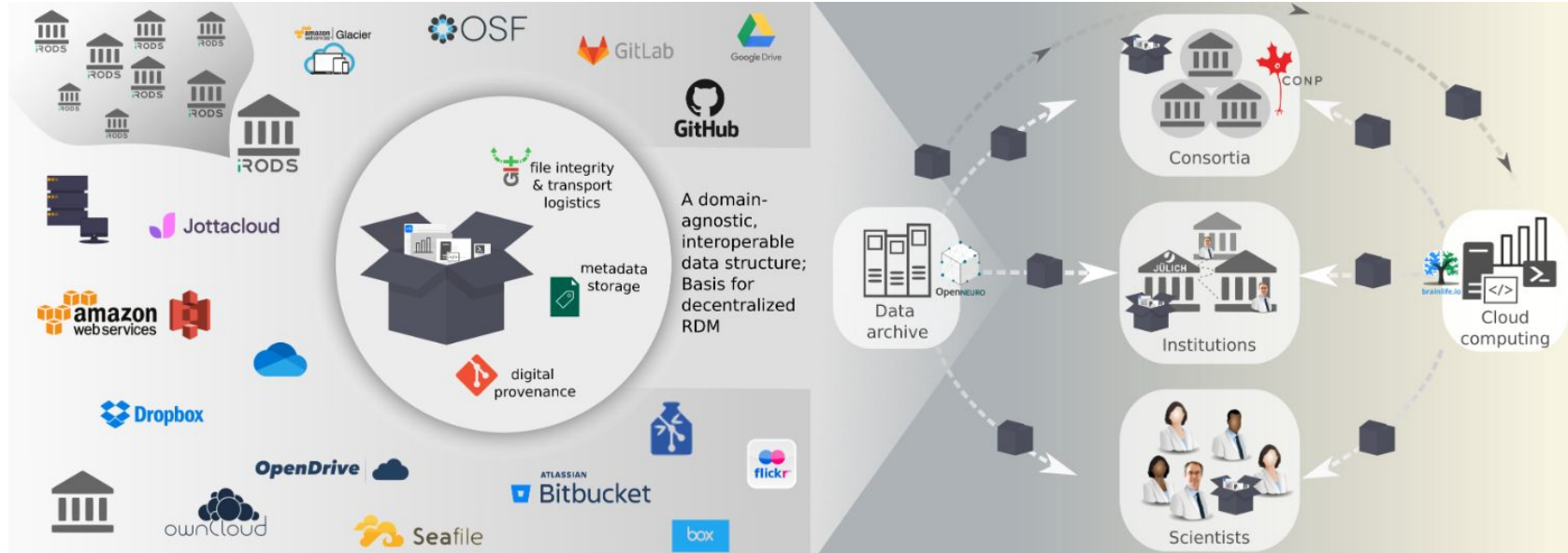


# FAIR data management

\*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.



## 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

# 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](#).





# 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
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- prerequisites - preregistration (standardized)



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\\_6pkJm](https://zenodo.org/record/3695300#_X8q1_6pkJm)

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





## ReproNim: A Center for Reproducible Neuroimaging Computation

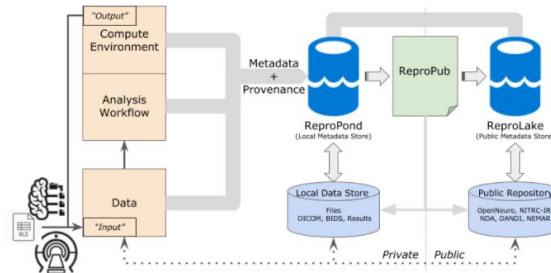
### Welcome to ReproNim!

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.





## 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.

## Welcome to INCF Neurostars.

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

## 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

### Announcements

162

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

### Community Chat

10

A place for friendly conversation, ideas, discussion, and otherwise un-categorizable things in the NeuroStars community.

1 unread

### Institutions

37

Category for Institutions that are partnered with INCF to discuss the institution-specific topics.

■ OCNS

### ABCD ReproNim

13

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



Datalad on github + cluster storage • datalad

2  
42m



BIDS and cookiecutter? • bids

0  
1h



OCNS Infrastructure/software/tools SIG: meet and greet, initial discussions • OCNS cns2020

64  
4h



Seed map - HELP from a newbie • Neuromatch Academy projects

0  
6h



Deadline extended to Nov. 15th Postdoctoral position for the development of acquisition, storage and processing pipelines for reproducible science building on the BIDS standard. • Announcements jobs

0  
20h

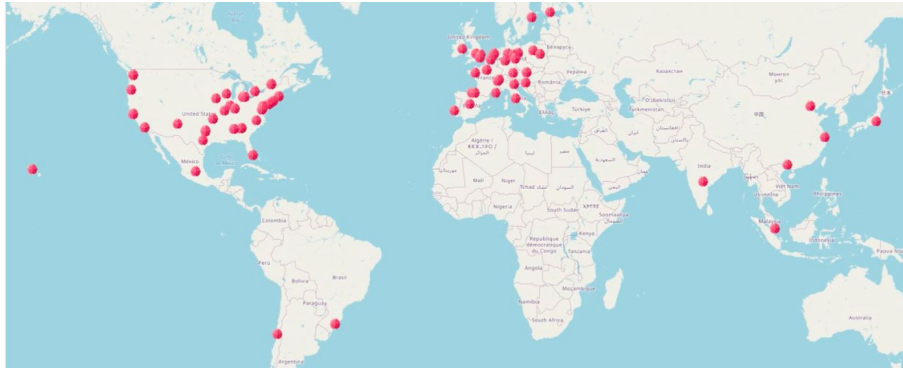


# FAIR initiatives & support

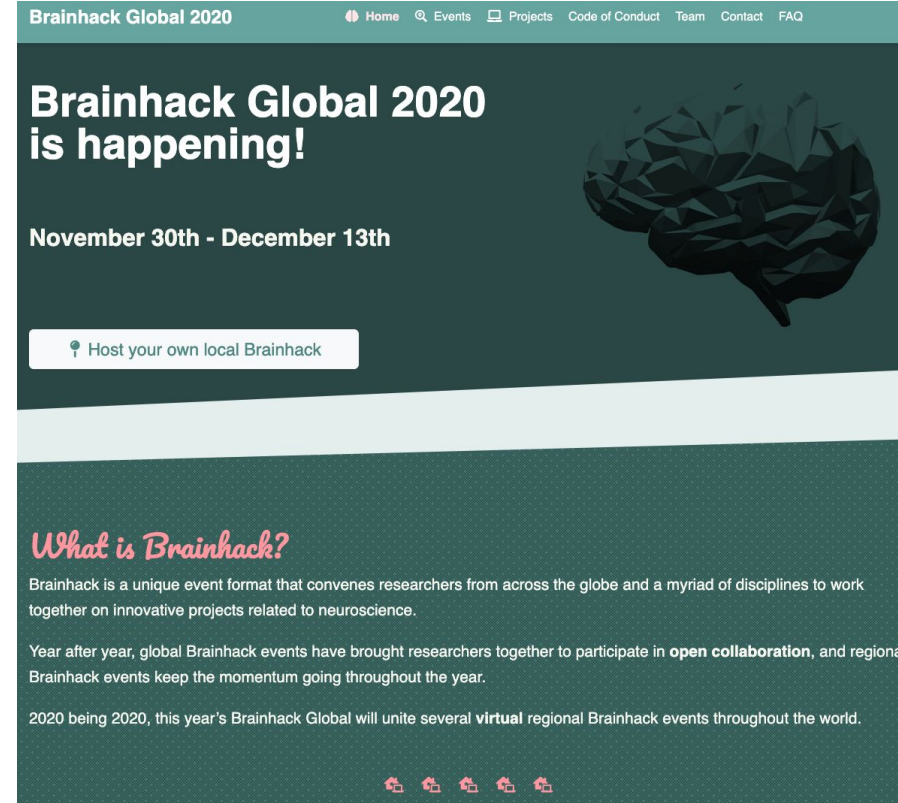
<https://brainhack.org/global2020/>



<http://brainhack.org/>



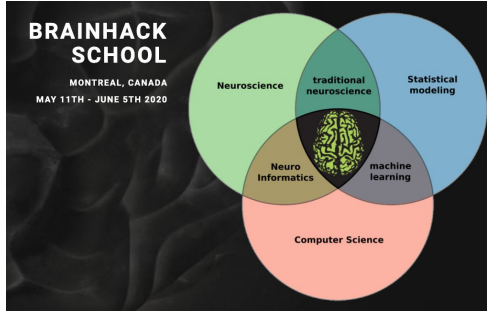
- started in 2012, spread across 25 countries, over 140 events, over 4000 members
- over 1000 attendees annually, focus on ECRs-collaboration-community-hacking-teaching
- model for various large scale teaching initiatives



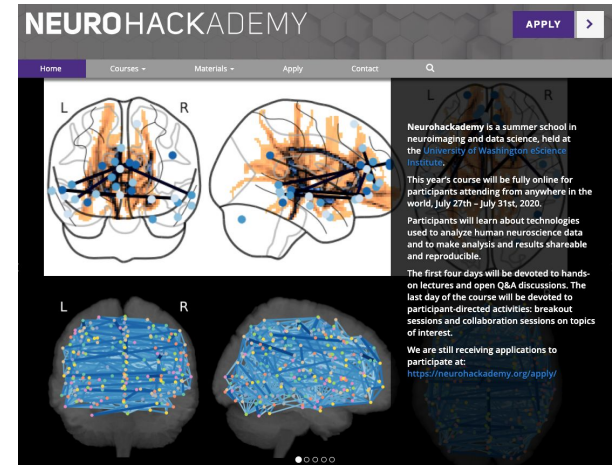


# FAIR initiatives & support

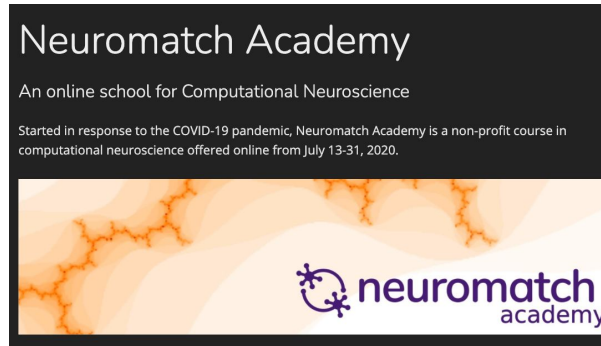
- **open** to everyone interested
- mostly **free** (if not, dedicated funding to support individuals from underrepresented and marginalized groups exists)
- **all materials publicly available** (including video recordings of lectures)
- **comprehensive, application oriented training**
- **decentralized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



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



<https://neurohackacademy.org/>



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



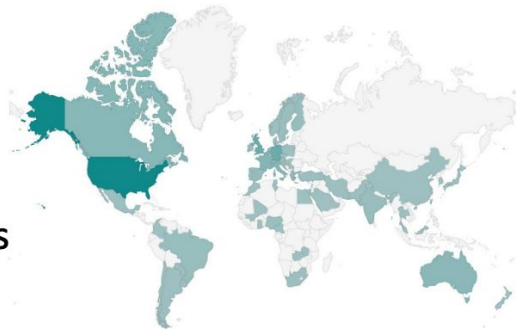
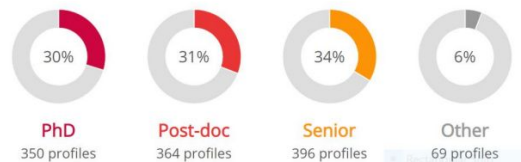


## Repository for Women in Neuroscience

- [www.winrepo.org](http://www.winrepo.org)
- over 1,100 profiles
- easy search
- recommendations

### Support the project:

- sign up
- spread the word
- submit recommendations



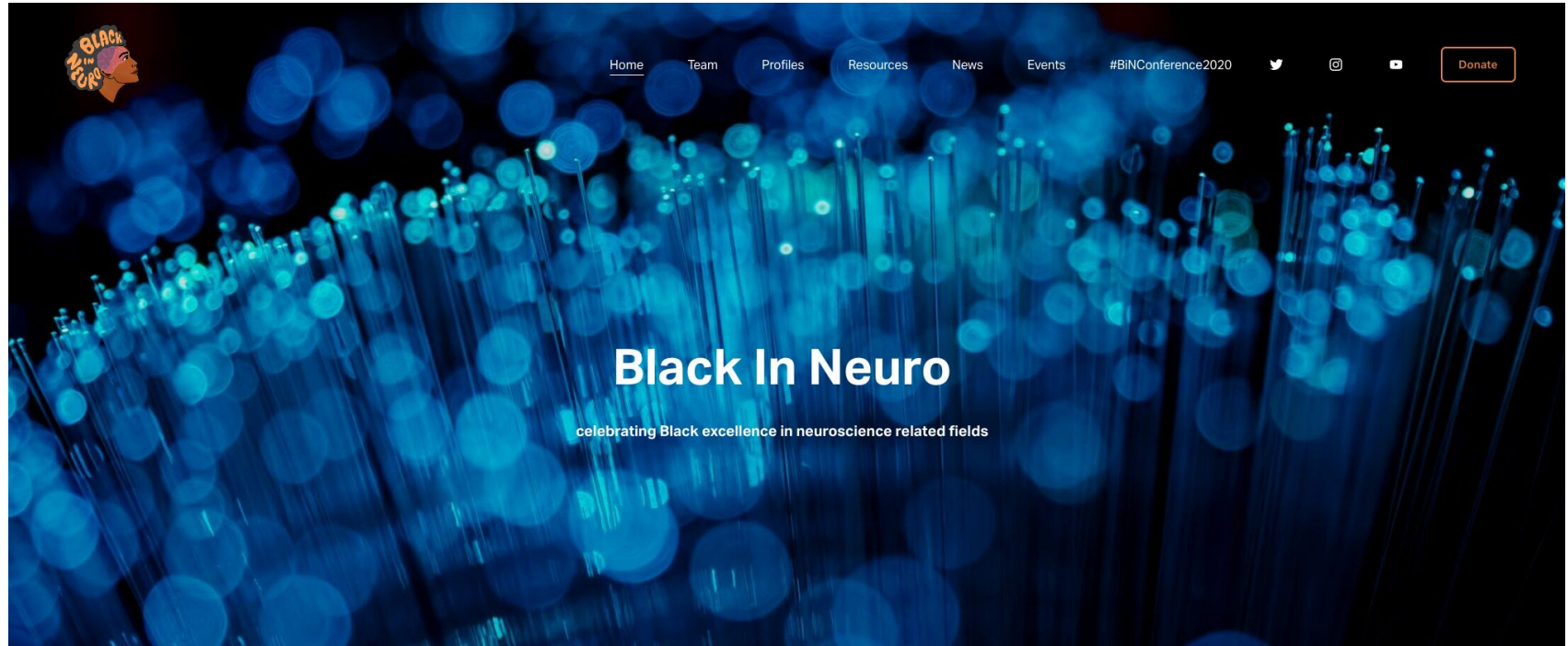
WINRePo1

[www.facebook.com/WiNRepository/](https://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/)



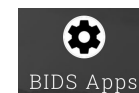
<https://ossig.netlify.app/>



# 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 ...



-----funding-----



-----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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