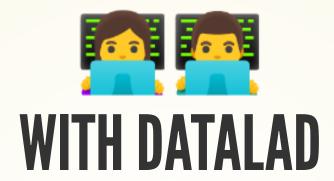
# DATA MANAGEMENT FOR NEUROIMAGING



### Adina Wagner



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Institute of Neuroscience and Medicine (INM-7)
Research Center Jülich

Slide sources: https://github.com/datalad-handbook/datalad-course/ Slide archive: doi.org/10.5281/zenodo.6880616

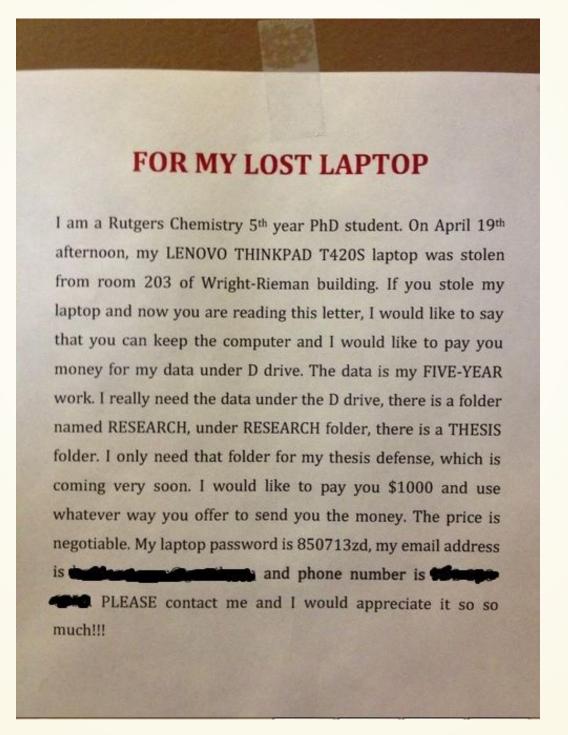
You write a paper & stay up late to generate good-looking figures, but you have to tweak many parameters and display options. The next morning, you have no idea which parameters produced which figures, and which of the figures fit to what you report in the paper.



# man on a state of the state of

# COMMON PROBLEMS IN SCIENCE

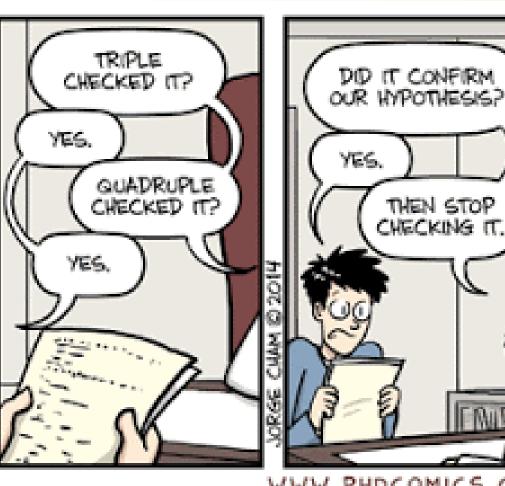
Your research project produces phenomenal results, but your laptop, the only place that stores the source code for the results, is stolen or breaks



A graduate student complains that a research idea does not work. Their supervisor can't figure out what the student did and how, and the student can't sufficiently explain their approach (data, algorithms, software). Weeks of discussion and mis-communication ensues because the supervisor can't first-hand explore or use the students project.







WWW.PHDCOMICS.COM

You wrote a script during your PhD that applied a specific method to a dataset.

Now, with new data and a new project, you try to reuse the script, but forgot how it worked.







www.phdcomics.com

You try to recreate results from another lab's published paper. You base your reimplementation on everything reported in their paper, but the results you obtain look nowhere like the original.



Scriberia

All these problems were paraphrased from Buckheit & Donoho, 1995

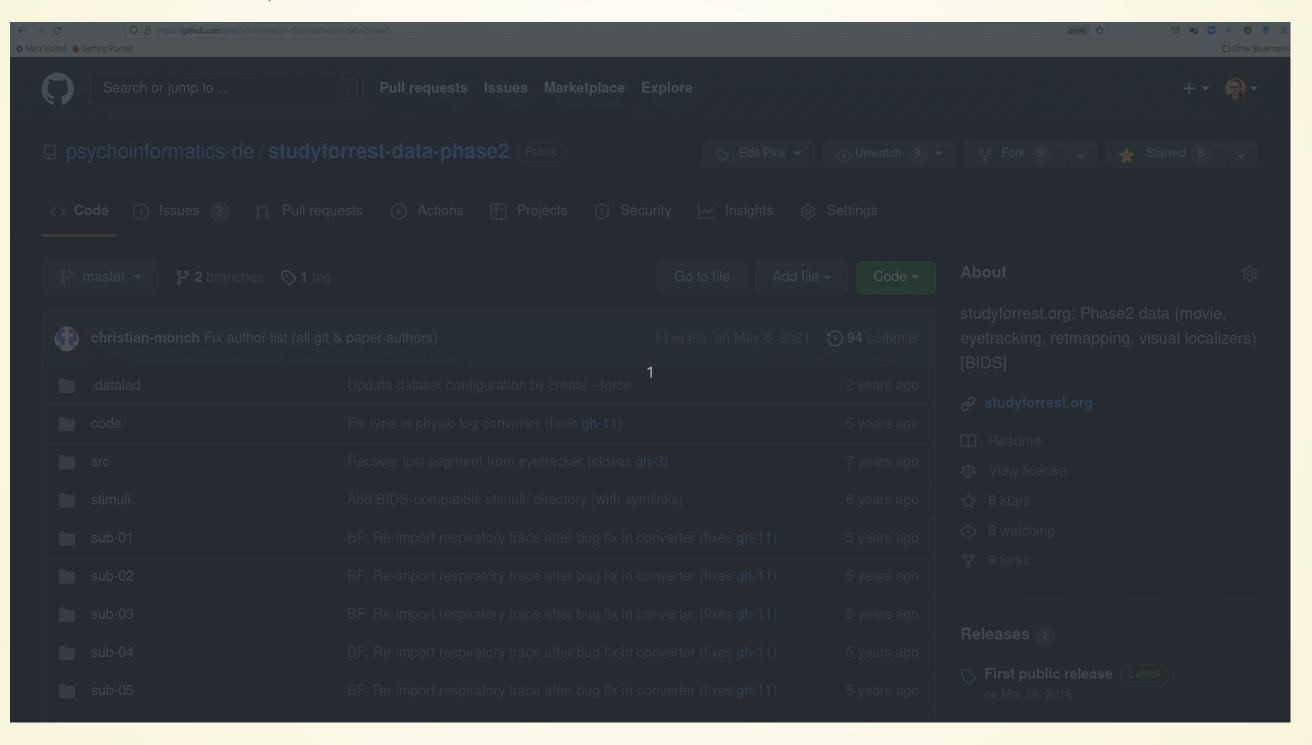


DataLad can help with small or large-scale data management

Free,
open source,
command line tool & Python API

Halchenko, Meyer, Poldrack, ... & Hanke, M. (2021). DataLad: distributed system for joint management of code, data, and their relationship. Journal of Open Source Software, 6(63), 3262.

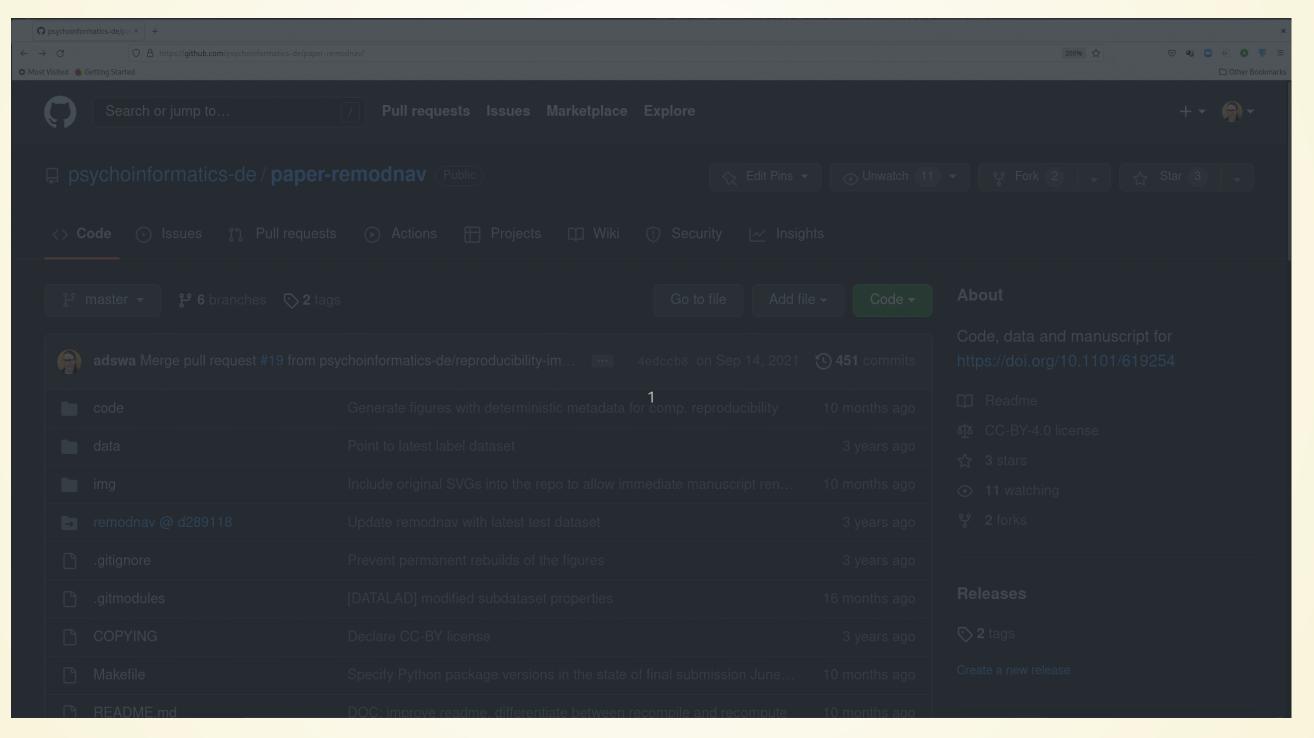
 Publish or consume datasets via GitHub, GitLab, OSF, the European Open Science Cloud, or similar services



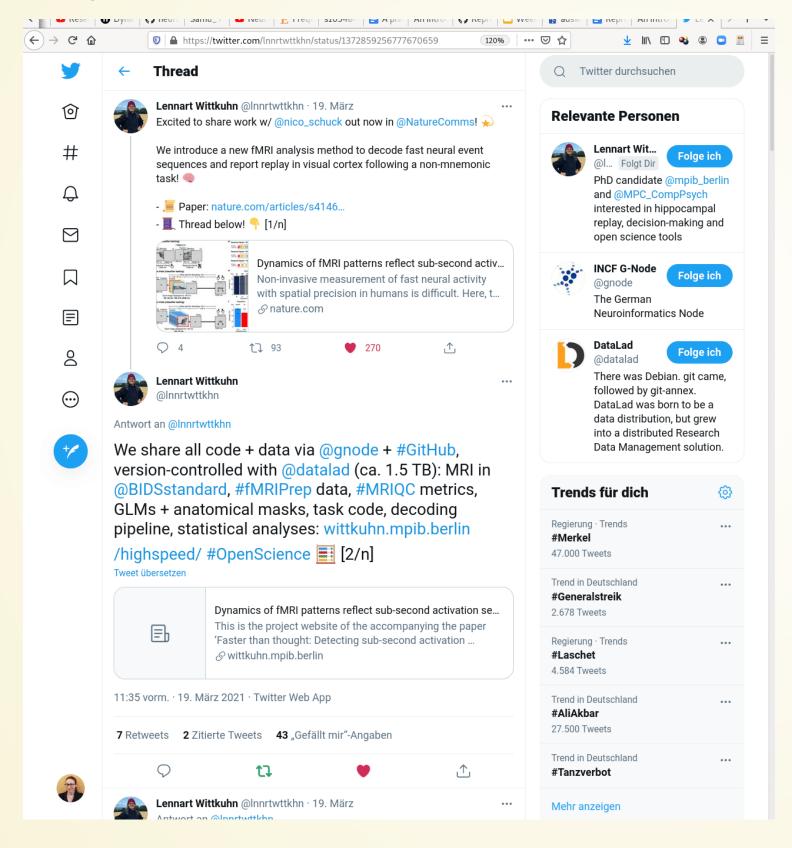
 Behind-the-scenes infrastructure component for data transport and versioning (e.g., used by OpenNeuro, brainlife.io, the Canadian Open Neuroscience Platform (CONP), CBRAIN)



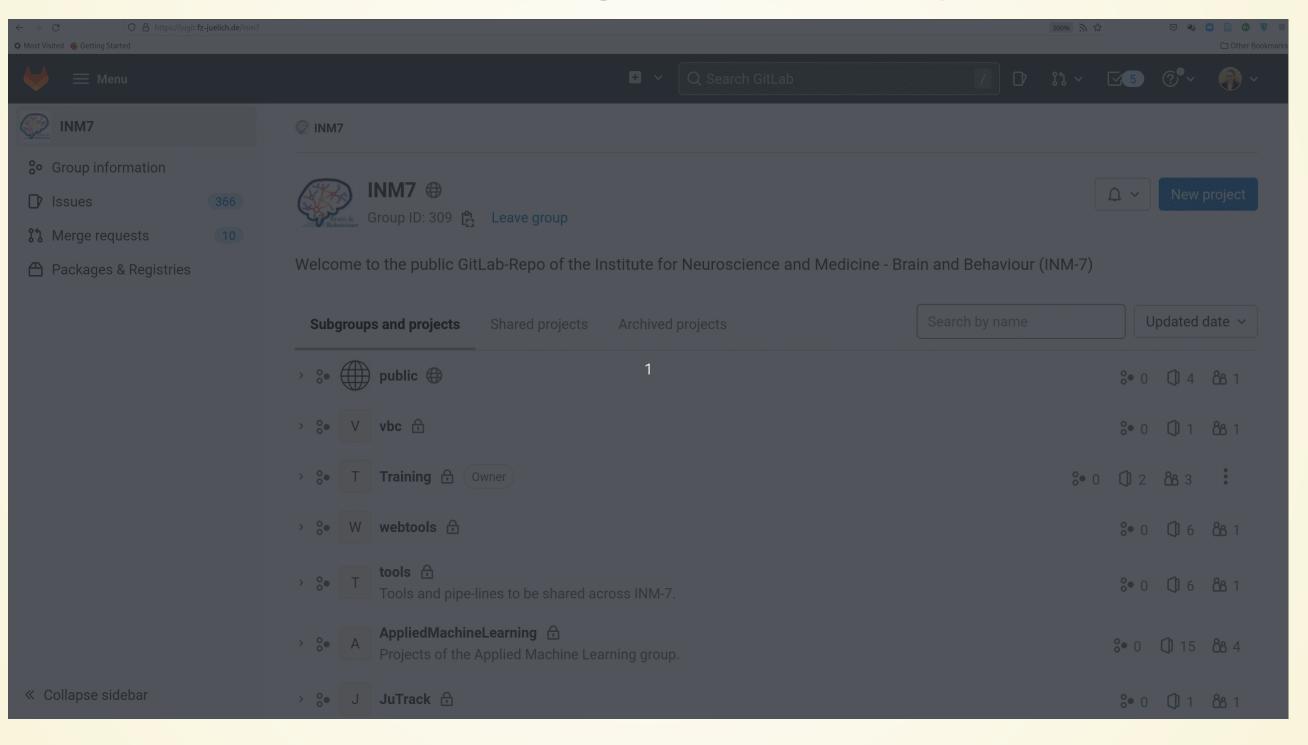
 Creating and sharing reproducible, open science: Sharing data, software, code, and provenance



 Creating and sharing reproducible, open science: Sharing data, software, code, and provenance



Central data management and archival system





- Joint version control (Git, git-annex): version control data & software alongside your code
- Provenance capture: Create and share machine-readable, re-executable provenance records for reproducible, transparent, and FAIR research
- decentral data transport mechanisms: Install, share and collaborate on scientific projects; publish, upgrade, and retrieve their contents in a streamlined fashion on demand, and distribute files in a decentral network on the services or infrastructures of your choice

Code for hands-on: handbook.datalad.org

# PREREQUISITES: TERMINAL

DataLad can be used from the command line

datalad create mydataset

... or with its Python API

```
import datalad.api as dl
dl.create(path="mydataset")
```

... and other programming languages can use it via system call

```
# in R
> system("datalad create mydataset")
```

# PREREQUISITES: USING DATALAD

 Every DataLad command consists of a main command followed by a sub-command. The main and the sub-command can have options.

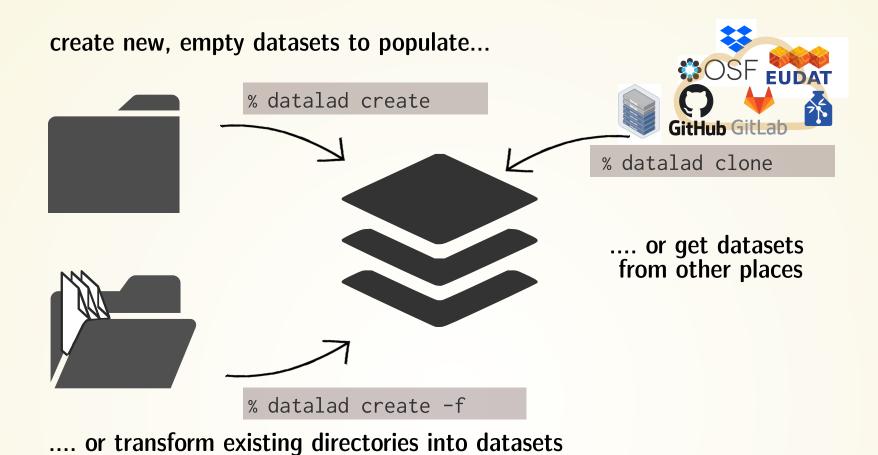
```
datalad [--GLOBAL-OPTION <opt. flag spec.>] COMMAND [ARGUMENTS] [--OPTION <opt. flag spec>]
                                                                                                                                        Each datalad
                                                                                                                                        invocation can
                                                                                                                                        have two sets of
                                                               -d/--dataset
                                                                                 Dataset location: path to root, or ^ for superdataset
                                                          COMMAND OPTIONS
-c KEY=VALUE
                                                                                                                                        options: general
   Set config variables (overrides configurations in files)
                                                              -D/--description A location description (e.g., "my backup server")
                                                                                                                                        options are given
-f/--output-format default|json|json_pp|tailored
                                                                                 Force execution of a command (Dangerzone!)
                                                              -f/--force
                                                                                                                                        first, command-
   Specify the format for command result renderung
                                                                                 A description about a change made to the dataset
                                                                                                                                        specific ones go
-l/--log-level critical|error|warning|info|debug
                                                                                 Perform an operation recursively across subdatasets
                                                                                                                                        after the
                                                              -r/--recursive
   Set logging verbosity level
                                                                                                                                        subcommand.
                                                              -R/--recursion-limit <n> Limit recursion to n subdataset levels
```

Example (main command, subcommand, several subcommand options):

```
$ datalad save -m "Saving changes" --recursive
```

 Use --help to find out more about any (sub)command and its options, including detailed description and examples (q to close). Use -h to get a short overview of all options

# **EVERYTHING HAPPENS IN DATALAD DATASETS**



- Look and feel like a directory on your computer
- content agnostic
- no custom data structures

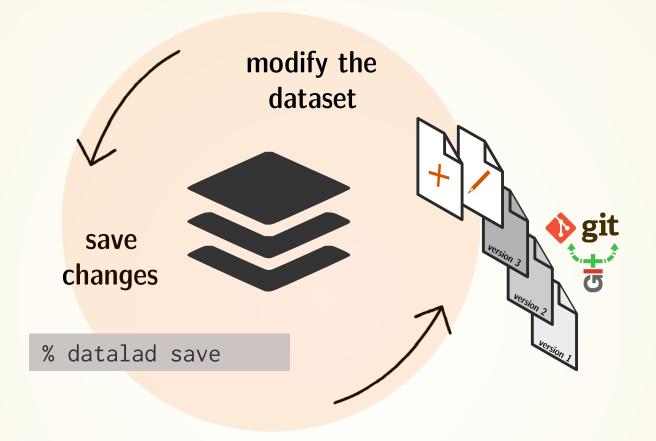


Terminal view

File viewer

# DATASET = GIT/GIT-ANNEX REPOSITORY

version control files regardless of size or type



### Stay flexible:

- Non-complex DataLad core API (easy for data management novices)
- Pure Git or git-annex commands (for regular Git or git-annex users, or to use specific functionality)

The building blocks of a scientific result are rarely static

Analysis code evolves

(Fix bugs, add functions, refactor, ...)

### The building blocks of a scientific result are rarely static

### Data changes

(errors are fixed, data is extended, naming standards change, an analysis requires only a subset of your data...)

The building blocks of a scientific result are rarely static

Data changes (for real)

(errors are fixed, data is extended, naming standards change, ...)

"Shit, which version of which script produced these outputs from which version of what data... and which software version?"



### 1. TRANSPARENCY - FOR DATA

Once you track changes to data with version control tools, you can find out why it changed, what has changed, when it changed, and which version of your data was used at which point in time.

```
o [DATALAD RUNCMD] add non-defaced commit 6da25fb6fee2c698d35f52066698b6f94850f4d2
                                      [DATALAD RUNCMD] reconvert DICOM
                                      o [master] {origin/HEAD} {origin/m
                                      o Enable DataLad metadata extracto AuthorDate: Fri Jan 19 14:09:53 2018 +0100
                                      [DATALAD] new dataset
                                      o [DATALAD] Set default backend fo
                                                                         CommitDate: Fri Jan 19 14:11:23 2018 +0100
                                      o <v1.5> Update changelog for 1.5
                                                                             BF: Re-import respiratory trace after bug fix in converter (fixes gh
                                      o BF: Re-import respiratory trace
                                      o Fix type in physio log converter
                                      o ENH: Report per-stimulus events
                                                                             .er task-movielocalizer run-1 recording-cardresp physio.tsv.gz |
                                      o Add BIDS-compatible stimuli/ dir
                                      o Minor tweaks to gaze overlay scr
                                      o Add "TaskName" meta data field f
                                      o Add task-* physio.json files
                                      o BF: Fix task label in file names
                                      O Update changelog
                                      o Add cut position information to
                                      o {origin/ } Mention openfmri as d
                                      O Update publication links
                                      o Disable invalid test
main] 6da25fb6fee2c698d35f52066698b6f94850f4d2 - commit 10 of
                                                                          [diff] 6da25fb6fee2c698d35f52066698b6f94850f4d2 - line 1 of 2391
```

# DIGITAL PROVENANCE

- = "The tools and processes used to create a digital file, the responsible entity, and when and where the process events occurred"
- Have you ever saved a PDF to read later onto your computer, but forgot where you got it from? Or did you ever find a figure in your project, but forgot which analysis step produced it?

# PROVENANCE AND REPRODUCIBILITY

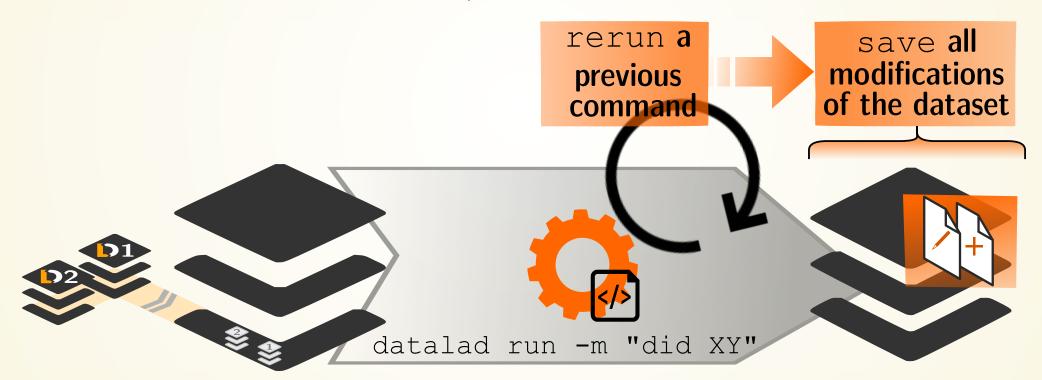
datalad run wraps around anything expressed in a command line call and saves the dataset modifications resulting from the execution



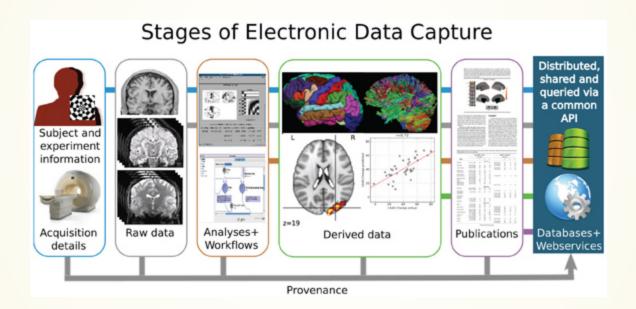
# PROVENANCE AND REPRODUCIBILITY

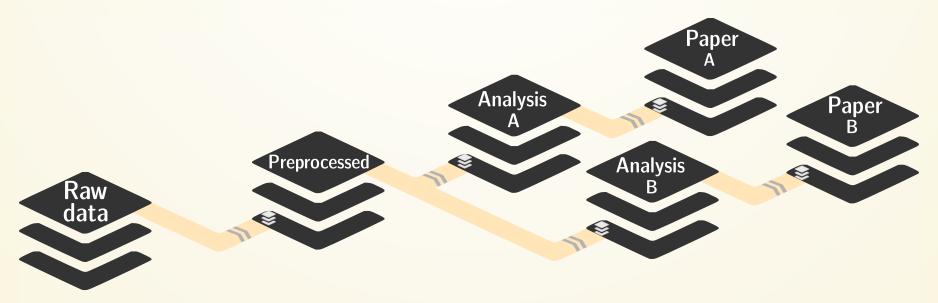
datalad rerun repeats captured executions.

If the outcomes differ, it saves a new state of them.



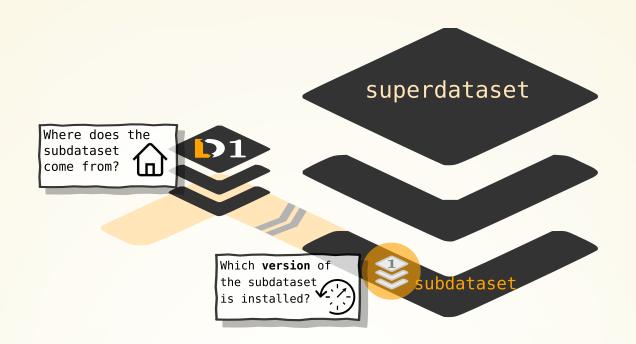
# SEAMLESS DATASET NESTING & LINKAGE





Nest modular datasets to create a linked hierarchy of datasets, and enable recursive operations throughout the hierarchy

# SEAMLESS DATASET NESTING & LINKAGE



\$ datalad clone --dataset . http://example.com/ds inputs/rawdata

# PLENTY OF DATA, BUT LITTLE DISK-USAGE

 Cloned datasets are lean. "Meta data" (file names, availability) are present, but no file content:

```
$ datalad clone git@github.com:psychoinformatics-de/studyforrest-data-phase2.git
install(ok): /tmp/studyforrest-data-phase2 (dataset)
$ cd studyforrest-data-phase2 && du -sh
18M .
```

files' contents can be retrieved on demand:

```
$ datalad get sub-01/ses-movie/func/sub-01_ses-movie_task-movie_run-1_bold.nii.gz
get(ok): /tmp/studyforrest-data-phase2/sub-01/ses-movie/func/sub-01_ses-movie_task-movie_run-1
```

Have access to more data on your computer than you have disk-space:

```
# eNKI dataset (1.5TB, 34k files):
$ du -sh
1.5G .
# HCP dataset (~200TB, >15 million files)
$ du -sh
48G .
```

# PLENTY OF DATA, BUT LITTLE DISK-USAGE

Drop file content that is not needed:

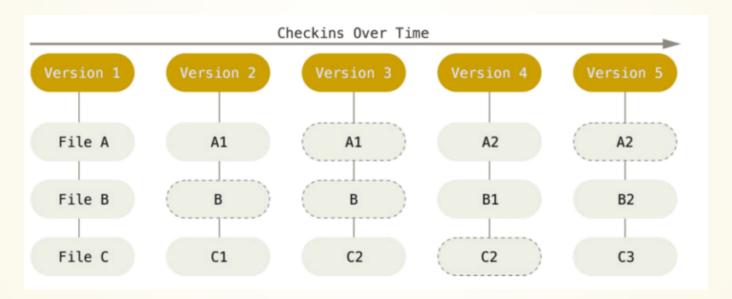
```
$ datalad drop sub-01/ses-movie/func/sub-01_ses-movie_task-movie_run-1_bold.nii.gz
drop(ok): /tmp/studyforrest-data-phase2/sub-01/ses-movie/func/sub-01_ses-movie_task-movie_run-1_
```

When files are dropped, only "meta data" stays behind, and they can be reobtained on demand.

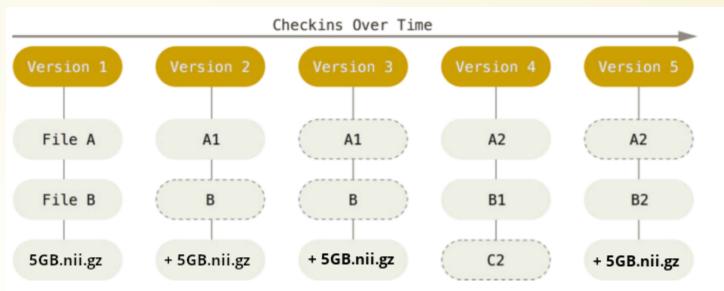
```
dl.get('input/sub-01')
[really complex analysis]
dl.drop('input/sub-01')
```

## THERE ARE TWO VERSION CONTROL TOOLS AT WORK - WHY?

Git does not handle large files well.



### THERE ARE TWO VERSION CONTROL TOOLS AT WORK - WHY?



Git does not handle large files well.

And repository hosting services refuse to handle large files:

```
adina@muninn in /tmp/myresearch on git:master
) git push gh-adswa master
Enumerating objects: 3, done.
Counting objects: 100% (3/3), done.
Delta compression using up to 8 threads
Compressing objects: 100% (2/2), done.
Writing objects: 100% (3/3), 497.87 KiB | 161.00 KiB/s, done.
Total 3 (delta 0), reused 0 (delta 0), pack-reused 0
remote: arror: Trace: 64a78dd4lece8e5493fe33f97397a7a90ef9c91260ba32786970dbdcf5c4e0dd
remote: error: See http://git.io/iEPt8g for more information.
remote: error: File output.dat is 500.00 MB; this exceeds GitHub's file size limit of 100.00 MB
remote: error: GH001: Large files detected. You may want to try Git Large File Storage - https://git-lfs.github.
To github.com:adswa/myresearch.git
! [remote rejected] master -> master (pre-receive hook declined)
error: failed to push some refs to 'dithub.com:adswa/myresearch.git'
```

git-annex to the rescue! Let's take a look how it works

# GIT VERSUS GIT-ANNEX



- dataset history (commit messages, run records)
- All files + content committed into Git (useful with code, text, ...)
- File identity information of all annexed files (file name, identity hash, storage locations where to retrieve it from)

- organized in the "annex" or "object tree" of the dataset

# DATASET INTERNALS

Where the filesystem allows it, annexed files are symlinks:

```
$ ls -l sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz
lrwxrwxrwx 1 adina adina 142 Jul 22 19:45 sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz ->
../../.git/annex/objects/kZ/K5/MD5E-s24180157--aeb0e5f2e2d5fe4ade97117a8cc5232f.nii.gz/MD5E-s24180
--aeb0e5f2e2d5fe4ade97117a8cc5232f.nii.gz
```

(PS: especially useful in datasets with many identical files)

The symlink reveals this internal data organization based on identity hash:

```
$ md5sum sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz
aeb0e5f2e2d5fe4ade97117a8cc5232f sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz
```

 The (tiny) symlink instead of the (potentially large) file content is committed version controlling precise file identity without checking contents into Git

```
diff --git a/sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz b/sub-02/func/sub-02_task-oneback_run-01_bold.nii.
new file mode 120000
index 00000000..398e7f1
--- /dev/null
+++ b/sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz
@@ -0,0 +1 @@
+../../.git/annex/objects/kZ/K5/MD5E-s24180157--aeb0e5f2e2d5fe4ade97117a8cc5232f.nii.gz/MD5E-s24180157--aeb0e5f2e2
```

 File contents can be shared via almost all standard infrastructure. File availability information is a decentral network. A file can exist in multiple different locations.

# GIT VERSUS GIT-ANNEX

# Data in datasets is either stored in Git or git-annex

By default, everything is annexed.

### Two consequences:

- Annexed contents are not available right after cloning, only content identity and availability information (as they are stored in Git). Everything that is annexed needs to be retrieved with datalad get from whereever it is stored.
- Files stored in Git are modifiable, annexed files are protected against accidental modifications

Git	git-annex
handles <b>small</b> files well (text, code)	handles <b>all</b> types and sizes of files well
file contents are in the Git history and will be shared upon git/datalad push	file contents are in the annex. Not necessarily shared
Shared with every dataset clone	Can be kept private on a per-file level when sharing the dataset
Useful: Small, non-binary, frequently modified, need-to-be-accessible (DUA, README) files	Useful: Large files, private files

Useful background information for demo later. Read this handbook chapter for details

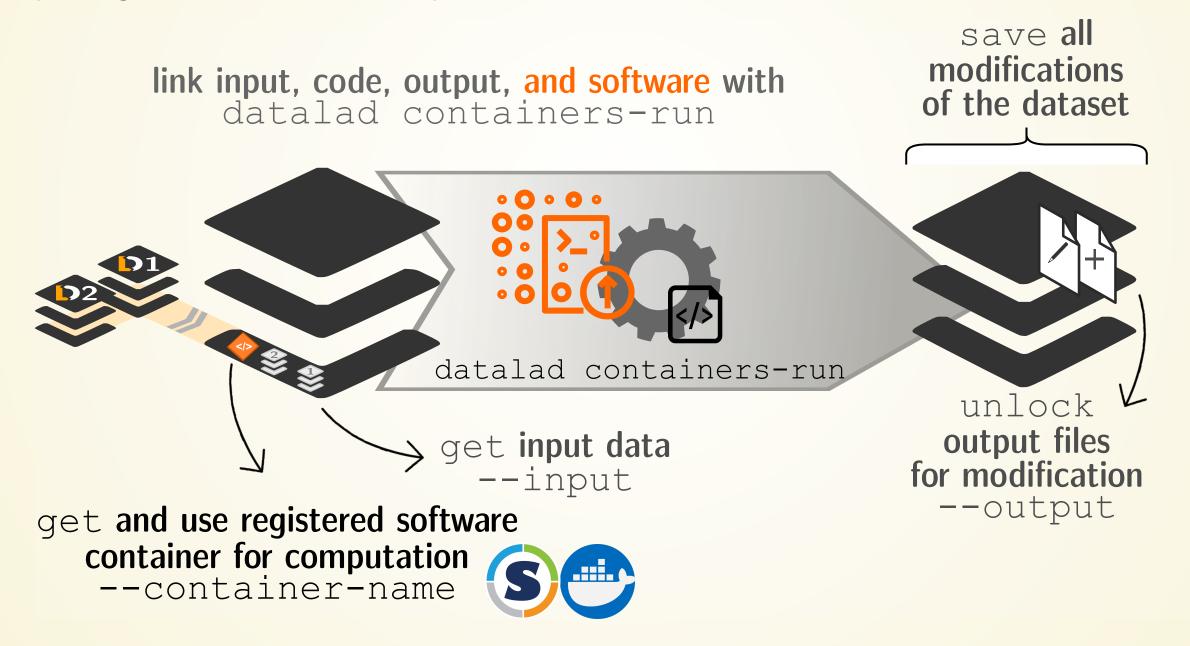
# GIT VERSUS GIT-ANNEX

Users can decide which files are annexed:

- Pre-made run-procedures, provided by DataLad (e.g., text2git, yoda) or created and shared by users (Tutorial)
- Self-made configurations in .gitattributes (e.g., based on file type, file/path name, size, ...; rules and examples)
- Per-command basis (e.g., via datalad save --to-git)

### COMPUTATIONAL PROVENANCE

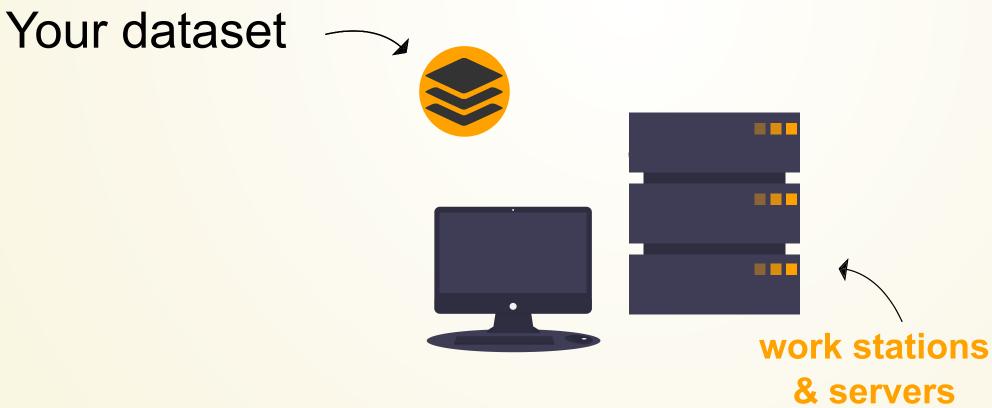
The datalad-container extension gives DataLad commands to register software containers
as "just another file" to your dataset, and datalad containers-run analysis inside the container,
capturing software as additional provenance

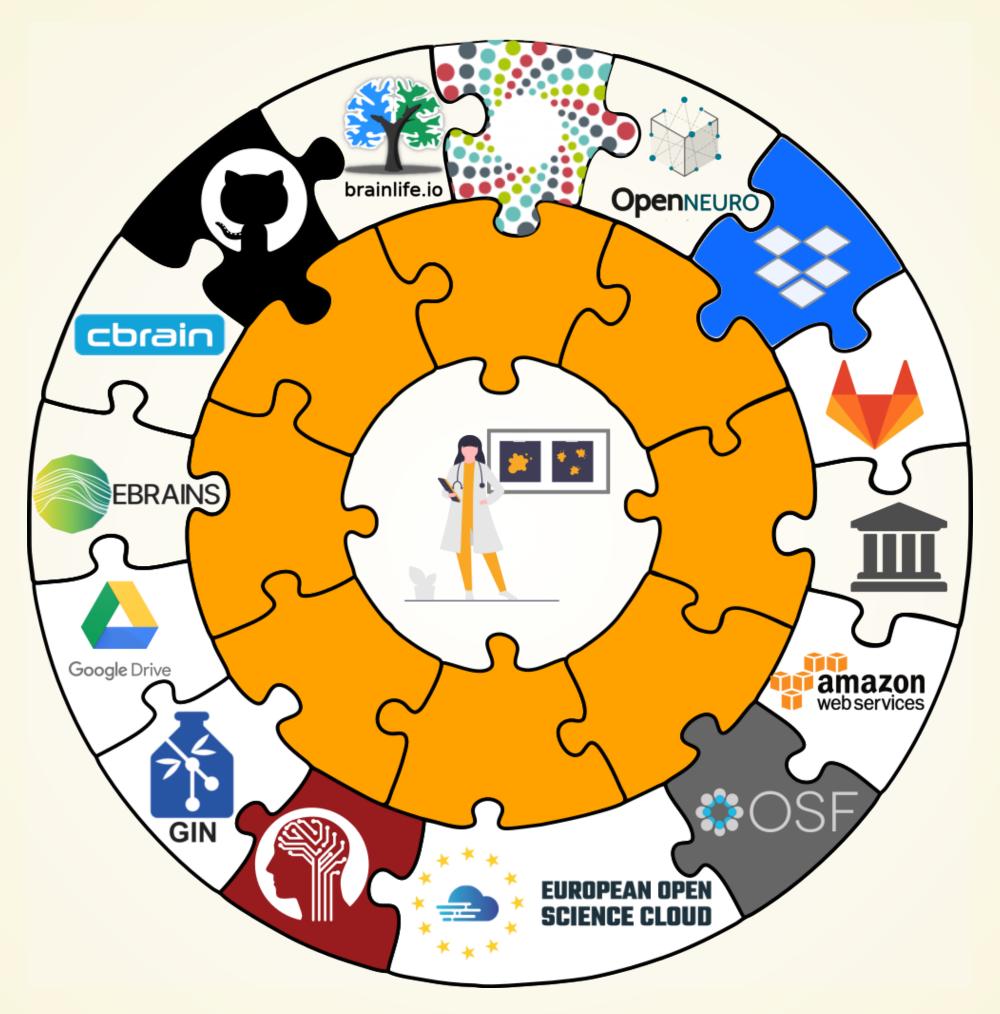


## **SHARING DATASETS**



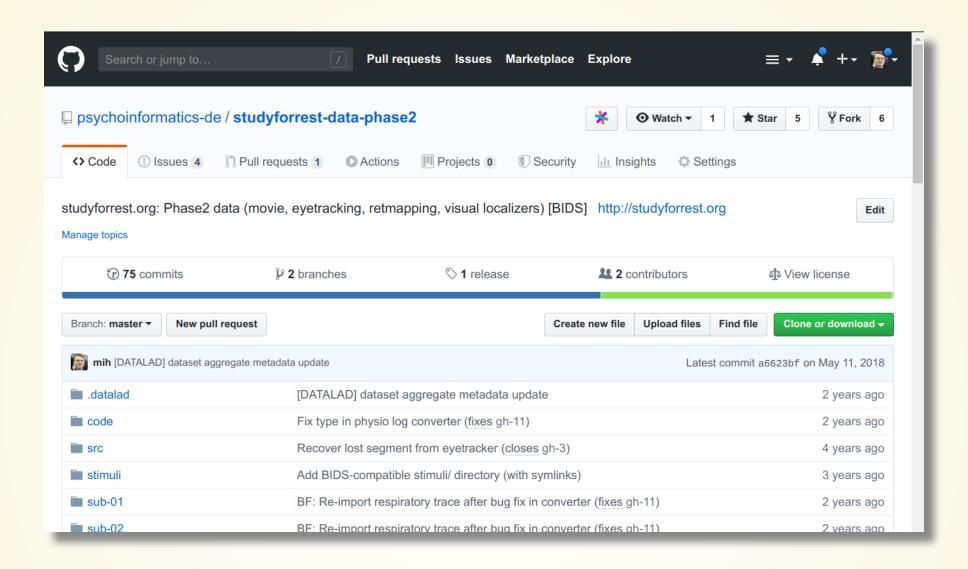






Apart from local computing infrastructure (from private laptops to computational clusters), datasets can be hosted in major third party repository hosting and cloud storage services. More info: Chapter on Third party infrastructure.

## **SERVICES**

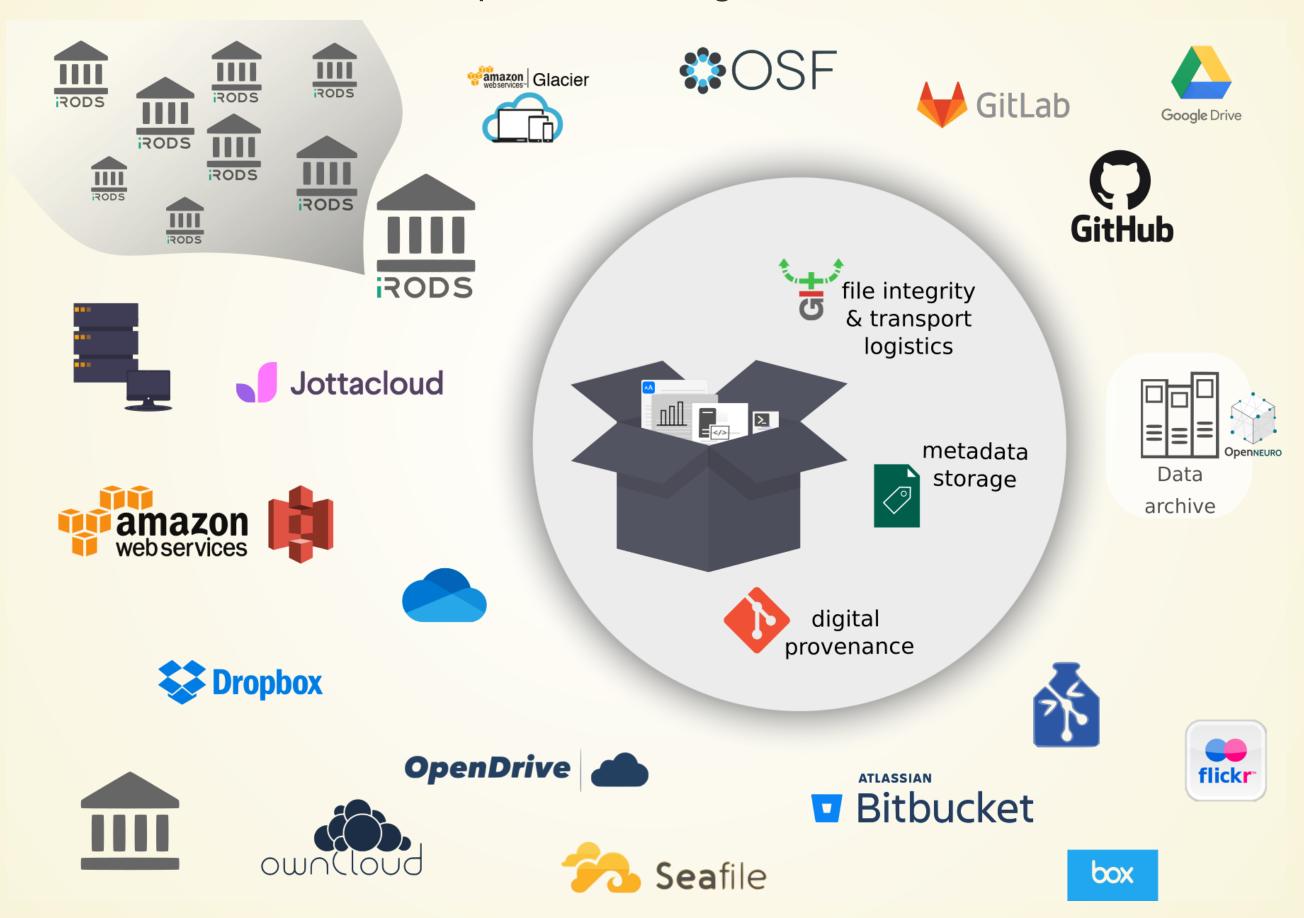


- make the difference for advertisment, discovery, convenience
- but imply gigantic dependencies
- often impossible to "take over"

Make sure data/metadata are self-contained to facilitate/enable transition to another service

### **SECURITY AND RELIABILITY - FOR DATA**

Decentral version control for data integrates with a variety of services to let you store data in different places - creating a resilient network for data



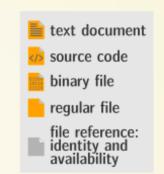
### **COLLABORATION**

#### Teamscience on more than code:



### **EXHAUSTIVE TRACKING OF RESEARCH COMPONENTS**



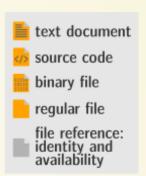


Well-structured datasets (using community standards), and portable computational environments — and their evolution — are the precondition for reproducibility

```
# turn any directory into a dataset
# with version control
# file content of any size
% datalad create <directory>
% datalad save
```

## CAPTURE COMPUTATIONAL PROVENANCE



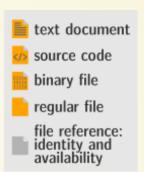


Which data was needed at which version, as input into which code, running with what parameterization in which computional environment, to generate an outcome?

```
# execute any command and capture its output
# while recording all input versions too
% datalad run --input ... --output ... <command>
```

### **EXHAUSTIVE CAPTURE ENABLES PORTABILITY**



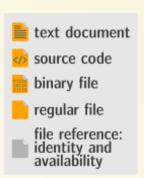


Precise identification of data and computational environments combined with provenance records form a comprehensive and portable data structure, capturing all aspects of an investigation.

```
# transfer data and metadata to other sites and services
# with fine-grained access control for dataset components
% datalad push --to <site-or-service>
```

### REPRODUCIBILITY STRENGTHENS TRUST





Outcomes of computational transformations can be validated by authorized 3rd-parties. This enables audits, promotes accountability, and streamlines automated "upgrades" of outputs

```
# obtain dataset (initially only identity,
# availability, and provenance metadata)

% datalad clone <url>
# immediately actionable provenance records
# full abstraction of input data retrieval
% datalad rerun <commit|tag|range>
```

# **ULTIMATE GOAL: (RE-)USABILITY**



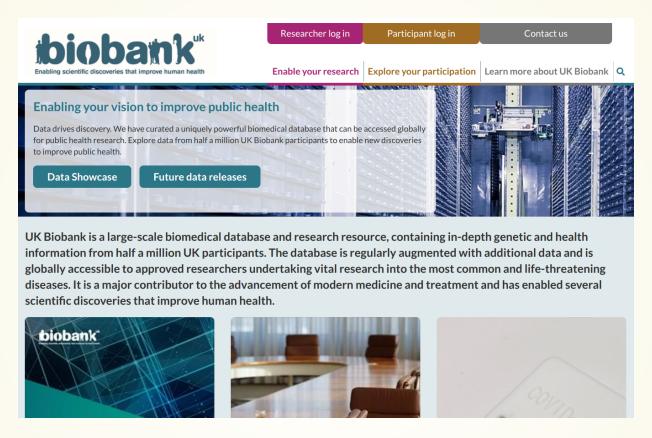
Verifiable, portable, self-contained data structures that track all aspects of an investigation exhaustively can be (re-)used as modular components in larger contexts — propagating their traits

```
# declare a dependency on another dataset and
# re-use it a particular state in a new context
% datalad clone -d <superdataset> <url> <path-in-dataset>
```

# **BIG DATA**

### FAIRLY BIG: SCALING UP

Objective: Process the UK Biobank (imaging data)

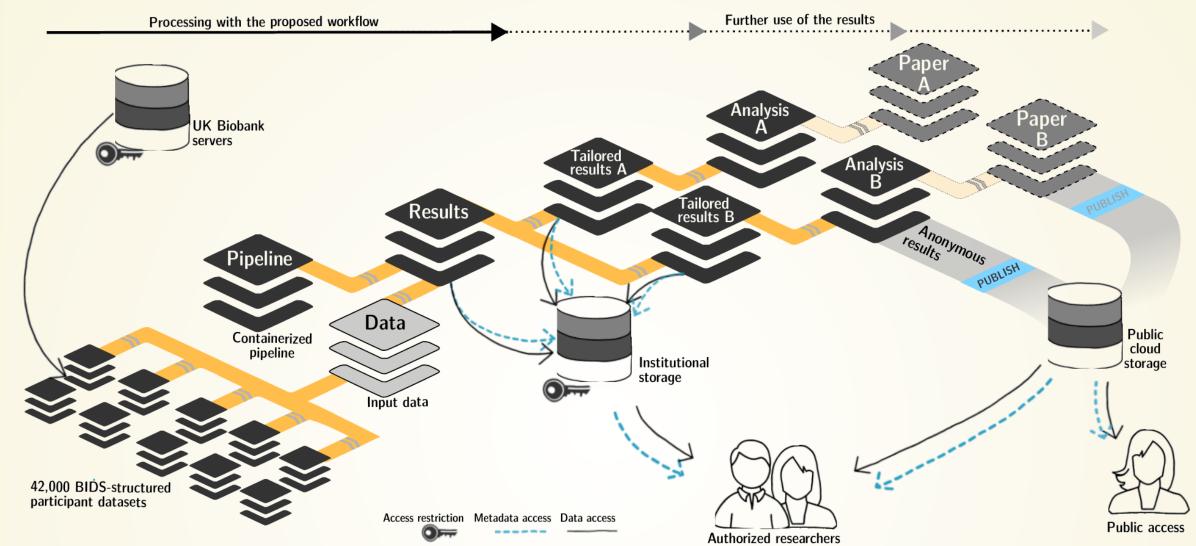


- 76 TB in 43 million files in total
- 42,715 participants contributed personal health data
- Strict DUA
- Custom binary-only downloader
- Most data records offered as (unversioned) ZIP files

### **CHALLENGES**

- Process data such that
  - Results are computationally reproducible (without the original compute infrastructure)
  - There is complete linkage from results to an individual data record download
  - It scales with the amount of available compute resources
- Data processing pipeline
  - Compiled MATLAB blob
  - 1h processing time per image, with 41k images to process
  - 1.2 M output files (30 output files per input file)
  - 1.2 TB total size of outputs

## FAIRLY BIG SETUP

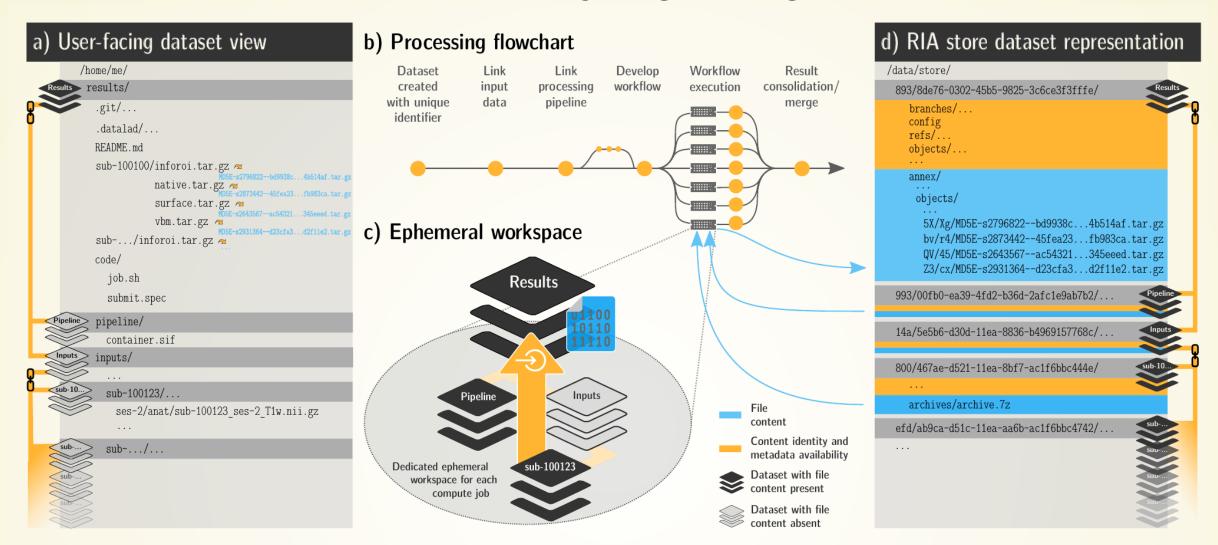


#### **Exhaustive tracking**

- datalad-ukbiobank extension downloads, transforms & track the evolution of the complete data release in DataLad datasets
- Native and BIDSified data layout (at no additional disk space usage)
- Structured in 42k individual datasets, combined to one superdataset
- Containerized pipeline in a software container
- Link input data & computational pipeline as dependencies

Wagner, Waite, Wierzba et al. (2021). FAIRly big: A framework for computationally reproducible processing of large-scale data.

### **FAIRLY BIG WORKFLOW**

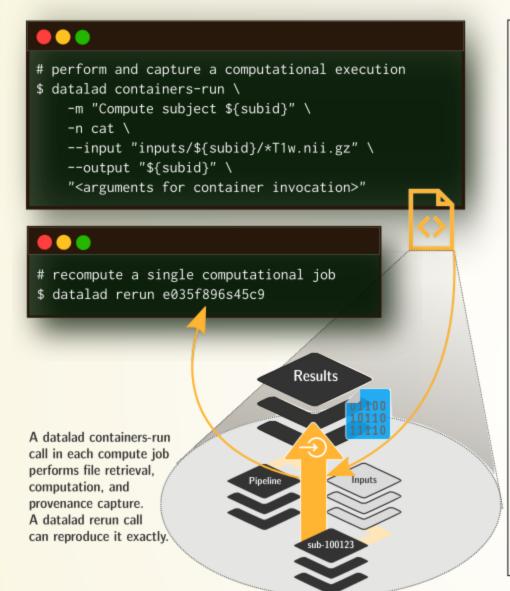


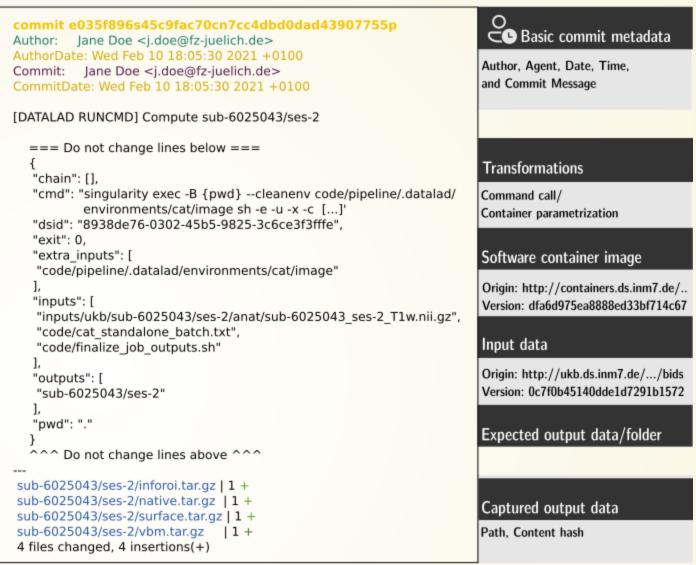
#### portability

- Parallel processing: 1 job = 1 subject (number of concurrent jobs capped at the capacity of the compute cluster)
- Each job is computed in a ephemeral (short-lived) dataset clone, results are pushed back: Ensure exhaustive tracking & portability during computation
- Content-agnostic persistent (encrypted) storage (minimizing storage and inodes)
- Common data representation in secure environments

Wagner, Waite, Wierzba et al. (2021). FAIRly big: A framework for computationally reproducible processing of large-scale data.

### FAIRLY BIG PROVENANCE CAPTURE



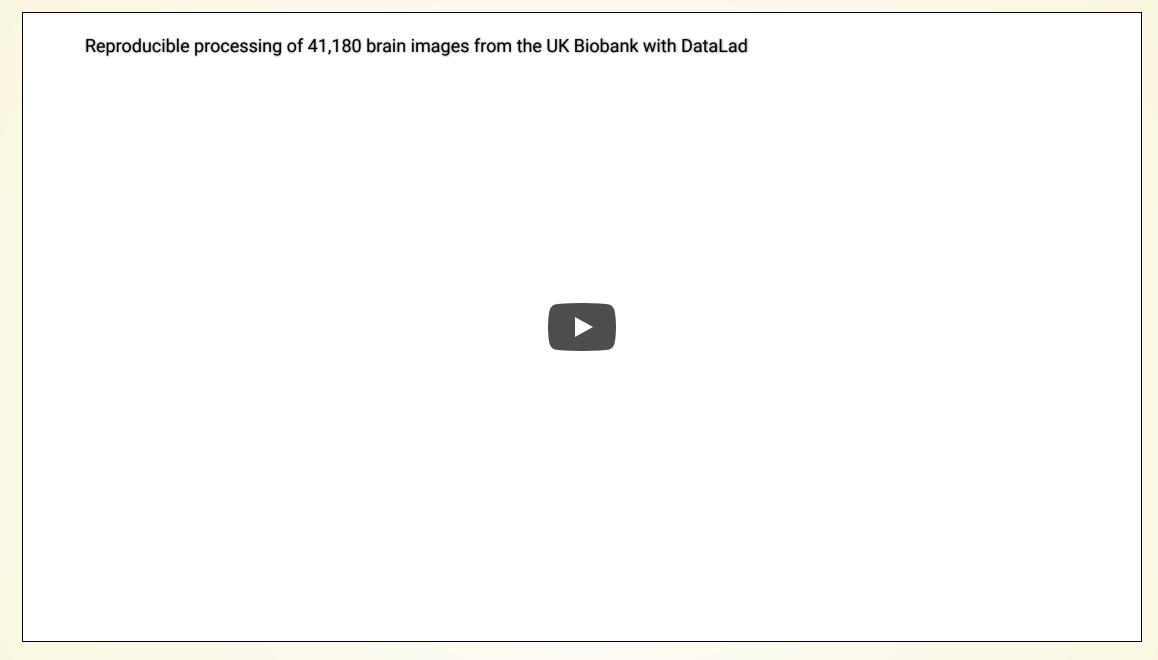


#### **Provenance**

- Every single pipeline execution is tracked
- Execution in ephemeral workspaces ensures results individually reproducible without HPC access

Wagner, Waite, Wierzba et al. (2021). FAIRly big: A framework for computationally reproducible processing of large-scale data.

### FAIRLY BIG MOVIE



- Two computations on clusters of different scale (small cluster, supercomputer).
   Full video: https://youtube.com/datalad
- Two full (re-)computations, programmatically comparable, verifiable, reproducible -- on any system with data access

### TAKE HOME MESSAGES

#### Data deserves version control

It changes and evolves just like code

#### Science, especially on big data, relies on good data management

But effort pays off: Increased transparency, better reproducibility, easier accessibility, efficiency through automation and collaboration, streamlined procedures for synchronizing and updating your work, ...

#### DataLad can help with some things

Have access to more data than you have disk space

Who needs short-term memory when you can have automatic provenance capture?

Link versioned data to your analysis at no disk-space cost

• • •

### HELP?!

If you have a question, you can reach out for help any time:

#### Reach out to to the DataLad team via

- Matrix (free, decentralized communication app, no app needed). We run a weekly Zoom office hour (Thursday, 4pm Berlin time) from this room as well.
- the development repository on GitHub (github.com/datalad/datalad)

#### Reach out to the user community with

A question on neurostars.org with a datalad tag

#### Find more user tutorials or workshop recordings

- On DataLad's YouTube channel (www.youtube.com/channel/datalad)
- In the DataLad Handbook (handbook.datalad.org)
- In the DataLad RDM course (psychoinformatics-de.github.io/rdm-course)
- In the Official API documentation (docs.datalad.org)

### **ACKNOWLEDGEMENTS**

#### Software

- Joey Hess (git-annex)
- The DataLad team & contributors

#### Illustrations

The Turing Way project & Scriberia



#### Science

- Psychoinformatics Lab & INM-7
- Countless open scientists

#### **Funders**







#### **Collaborators**













# LET'S CLEAN UP

- Removing files from a version control system can be unintuitive and harder than expected
- Let's clean up!

datalad drop removes annexed file contents from a local dataset annex and frees up disk space.
 It is the antagonist of get (which can get files and subdatasets).

```
$ datalad drop inputs/sub-02
drop(ok): input/sub-02/func/sub-02_task-oneback_run-01_bold.nii.gz (file)
drop(ok): input/sub-02 (directory)
action summary:
    drop (ok: 2)
```

- But: Default safety checks require that dropped files can be re-obtained to prevent accidental
  data loss. git annex whereis reports all registered locations of a file's content
- drop does not only operate on individual annexed files, but also directories, or globs, and it can uninstall subdatasets:

```
$ datalad drop --what all input
uninstall(ok): input (dataset)
```

datalad remove removes complete dataset or dataset hierarchies and leaves no trace of them. It
is the antagonist to clone.

```
# The command operates outside of the to-be-removed dataset!
$ datalad remove inputs
uninstall(ok): inputs (dataset)
```

 But: Default safety checks require that it could be re-cloned in its most recent version from other places, i.e., that there is a sibling that has all revisions that exist locally datalad siblings reports all registered siblings of a dataset.

 datalad drop refuses to remove annexed file contents if it can't verify that datalad get could reretrieve it

```
$ datalad drop figures/sub-02_mean-epi.png
drop(error): figures/sub-02_mean-epi.png (file) [unsafe; Could only verify the existence of 0 out of 1 necessary
copy; (Use --reckless availability to override this check, or
adjust numcopies.)]
```

Adding --reckless availability overrides this check

```
$ datalad drop figures/sub-02_mean-epi.png --reckless availability
```

 Be mindful that drop will only operate on the most recent version of a file - past versions may still exist afterwards unless you drop them specifically. git annex unused can identify all files that are left behind

datalad remove refuses to remove datasets without an up-to-date sibling

Adding --reckless availability overrides this check

```
$ datalad remove -d my-analysis --reckless availability
```

### **REMOVING WRONGLY**

Removing datasets the wrong way causes chaos and leaves an usuable dataset corpse behind:

```
$ rm -rf local-dataset
rm: cannot remove 'local-dataset/.git/annex/objects/Kj/44/MD5E-s42--8f008874ab52d0ff02a5bbd0174ac95e.txt/
MD5E-s42--8f008874ab52d0ff02a5bbd0174ac95e.txt': Permission denied
```

The dataset can't be fixed, but to remove the corpse chmod (change file mode bits) it (i.e., make it writable)

```
$ chmod +w -R local-dataset
$ rm -rf local-dataset
```