



Nipopy and **Neurobagel**: Practical tools to navigate barriers for FAIR research

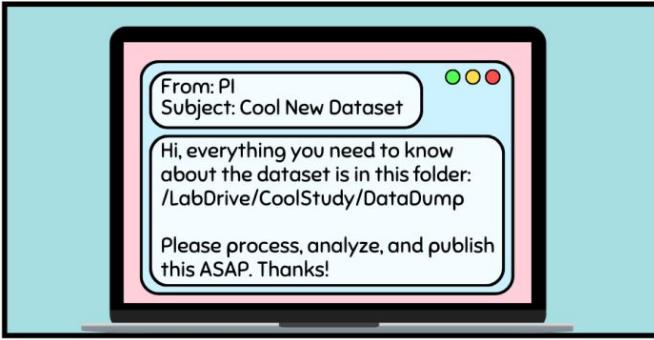
Michelle Wang
PhD candidate, NeuroDataScience-ORIGAMI Lab (JB Poline)

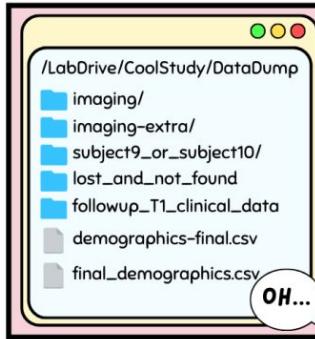
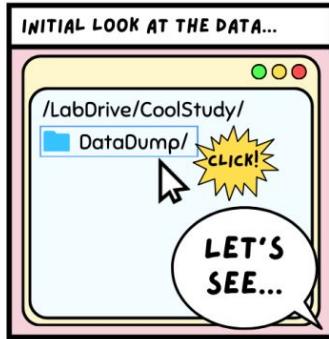
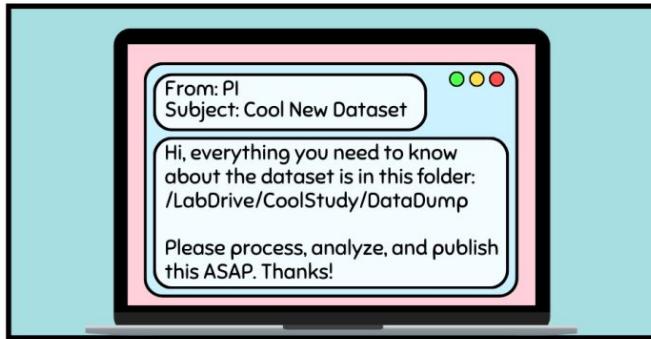
2025 June 18





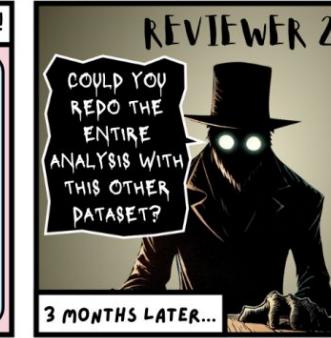
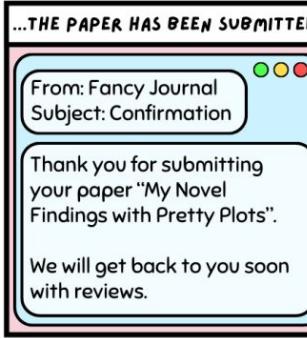
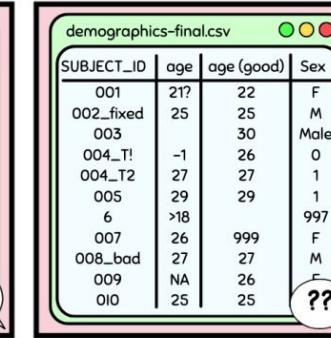
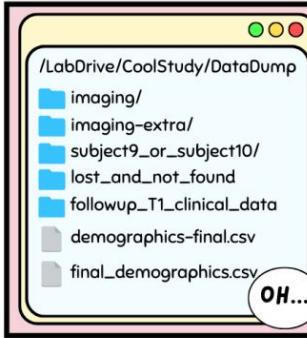
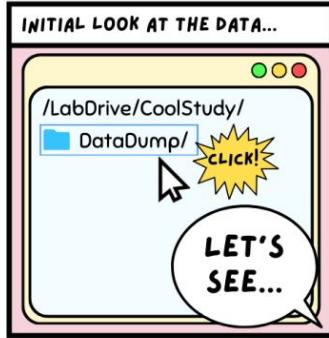
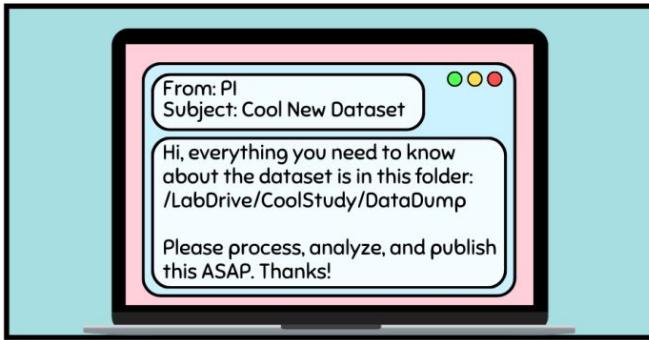
FIRST WEEK IN A NEW LAB...





demographics-final.csv				
SUBJECT_ID	age	age (good)	Sex	
001	21?	22	F	
002_fixed	25	25	M	
003		30	Male	
004_T!	-1	26	O	
004_T2	27	27	1	
005	29	29	1	
6	>18		997	
007	26	999	F	
008_bad	27	27	M	
009	NA	26	F	
010	25	25		??





The ORIGAMI Lab

- **Promote FAIR neuro-data-science**
- **Build neuroinformatic workflows**
- **Improve clinical research practices**



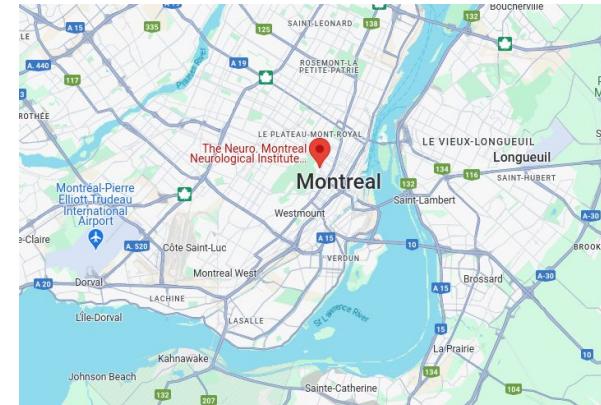
JB Poline



ORIGAMI
Lab



Montreal Neurological
Institute-Hospital





Data challenges in **multicentric** studies

- **Curation:** inevitability of heuristics
- **Processing:** reproducibility and versioning
- **Harmonization:** clinical knowledge
- **Sharing:** privacy, access control



A lot of time-consuming manual work that is difficult to coordinate!



ORIGAMI solutions

Nipopy (curate + process)



A protocol for organizing and processing neuroimaging data

Neurobagel (annotate + search)



Tools for decentralized data harmonization and search



ORIGAMI solutions

Nipopy (curate + process)



A protocol for organizing and processing neuroimaging data

Neurobagel (annotate + search)



Tools for decentralized data harmonization and search



Bagel



ORIGAMI solutions

Nipopy (curate + process)

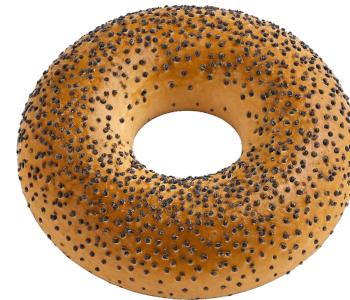


A protocol for organizing and processing neuroimaging data

Neurobagel (annotate + search)



Tools for decentralized data harmonization and search



Poppy seed bagel



Michelle Wang



Nikhil Bhagwat



Nipopy: A framework for the organization and decentralized processing of neuroimaging-clinical studies

Mathieu Dugré



Julia Pfeifer



Missa Dai
Brent McPherson



Jean-Baptiste Poline



<https://nipopy.readthedocs.io/>



<https://github.com/nipopy/nipopy>

Neuroimaging-clinical datasets are **messy**



- **Curating/processing** the data is time-consuming and error-prone
 - Reproducibility concerns
- **Linking** data is difficult



How many participants with

- Parkinson's disease diagnosis
- age under 65
- two imaging sessions
- processed with FreeSurfer

are in my dataset?



How can we make neuroimaging data more FAIR?

- FAIR: findable, accessible, interoperable, reusable (Wilkinson *et al.*, 2016)

Existing open standards/tools developed by the community



(Gorgolewski *et al.*, 2016)



(Kurtzer *et al.*, 2017)



Boutiques

(Glatard *et al.*, 2018)

We leveraged existing open science tools to build a flexible framework for data organization and processing of neuroimaging-clinical data

Introducing the **Nipopy** framework



1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets

Introducing the **Nipopy** framework



1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets

Capture

Take data **as they are**



DICOM



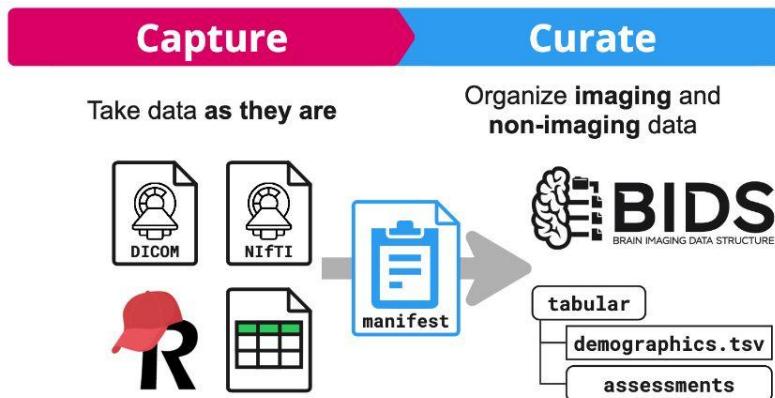
NIFTI



Introducing the **Nipopy** framework



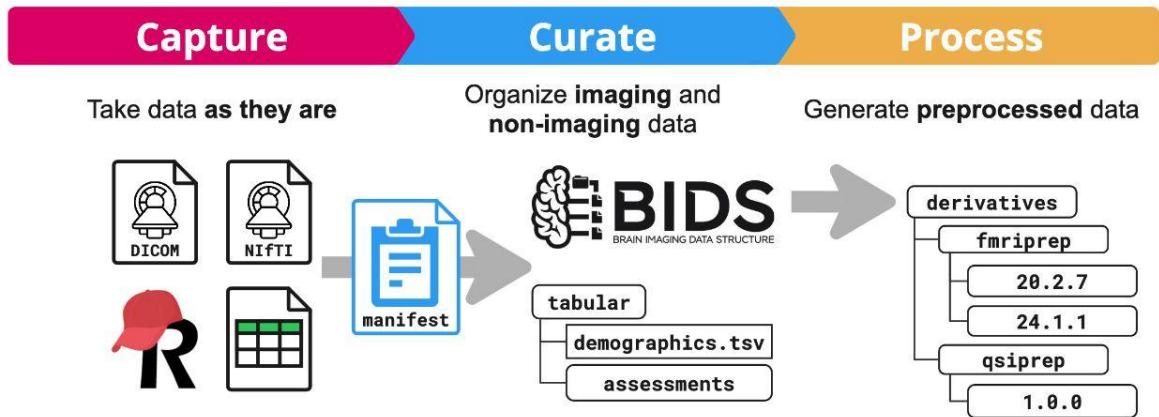
1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets



Introducing the **Nipopy** framework



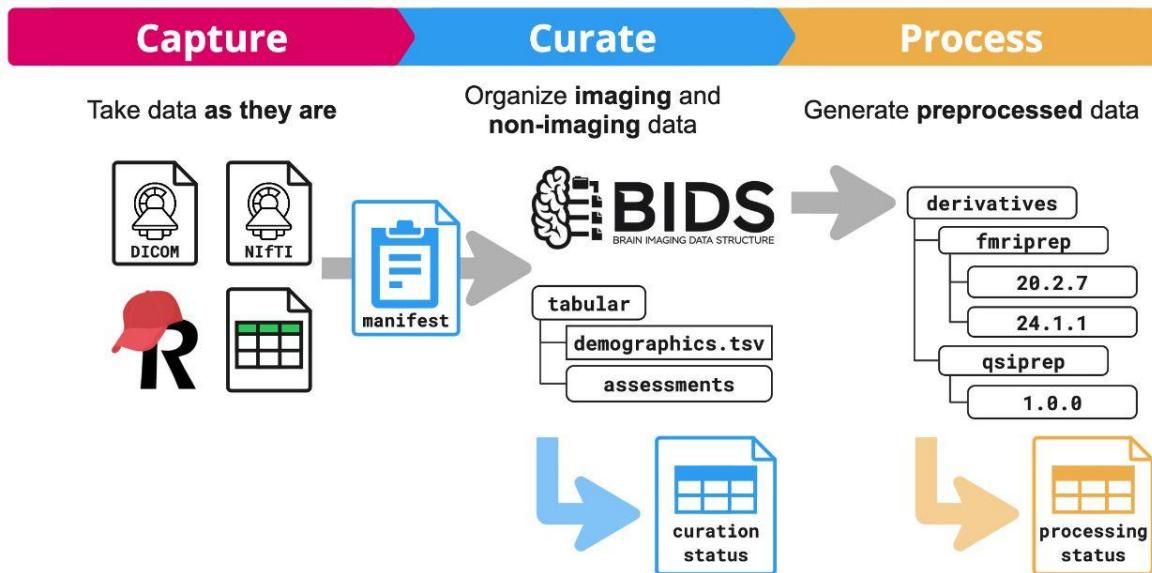
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Introducing the **Nipopy** framework



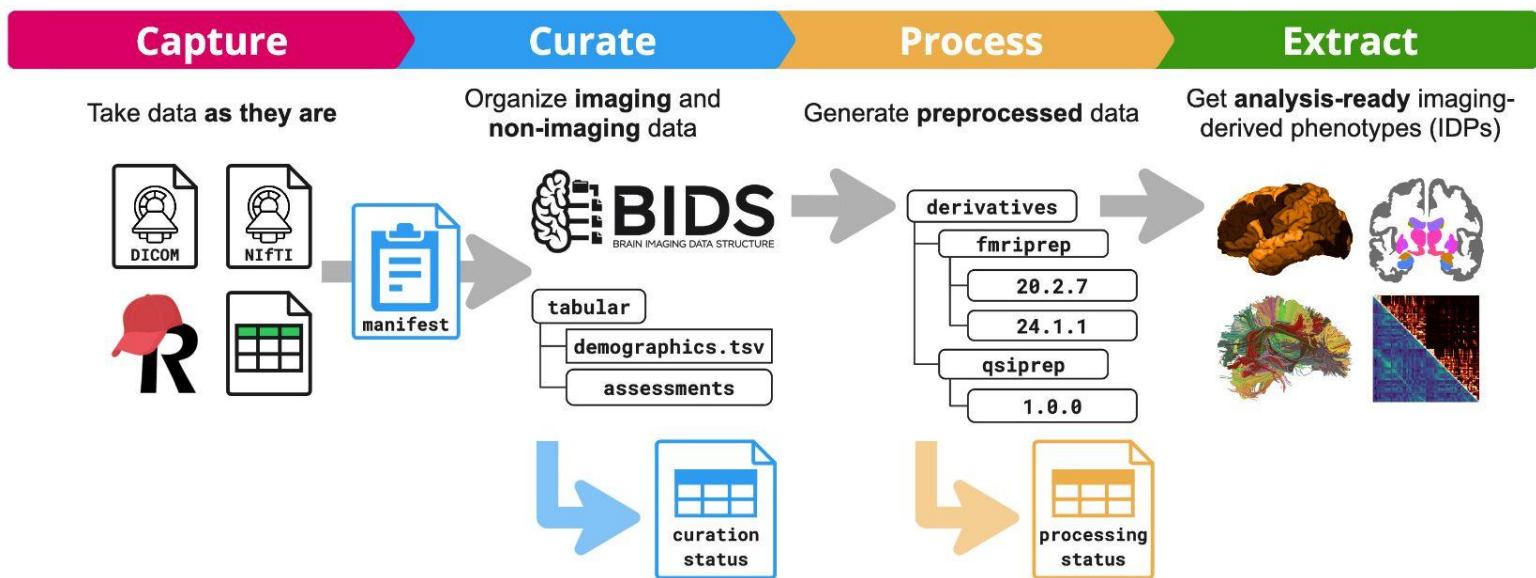
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Introducing the **Nipopy** framework



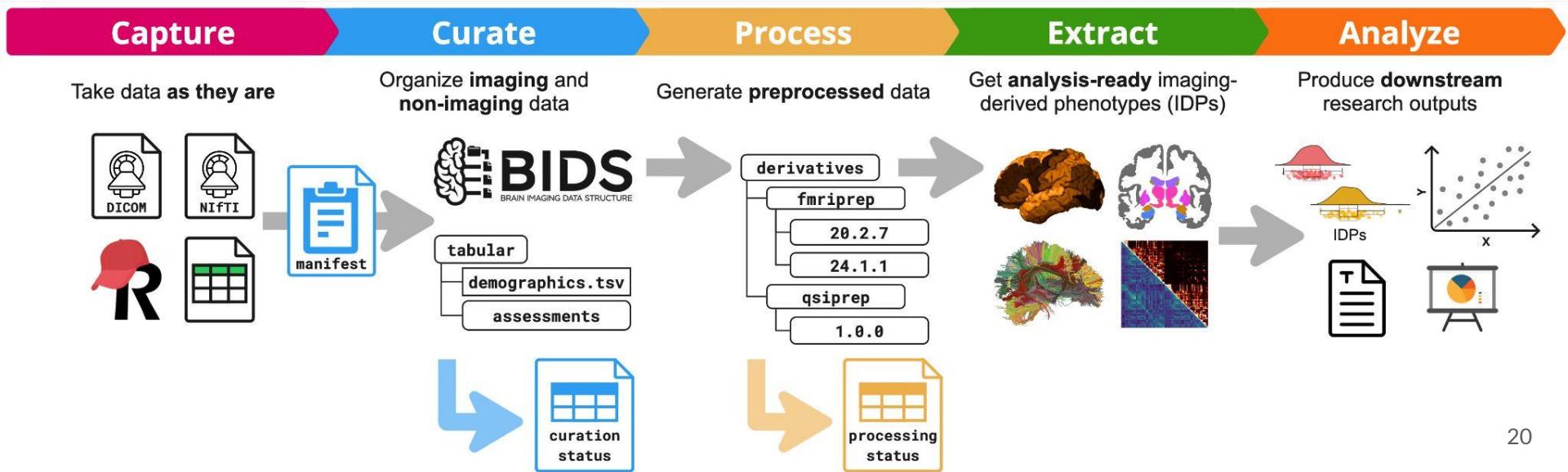
1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets



Introducing the **Nipopy** framework



1. Protocol for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets

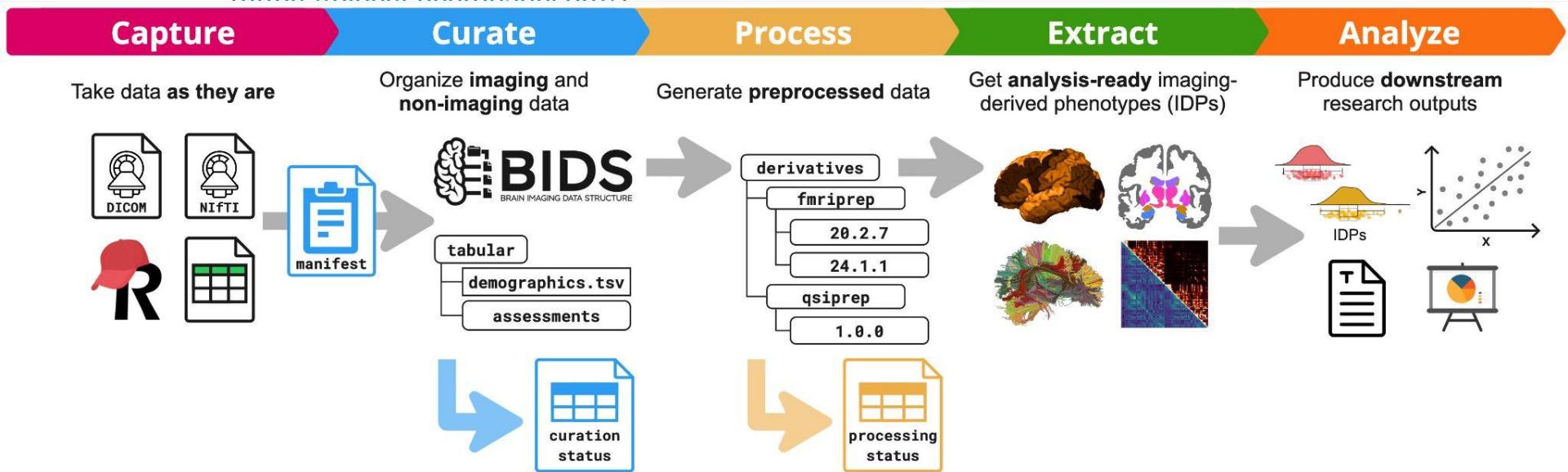


Introducing the **Nipopy** framework



1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets
 - a. From raw scanner data to extracted imaging-derived phenotypes (IDPs)
 - b. Visualizing tabular data/metadata on a dashboard

(<https://dscntt.neurobboal.org/>)



<https://digest.neurobagel.org/>



Neuroimaging and phenotypic dataset exploration beta

Input schema Example input files GitHub

Upload your own digest file:

Select imaging CSV file...

Select phenotypic CSV file...

Load an available digest file:

Available imaging digests ▾ Available phenotypic digests ▾



Introducing the **Nipopy** framework



1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets
 - a. From raw scanner data to extracted imaging-derived phenotypes (IDPs)
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(<https://dicoat.neurobogal.org/>)

Capture

Curate

Process

Extract

Analyze

Introducing the **Nipopy** framework



1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets
 - a. From raw scanner data to extracted imaging-derived phenotypes (IDPs)
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(<https://dicoat.neurobigdata.org/>)

Capture

Curate

Process

Extract

Analyze

2. Data organization **specification**

- a. For imaging and non-imaging data
 - b. At the whole study level

Introducing the **Nipopy** framework



1. **Protocol** for best practices (FAIR principles) in organization/processing of neuroimaging-clinical datasets
 - a. From raw scanner data to extracted imaging-derived phenotypes (IDPs)
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Capture

Curate

Process

Extract

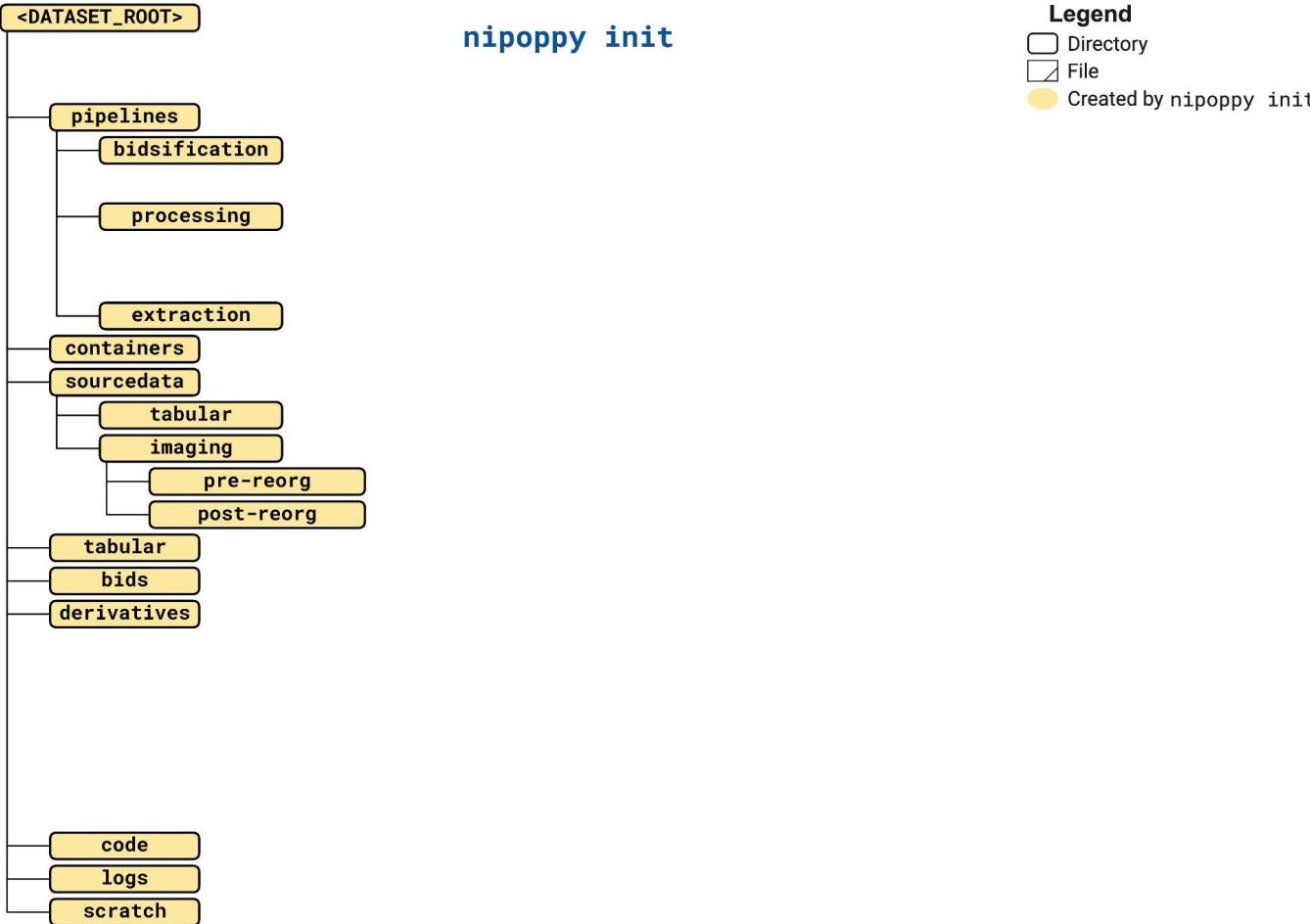
Analyze

2. Data organization **specification**
 - a. For imaging and non-imaging data
 - b. At the whole study level
3. Software package with **tools** to help work with this framework **locally**

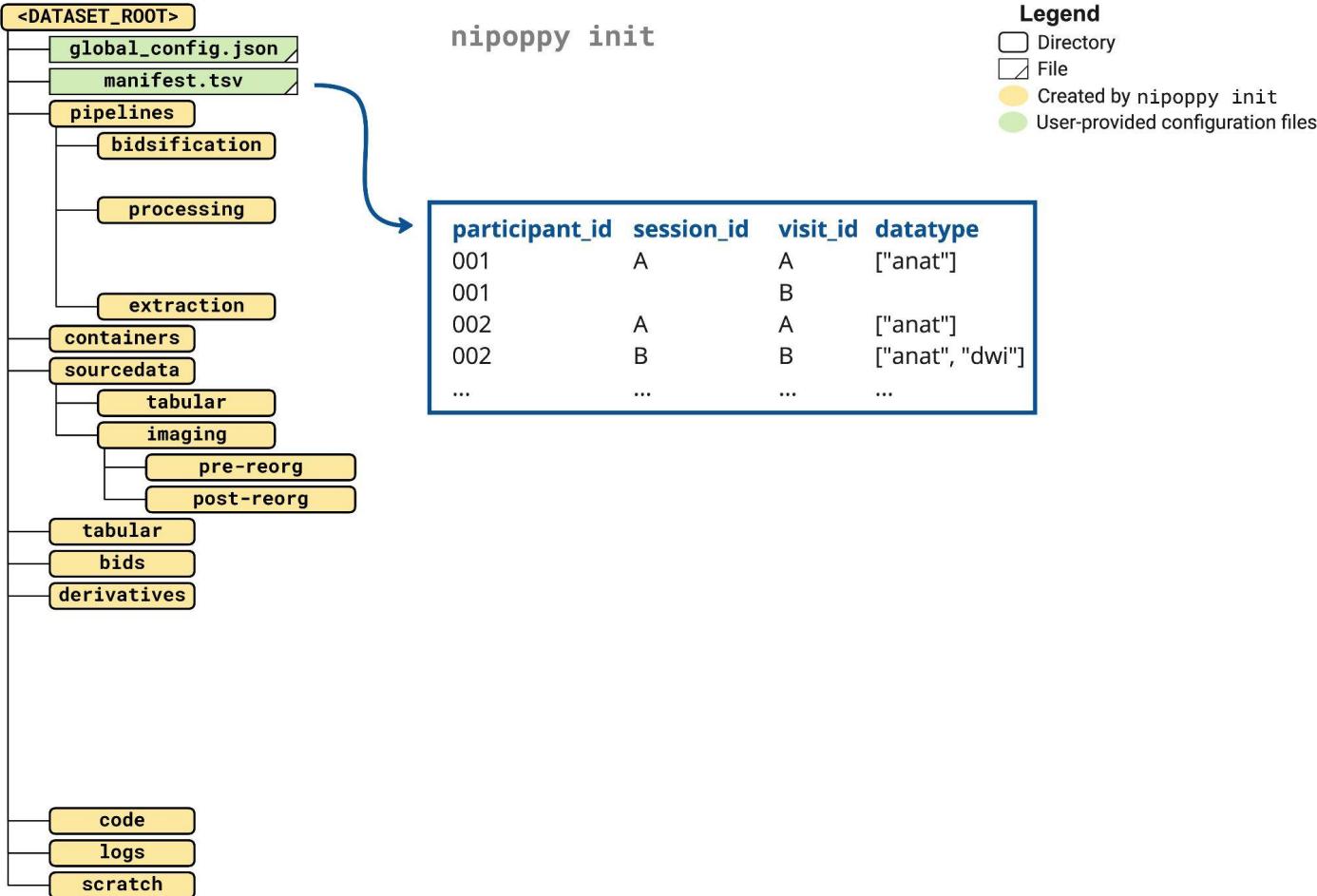
A typical **Nipopy** workflow



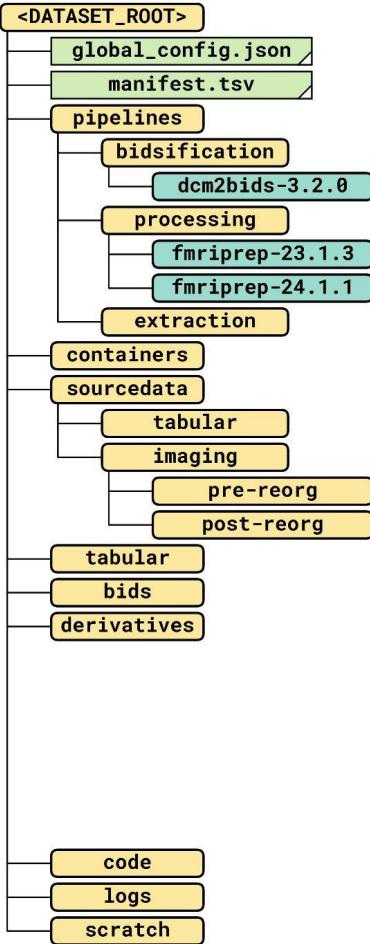
A typical **Nipoppy** workflow



A typical Nipoppyp workflow



A typical Nipoppyp workflow



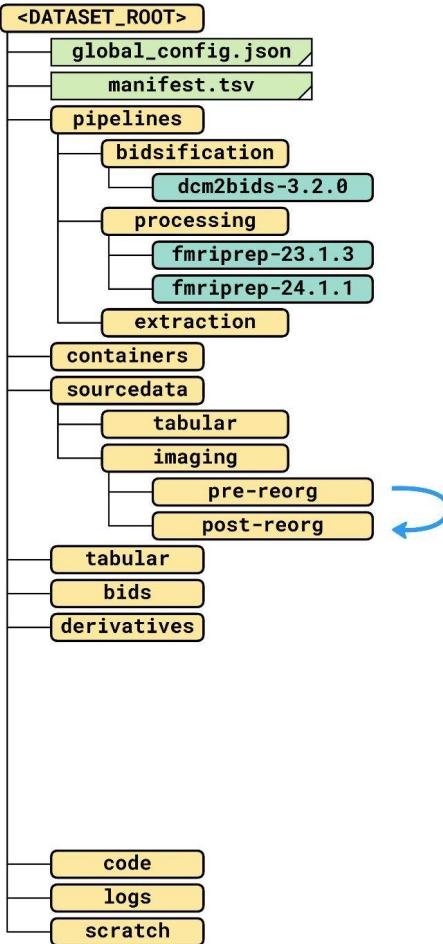
`nipoppyp init`

`nipoppyp pipeline install`

Legend

- Directory
- File
- Created by nipoppyp init
- User-provided configuration files
- Pipeline configuration files

A typical Nipopp y workflow



`nipopp y init`

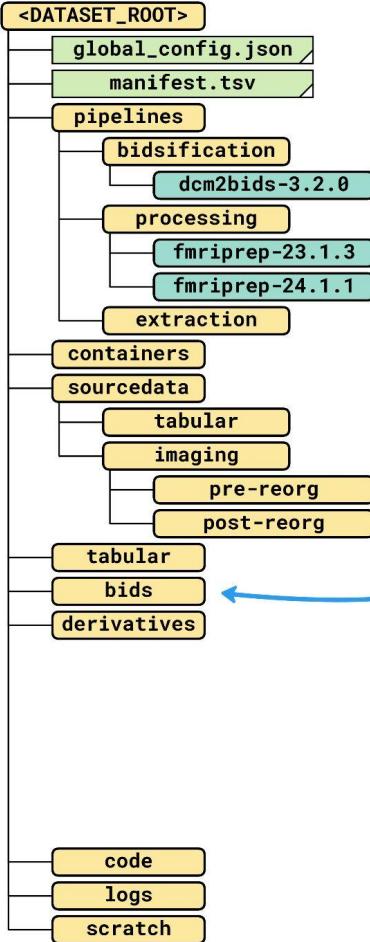
`nipopp y pipeline install`

`nipopp y reorg`

Legend

- Directory
- File
- Created by nipopp y init
- User-provided configuration files
- Pipeline configuration files

A typical Nipoppyp workflow



nipoppyp init

nipoppyp pipeline install

nipoppyp reorg

nipoppyp bidsify

```
{  
    "sourcefolder": "[[NIPOPPYP_DPATH_SOURCEDATA]]",  
    "bidsfolder": "[[NIPOPPYP_DPATH_BIDS]]",  
    "participant_label": [  
        "[[NIPOPPYP_PARTICIPANT_ID]]"  
    ]  
}
```

Legend

Directory

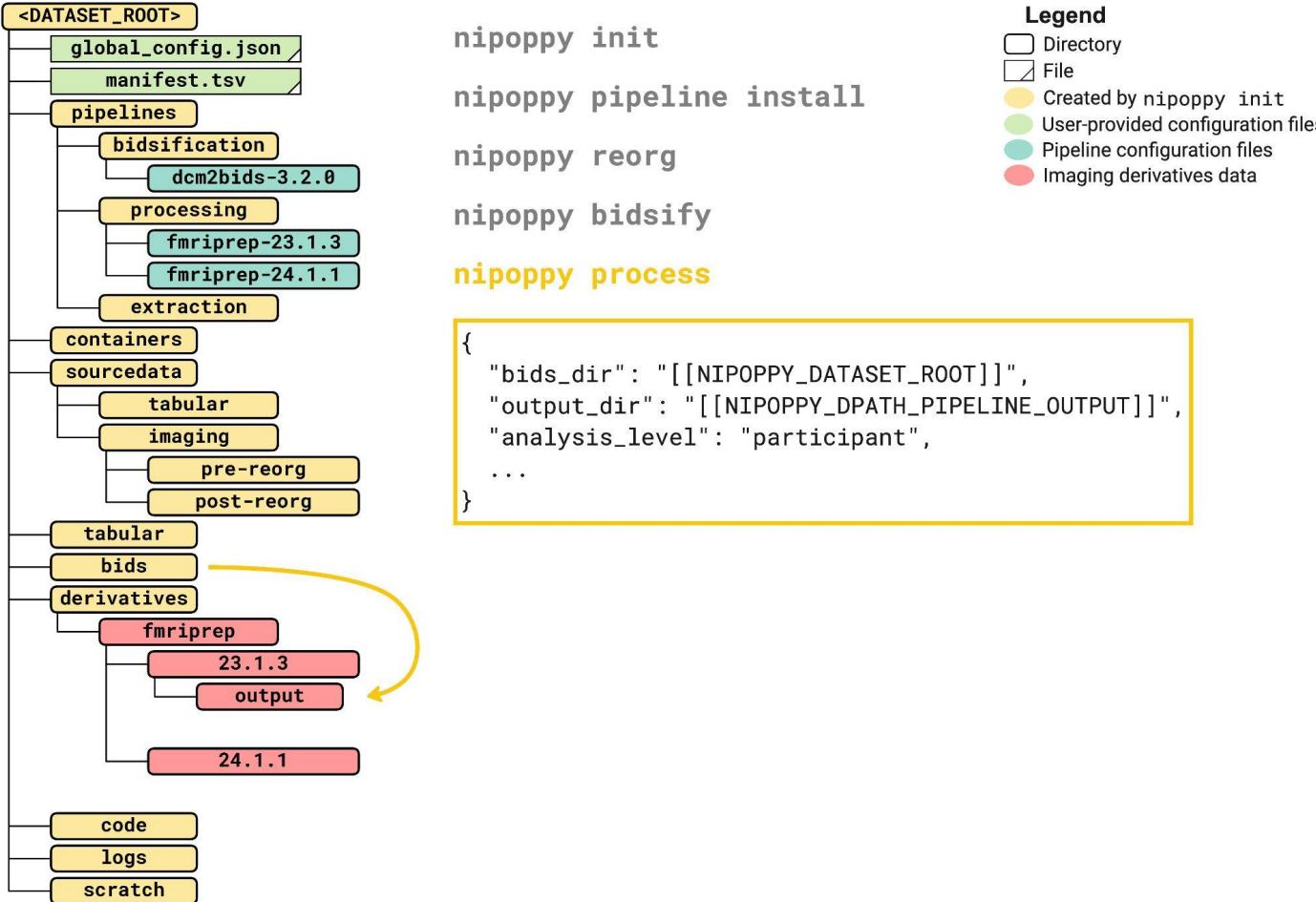
File

Created by nipoppyp init

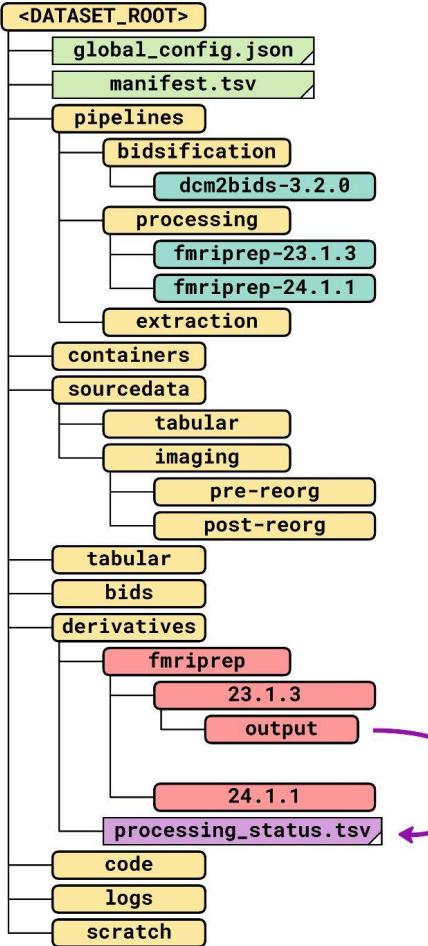
User-provided configuration files

Pipeline configuration files

A typical Nipoppyp workflow



A typical Nipoppyp workflow



nipoppyp init
nipoppyp pipeline install
nipoppyp reorg
nipoppyp bidsify
nipoppyp process
nipoppyp track-processing

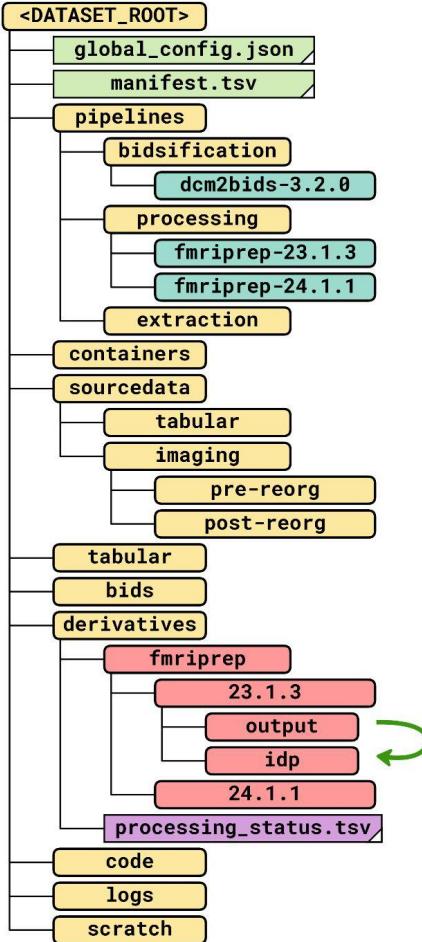
Legend

- Directory
- File
- Created by nipoppyp init
- User-provided configuration files
- Pipeline configuration files
- Imaging derivatives data
- Tracking files

participant_id	session_id	pipeline	version	status
01	A	freesurfer	6.0.0	SUCCESS
01	A	freesurfer	7.4.1	SUCCESS
02	A	freesurfer	6.0.0	FAIL
02	A	freesurfer	7.4.1	FAIL
02	B	freesurfer	6.0.0	INCOMPLETE
02	B	freesurfer	7.4.1	INCOMPLETE
...



A typical Nipoppyp workflow



nipoppyp init
nipoppyp pipeline install
nipoppyp reorg
nipoppyp bidsify
nipoppyp process
nipoppyp track-processing
nipoppyp extract

Legend

- Directory
- File
- Created by nipoppyp init
- User-provided configuration files
- Pipeline configuration files
- Imaging derivatives data
- Tracking files

participant_id	session_id	gm_vol	wm_vol	...
001	A	399445	463167	...
002	A	432054	422803	...
002	B	432913	422634	...
...



Demo

The **Nipopp**y pipeline catalog



Available on the [Zenodo](#) data repository

- Community-driven
- Seamless pipeline configuration sharing (container, runtime parameters)

```
(nipopp) mathdugre@osx ~ % nipopp_demo % py demo.py
```

Documentation and video tutorials



Written documentation available at <https://nipoppy.readthedocs.io/en/latest/>

Coming soon: video tutorials on YouTube

The screenshot shows a YouTube video player interface. At the top left, it says "nipoppy init". Below that, the title "In this video:" is displayed. To the right of the title is a flowchart diagram illustrating the nipoppy workflow: Capture (Take files as they are), Curate (Organize imaging and non-imaging data), Process (Generate preprocessed data), Extract (Get analysis-ready imaging data), and Analyze (Produce downstream reports). A red box highlights the "Capture" step. Below the title and flowchart is a list of topics:

- nipoppy init
- Nipoppy specification
- nipoppy status
- manifest.tsv
- track curation

At the bottom of the video player, there is a progress bar showing "0:53 / 7:42" and a "Scroll for details" button.

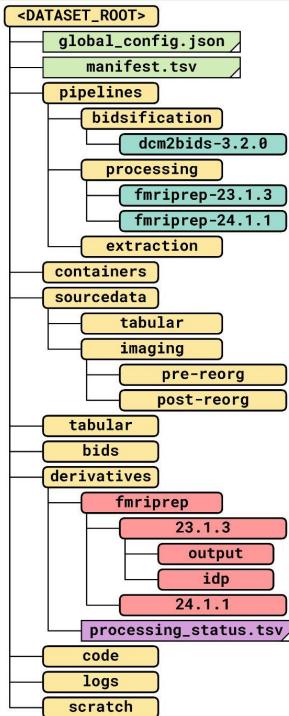


Nipoppyp takeaways

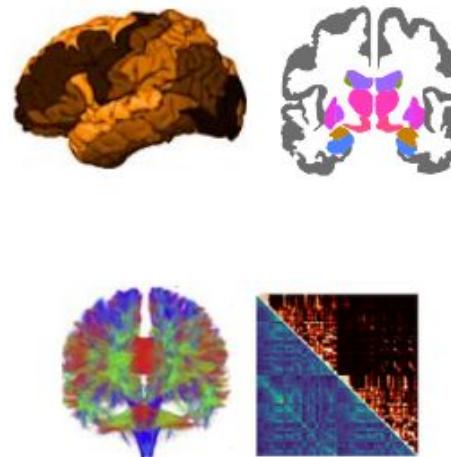
Framework combining existing tools



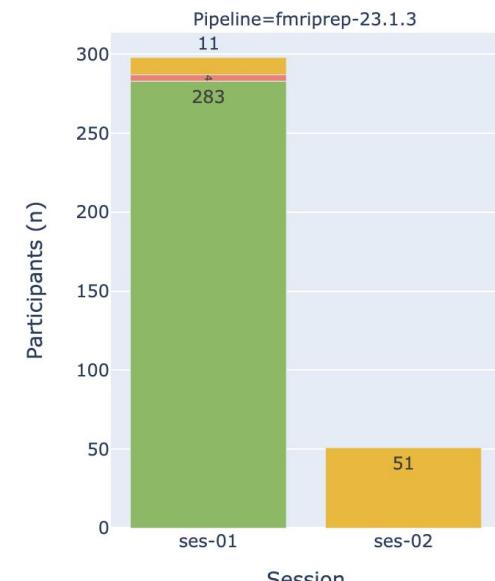
Local data management



Reproducible MRI processing and IDP extraction



Progress tracking



Nipoppy takeaways



For trainees



Best/FAIR practices

For labs



Efficiency and consistency

For institutions



Data-sharing
Metadata discovery
with **Neurobagel** 



Alyssa Dai



Rémi Gau



Arman Jahanpour



Brent McPherson



Huda Bhagat, Baptiste Poline



Sebastian Urchs



Michelle Wang



 neurobagel

 neurobagel.org

Neurobagel © 2025 by Neurobagel Team, Origami Lab, McGill
slides are licensed under CC BY 4.0



ORIGAMI
Lab



A neuroinformatics framework for
decentralized data harmonization and search



HEALTHY BRAINS
HEALTHY LIVES



neuro
Montreal Neurological
Institute-Hospital



CONP
PCNO



National Institutes
of Health



Fondation
Brain Canada
Foundation

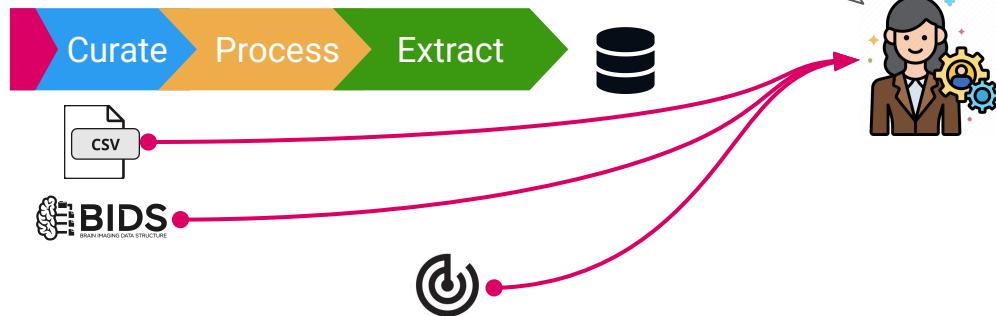




How many participants

- with PD diagnosis
- under 65
- 2 imaging sessions
- run with freesurfer v6

are in **this one dataset?**

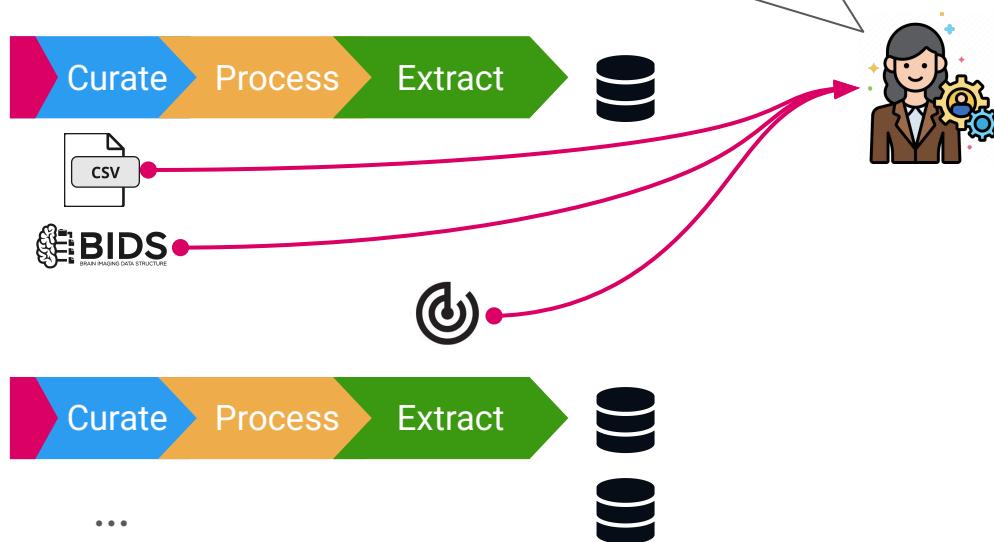




How many participants

- with PD diagnosis
- under 65
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do our PIs have?

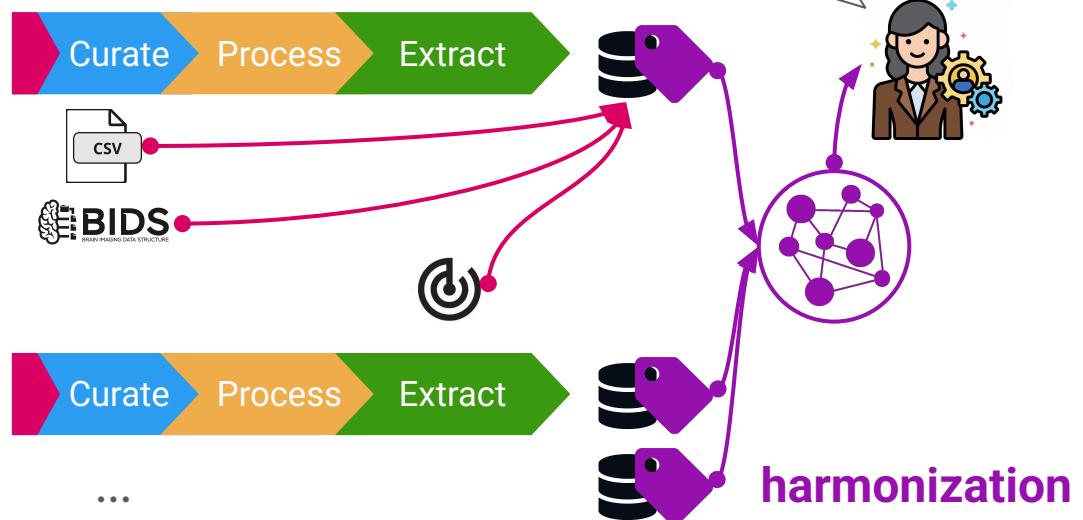




How many participants

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do our PIs have?

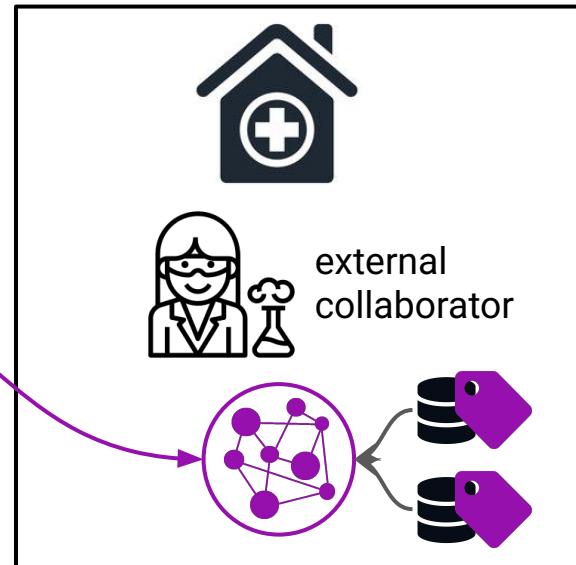
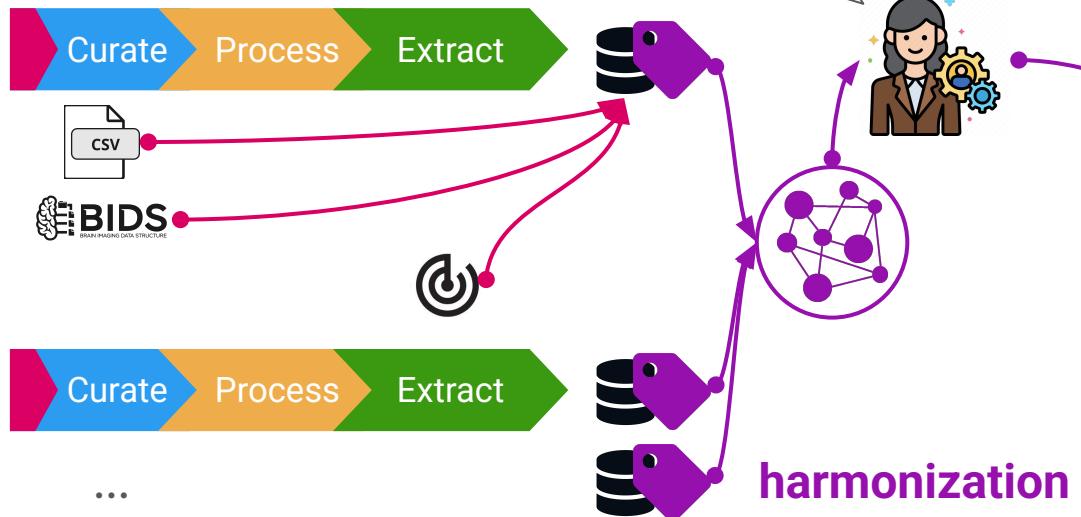




How many participants

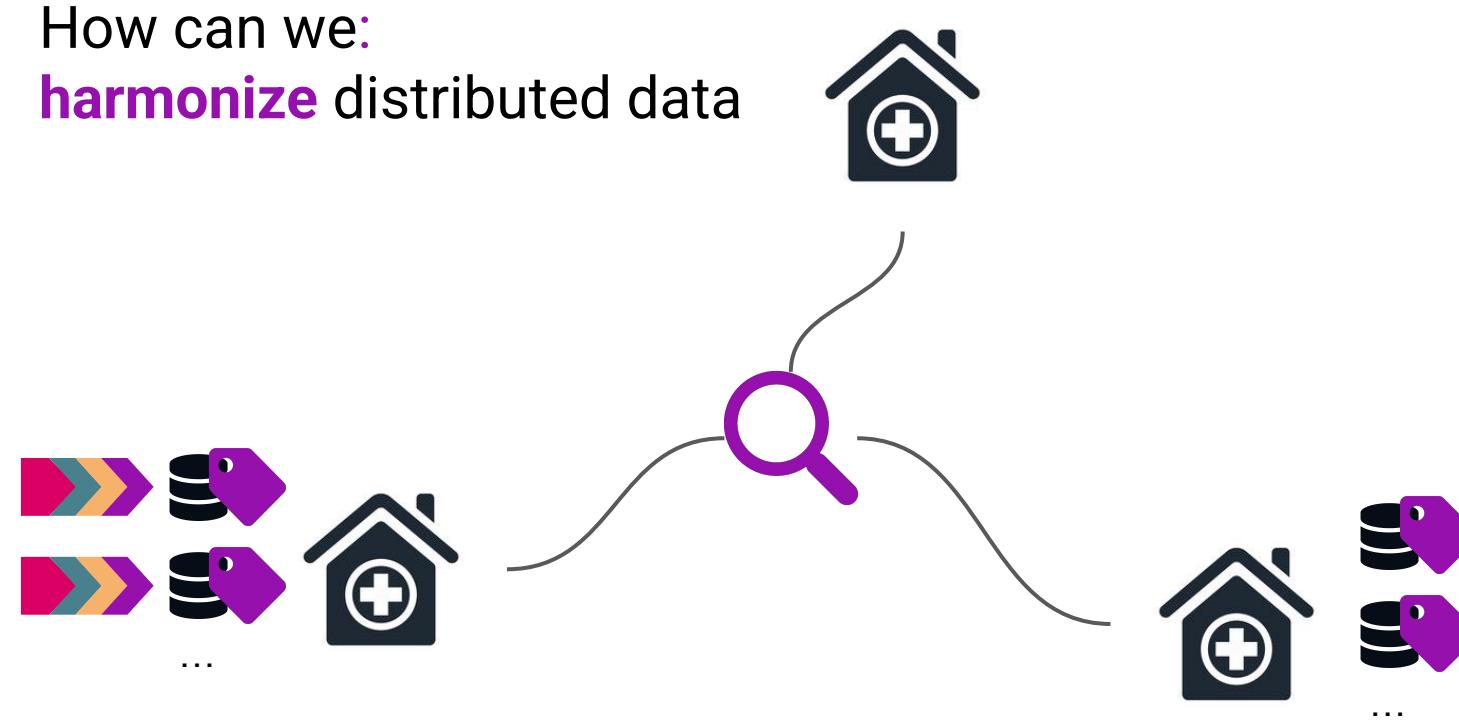
- with PD diagnosis
- under 65
- 2 imaging sessions
- run with freesurfer v6

exist within our network?





How can we:
harmonize distributed data



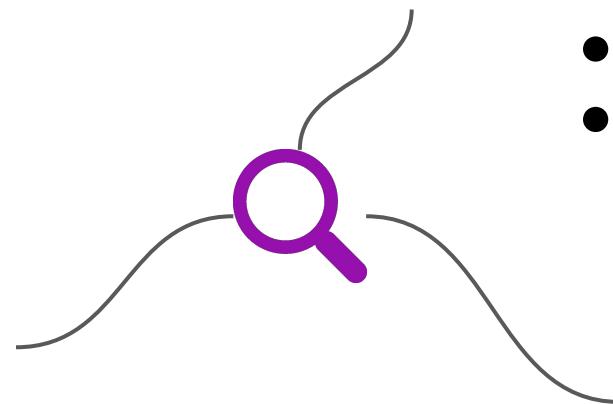


How can we:
harmonize distributed data



Age
Sex
Diagnosis
...

- use common **data model**
- distribute **intuitive GUI**



SNOMED CT
The global language of healthcare



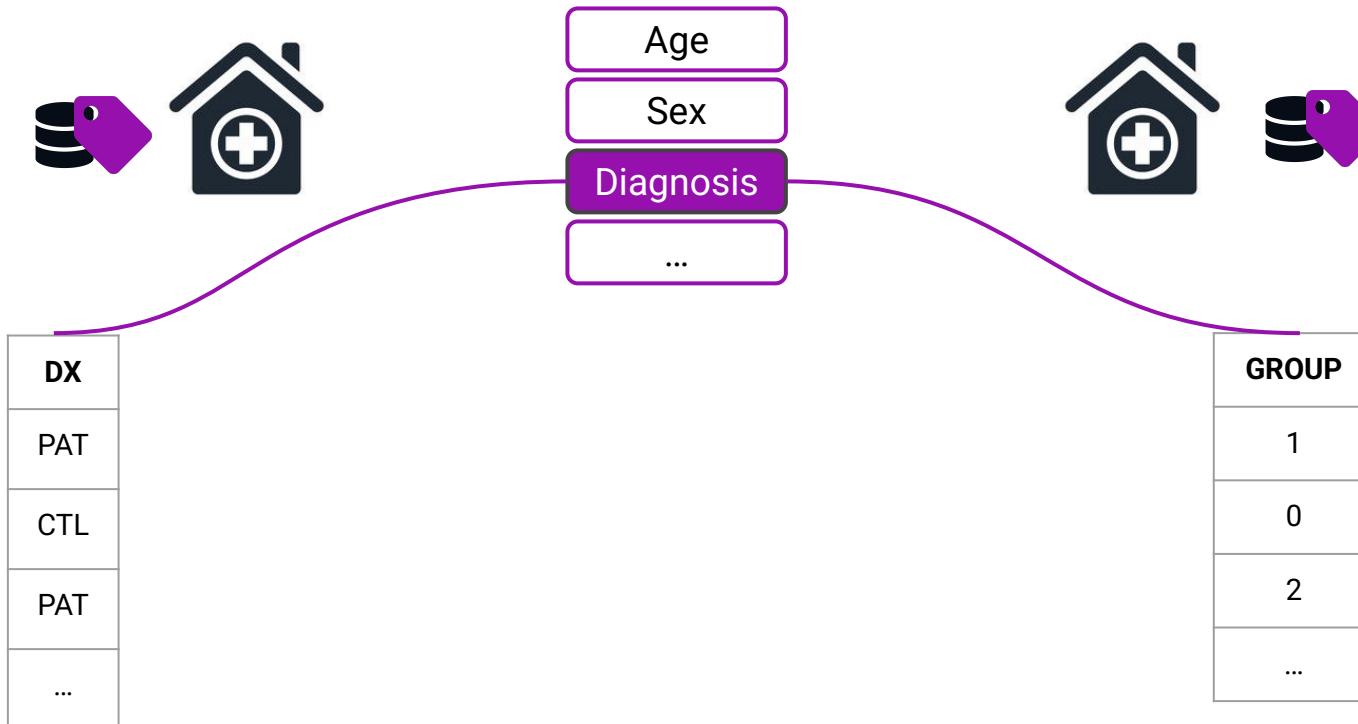
W3C
Word Wide Web Consortium



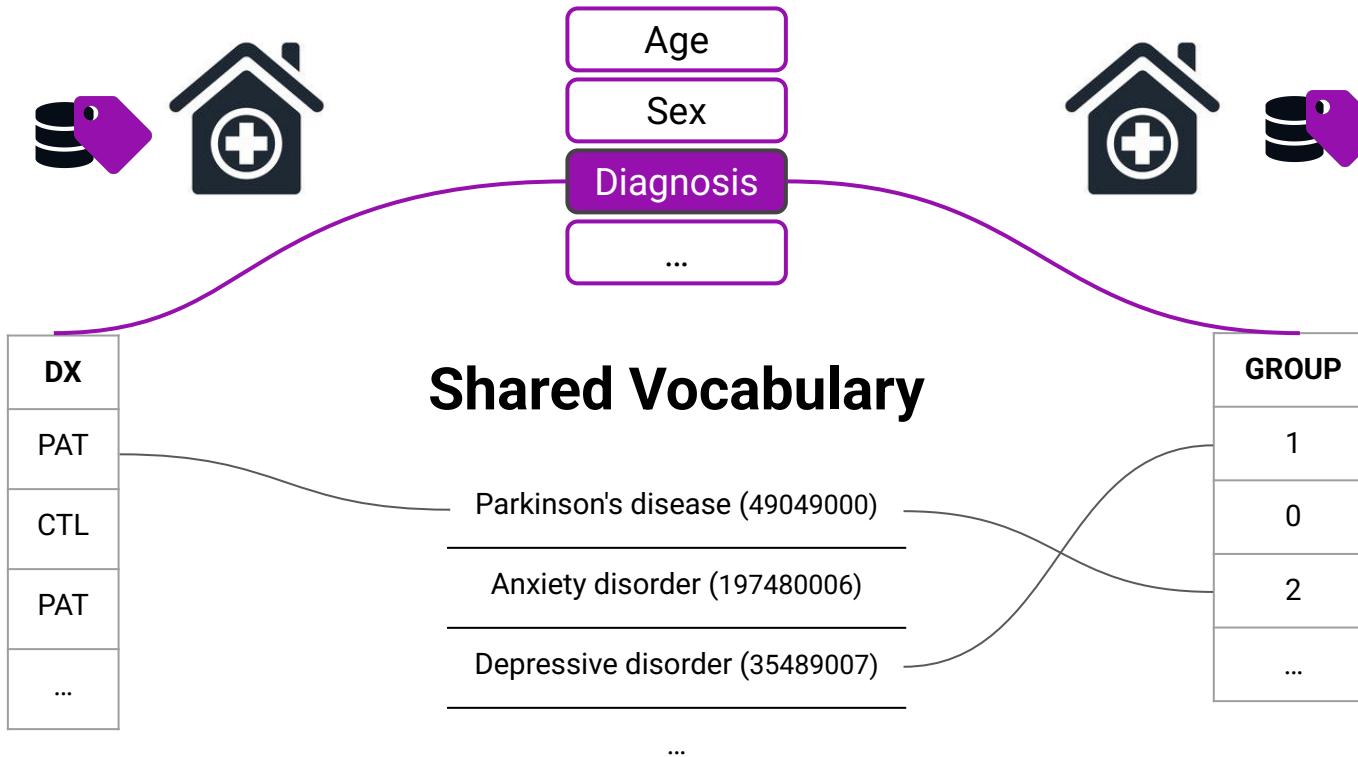
BIDS
BRAIN IMAGING DATA STRUCTURE

Findable
A
ccessible
I
nteroperable
R
eusable

Common participant level data model



Common participant level data model



Intuitive, web-based annotation tool



← → 🔍 🔒 https://annotate.neurobagel.org

 **Neurobagel Annotate** beta
Harmonize phenotypic data

home categorization annotation download | v0.2.0 Documentation

Welcome to the neurobagel annotation tool

This tool allows you to create a machine readable data dictionary in .json format for a tabular phenotypic file in .tsv format ("Data table"). If you already have a [BIDS-like .json data dictionary](#), you can provide it here (under "Data Dictionary") and augment it with more detailed annotations.

 This tool (like all of neurobagel) is under active development and we welcome any feedback or suggestions. Feel free to use our feedback widget on the side of the page.

[Summary of annotation steps](#)

Data table

example_synthetic.tsv

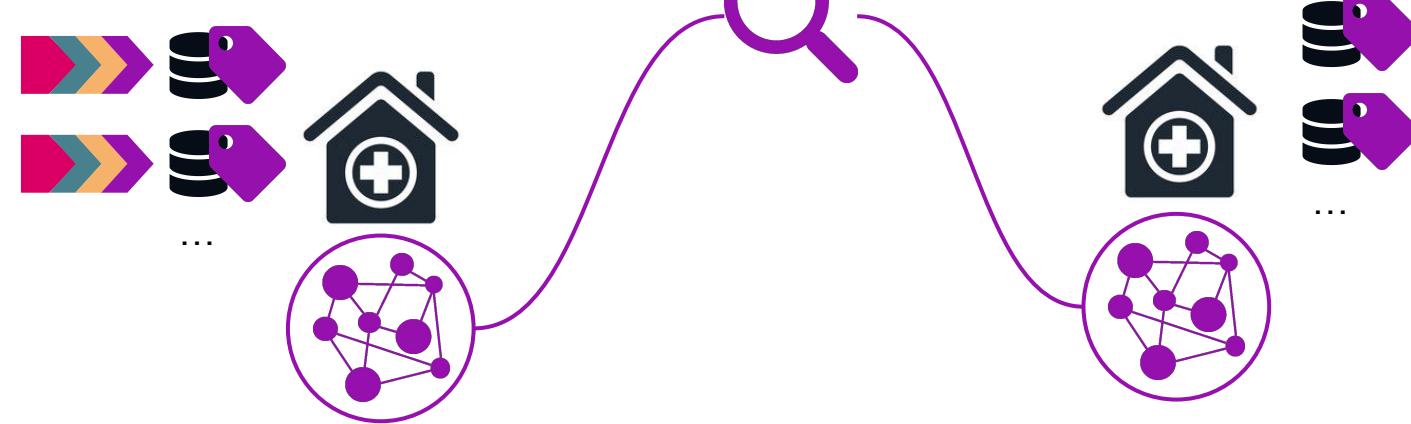
Participant Id	Session Id	Pheno Age	Pheno Sex	Pheno Group	Tool1 Item1	Tool1 Item2	Tool2 Item1
sub-01	ses-01	34,1	F	CTRL	good	far	hello
sub-01	ses-02	35,3	F	CTRL	bad	near	world
sub-02	ses-01	NA	M	PAT	ok	missing	hello

 We like Feedback

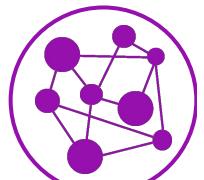
<https://annotate.neurobagel.org/>



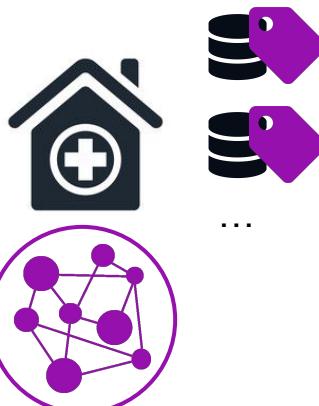
How can we:
search distributed data



How can we:
search distributed data



- **create query federation**
- **intuitive search interface**



Web-based federated query tool



← → ⌂ 🔒 https://query.neurobagel.org/?node>All

Neurobagel Query beta

Define and find cohorts at the subject level

v0.1.0 Documentation 🌐

Query fields

Neurobagel Graph

All

Phenotypic fields (at baseline)

Min Age Max Age

Sex

All

Diagnosis Healthy control

All

Minimum number of sessions

Assessment tool

Results

Select all datasets

Summary stats: 417 datasets, 24851 subjects

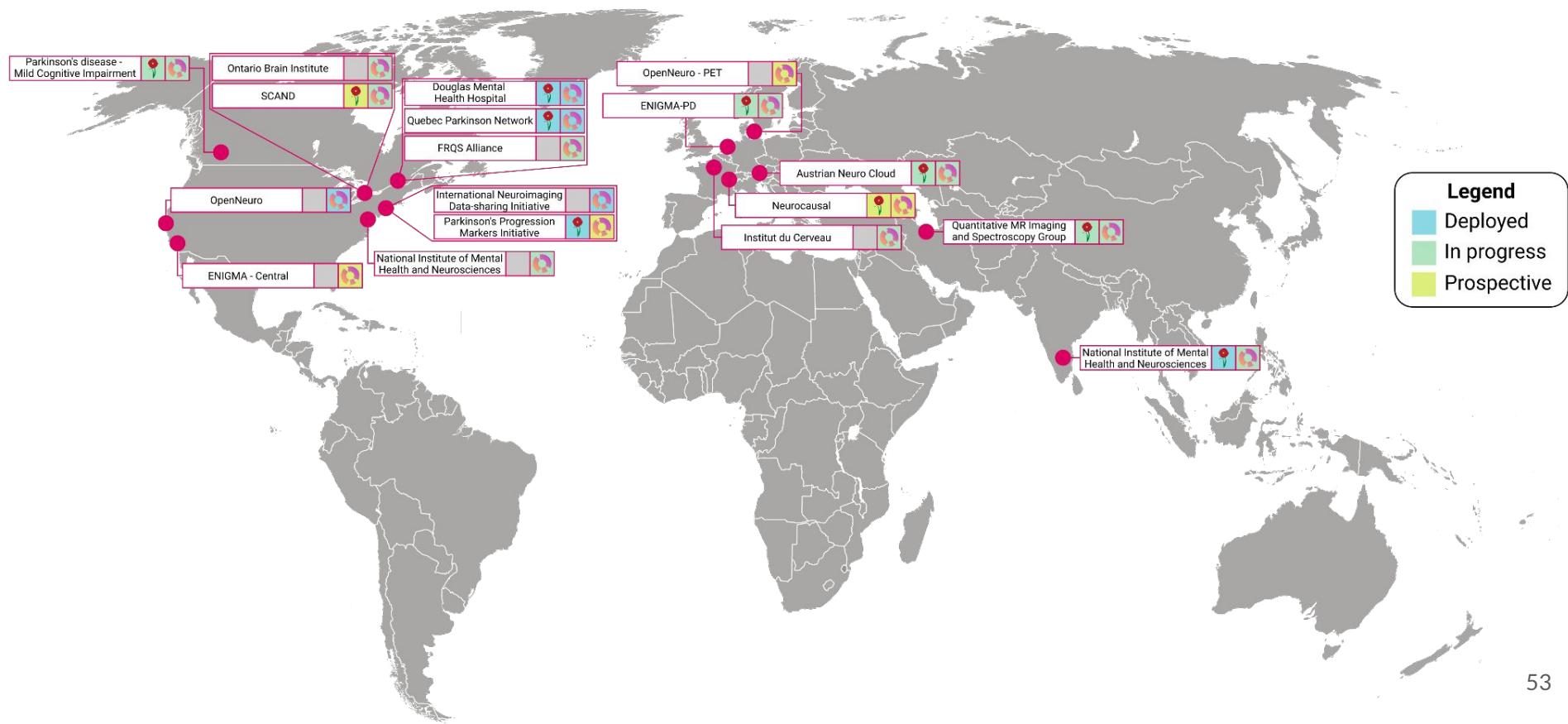
SharedStates from OpenNeuro
 26 subjects / 26 total subjects Flow

A longitudinal neuroimaging dataset on language processing i from OpenNeuro
 ...more DWI
322 subjects / 322 total subjects

MNE-somato-data-bids dataset (anonymized) from OpenNeuro
 1 subjects / 1 total subjects

We like Feedback

Nipoppy and Neurobagel adoption so far





Thank you!



Michelle
Wang



Mathieu
Dugré



Brent
McPherson



Nikhil
Bhagwat



Arman
Jahanpour



Sebastian
Urchs



Alyssa
Dai



Julia
Pfarr



Jean-Baptiste
Poline



ORIGAMI
Lab



neuro
Montreal Neurological
Institute-Hospital

CIHR IRSC
Canadian Institutes of
Health Research
Instituts de recherche
en santé du Canada

Québec
Fonds de recherche – Nature et technologies
Fonds de recherche – Santé
Fonds de recherche – Société et culture

PARKINSON
QUÉBEC

UNIQUE
CENTRE

NIH
National Institutes
of Health

THE MICHAEL J. FOX FOUNDATION
FOR PARKINSON'S RESEARCH



McGill
UNIVERSITY

Chan
Zuckerberg
Initiative

HBHL
NEUROHUB

P
QUEBEC
PARKINSON
NETWORK

Fondation
Brain Canada
Foundation