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22<sup>nd</sup> November 2023

Neuro Open-science Workshop



# Clinica

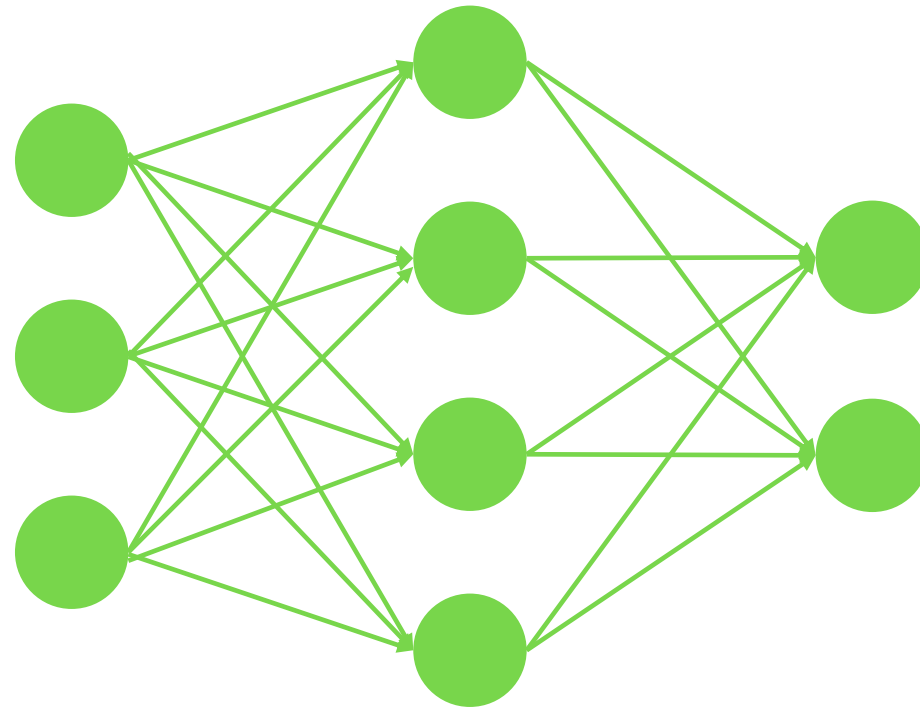
Open-source software platform  
for neuroimaging studies



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Once upon a time in a lab...



## Motivation

- Stop the waste of resources

## Objectives

- Spend less time on data management and processing
- Easily share data and results within institutions and with external collaborators
- Make research more reproducible
- Highlight methods developed in the team



Clinica

[User Documentation](#)

[Paper](#)

[Support](#)

[Development](#) ▾

[Around Clinica](#) ▾

[↓ Installation](#)

# Clinica

## Software platform for clinical neuroimaging studies

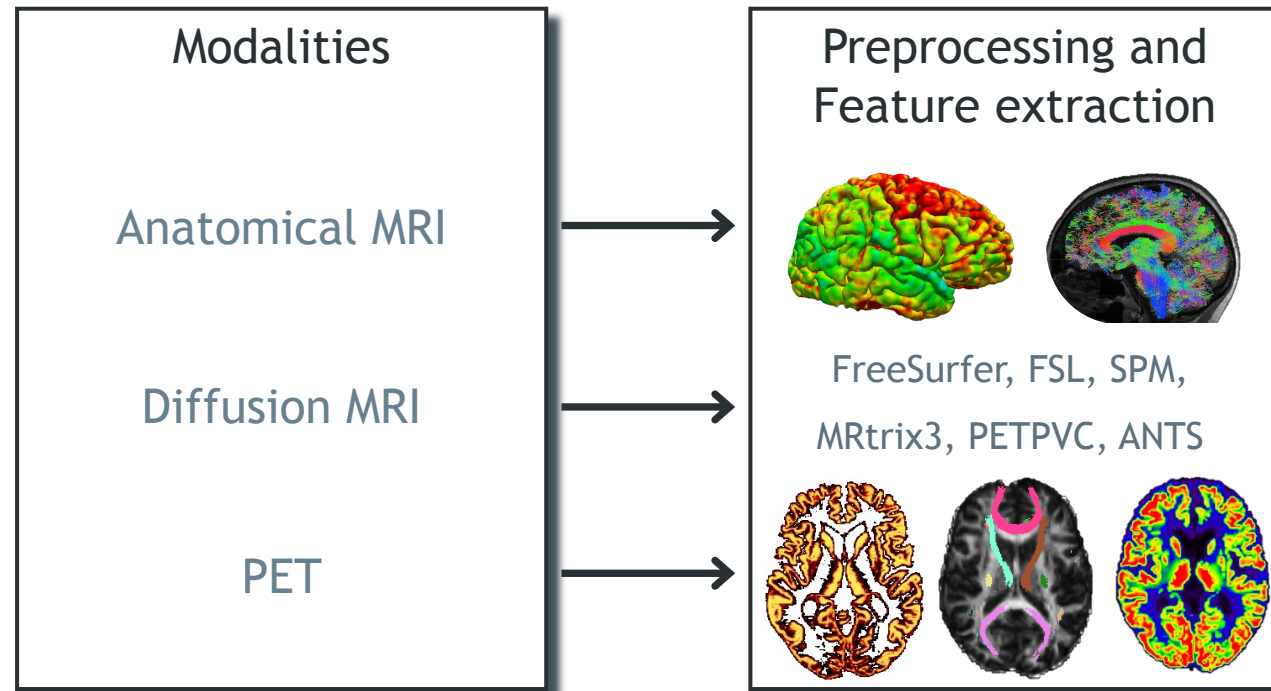
Clinica is a software platform for clinical research studies involving patients with neurological and psychiatric diseases and the acquisition of multimodal data (neuroimaging, clinical and cognitive evaluations, genetics...), most often with longitudinal follow-up.

The development of Clinica was initiated by the [ARAMIS Lab](#) at the [Paris Brain Institute](#).

[QUICK START](#)



## Software platform for neuroimaging studies



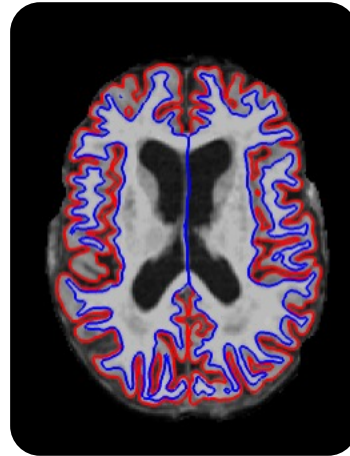




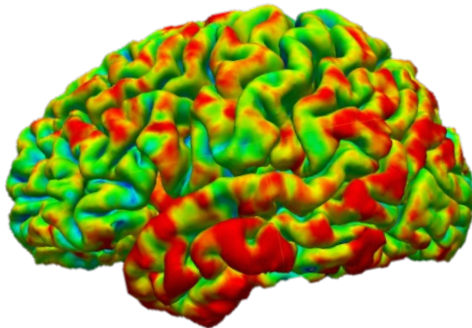
```
→ clinica run t1-freesurfer BIDS_Dataset CAPS_Dataset
```



Surface  
extraction



Cortical thickness  
estimation

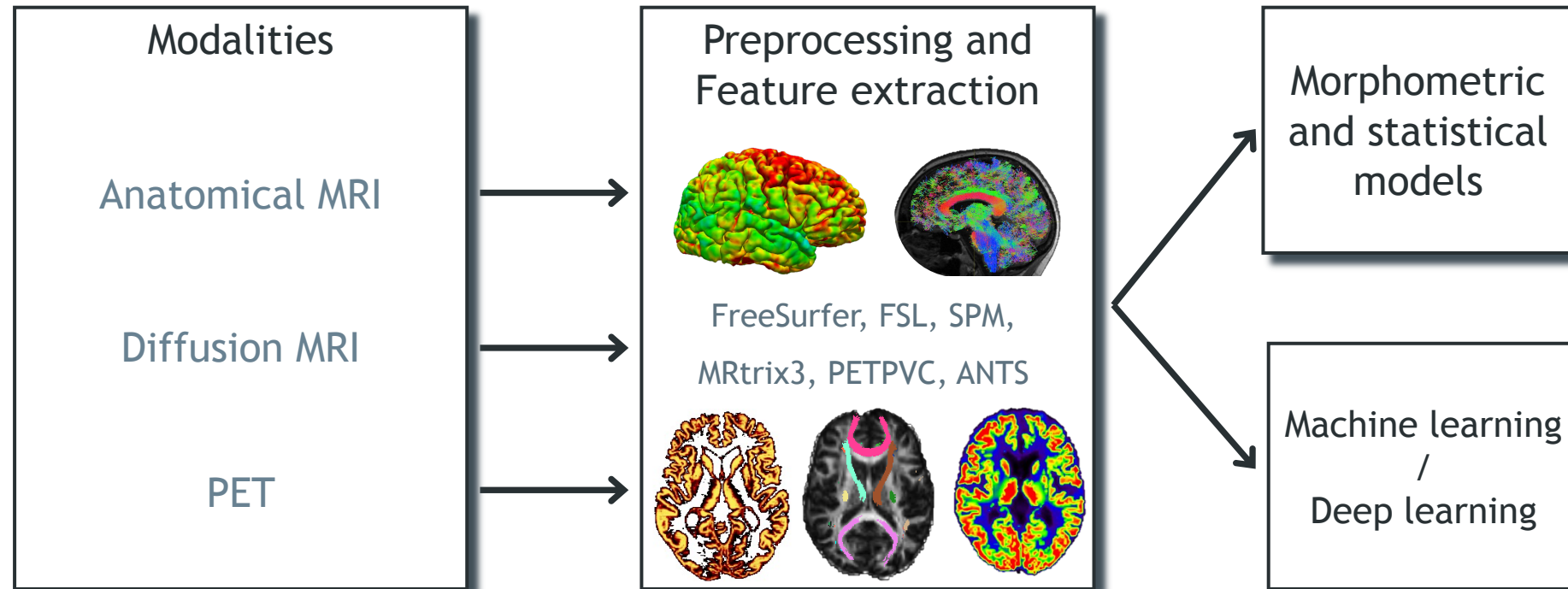


## FreeSurfer



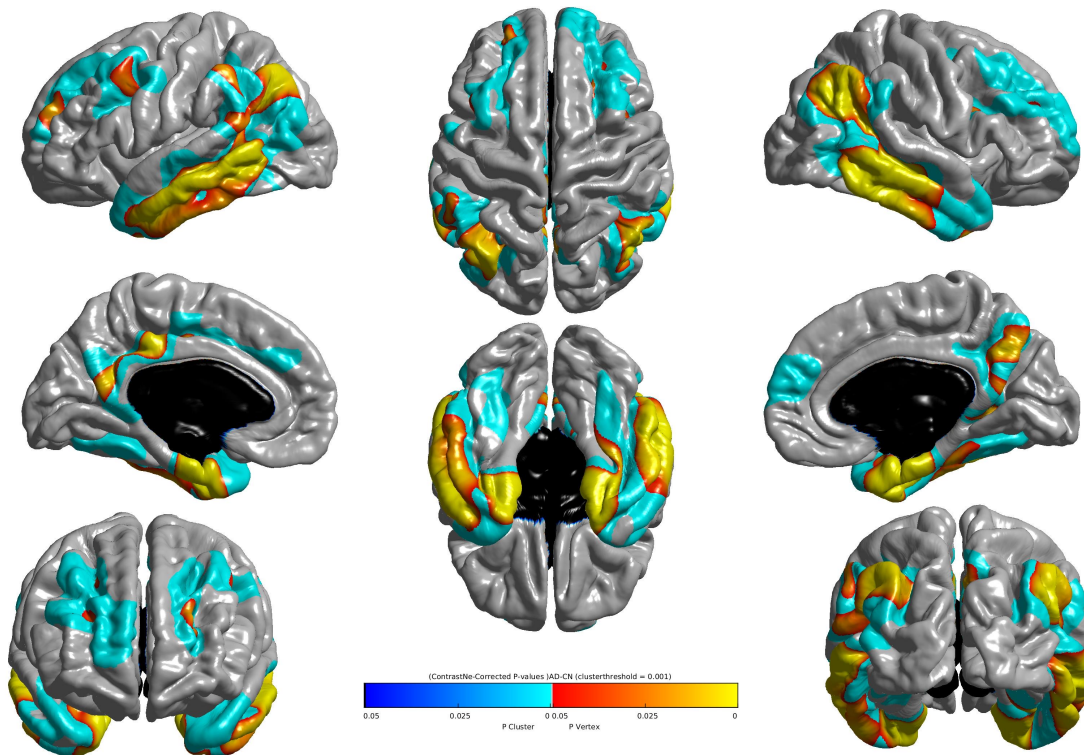
<https://surfer.nmr.mgh.harvard.edu>

# Software platform for neuroimaging studies





```
→ clinica run statistics-surface CAPS_Dataset ADvsHC ADvsHC_participants.tsv
<analysis_parameter>
```



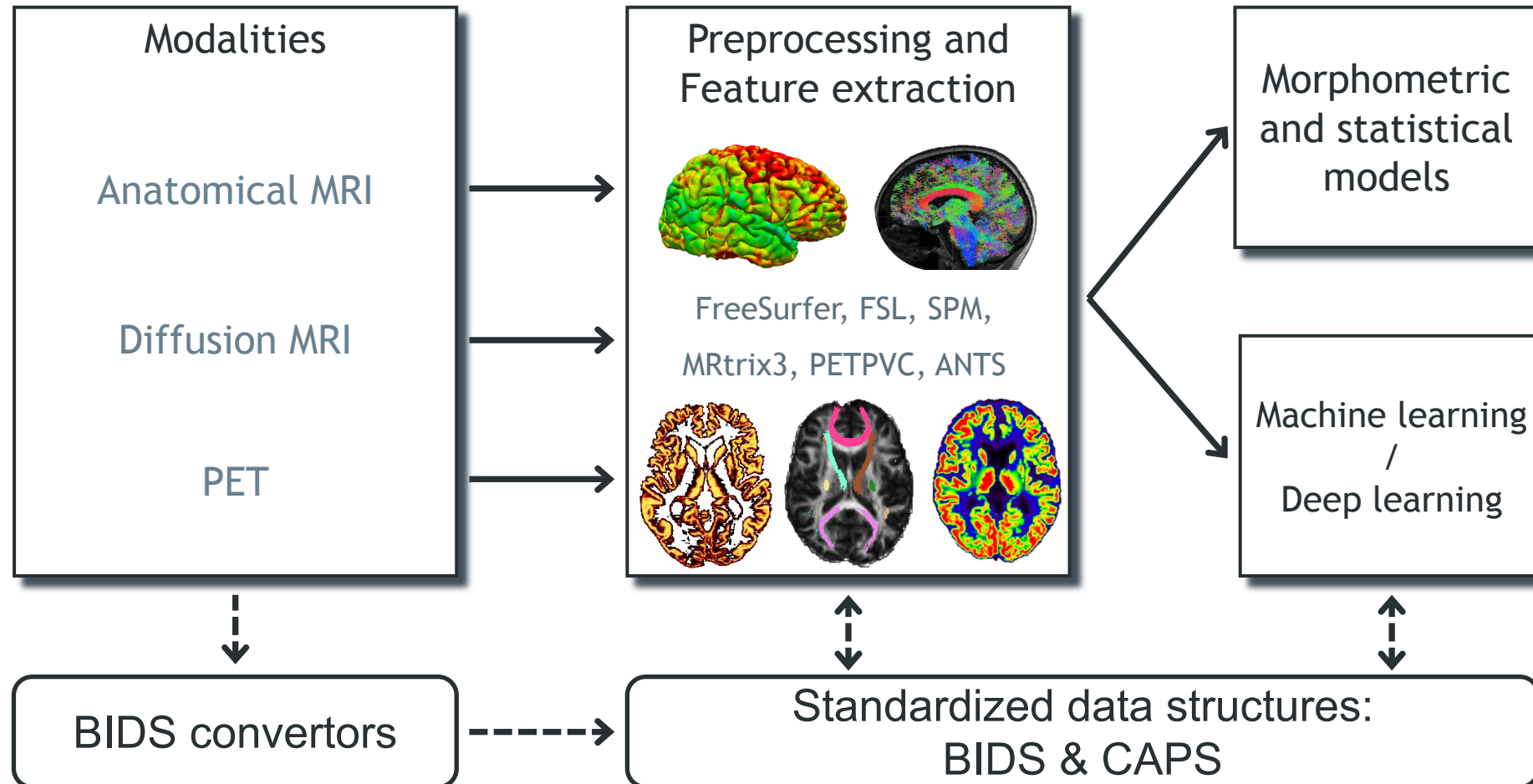
## SurfStat

<https://www.math.mcgill.ca/keith/surfstat>

Areas of significantly reduced cortical thickness in Alzheimer's disease patients compared with healthy controls



# Software platform for neuroimaging studies



```
→ clinica convert adni-to-bids ADNI_unorganized ADNI_BIDS
```

## ADNI\_unorganized

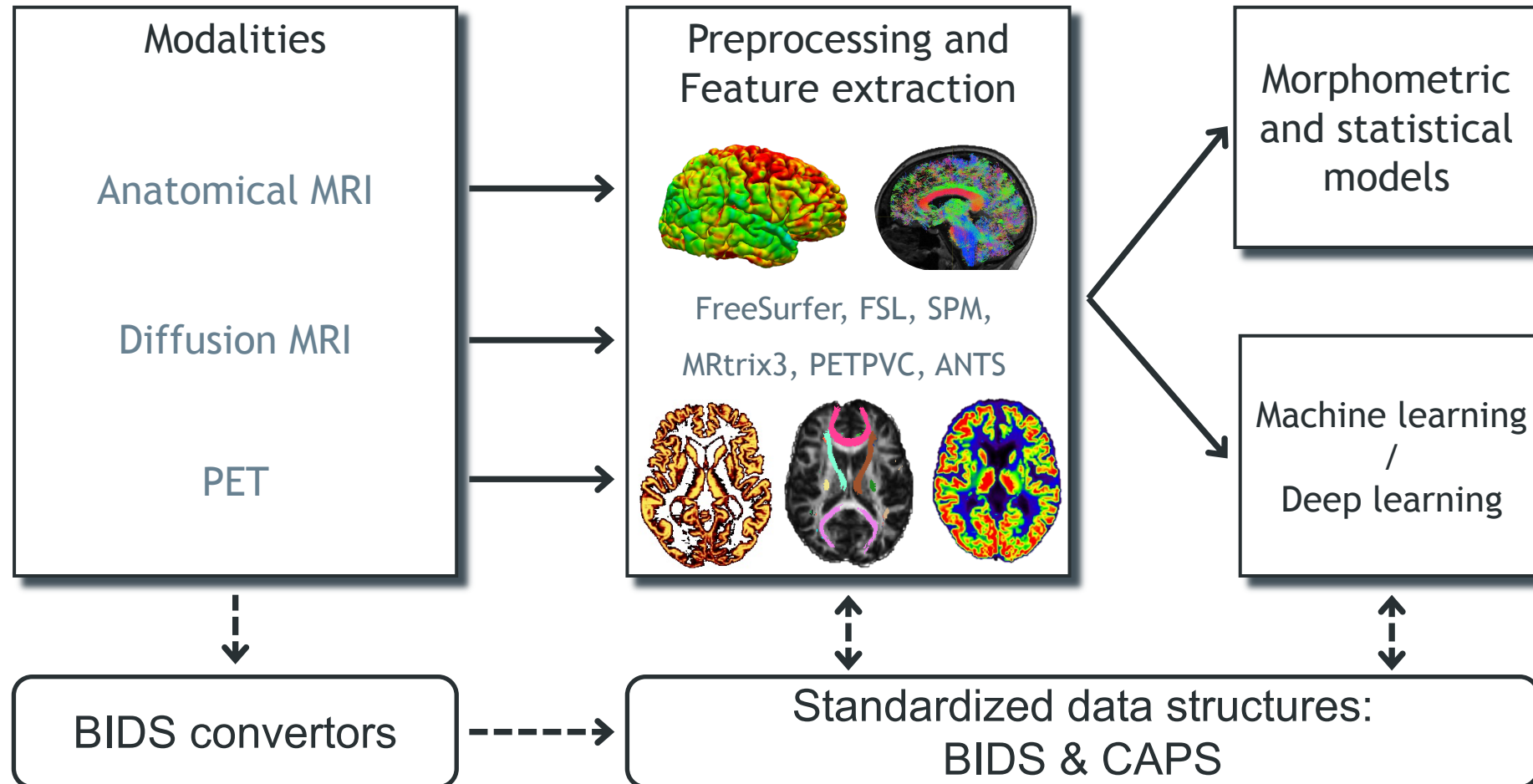
```
├─ 094_S_4089
│  │ ...
│  │ └─ Accelerated_SAG_IR-SPGR
│  │ └─ AV45_Coreg_Avg_Standardized_Image_and_Voxel_Size
│  │ ...
│  │ └─ Axial_DTI
│  │ └─ Axial_FLAIR
│  │ └─ Axial_T2_Star
│  │ └─ Calibration_Scan
│  │ └─ Coreg_Avg_Standardized_Image_and_Voxel_Size
│  │ ...
│  │ └─ MT1_GradWarp_N3m
│  │ └─ Sag_IR-SPGR
│  │   └─ 2011-10-18_12_15_56.0
│  │     └─ S125692
│  │       └─ ADNI_094_S_4089_MR_Sag_IR-SPGR_br_raw_20111019095510271_80_S125692_I261478.dcm
│  │       └─ ADNI_094_S_4089_MR_Sag_IR-SPGR_br_raw_20111019095512256_62_S125692_I261478.dcm
│  │       └─ ...
│  │   └─ 2011-12-14_15_53_24.0
│  │     └─ ...
│  │ └─ Sag_IR-SPGR_REPEAT
│  └─ Spatially_Normalized_Masked_and_N3_corrected_T1_image
├─ 094_S_4162
└─ ...
```

## ADNI\_BIDS

```
├─ sub-ADNI094S4089
│  │ └─ ses-M00
│  │   └─ anat
│  │     └─ sub-ADNI094S4089_ses-M00_T1w.nii.gz
│  │   └─ dwi
│  │     └─ sub-ADNI094S4089_ses-M00_acq-axial_dwi.bval
│  │     └─ sub-ADNI094S4089_ses-M00_acq-axial_dwi.bvec
│  │     └─ sub-ADNI094S4089_ses-M00_acq-axial_dwi.nii.gz
│  │   └─ pet
│  │     └─ sub-ADNI094S4089_ses-M00_trc-av45_pet.nii.gz
│  │     └─ sub-ADNI094S4089_ses-M00_trc-fdg_pet.nii.gz
│  │   └─ sub-ADNI094S4089_ses-M00_scans.tsv
│  └─ ses-M03
│  └─ ses-M12
│  └─ ses-M24
├─ sub-ADNI094S4162
└─ ...
```



# Software platform for neuroimaging studies



## BIDS Converters

```
clinica convert <dataset>-to-bids \
dataset_dir clinical_data_dir bids_dir
```

Conversion of datasets to the BIDS format:



+

## I/O Tools

```
clinica iotools create-subjects-visits ...
merge-tsv ...
check-missing-modalities ...
```

## Main Third-Party Dependencies

Advanced Normalization Tools

ANTs

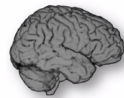
FreeSurfer



SPM

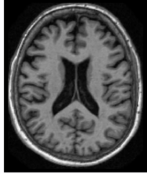
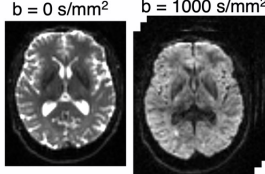
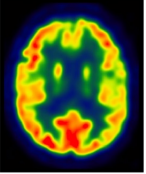

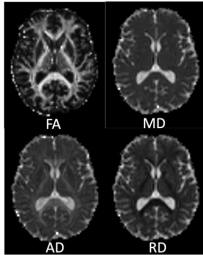
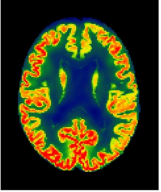
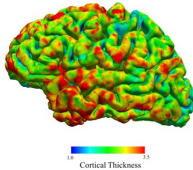
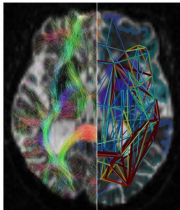
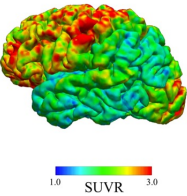


FSL



MRtrix

## Preprocessing and features extraction: `clinica run pipeline [BIDS dataset] [CAPS dataset] [options]`

Anatomical MRI	Diffusion MRI	PET
<p><code>{t1 flair}-linear</code></p> <p>Bias field correction, affine registration to standard space (MNI) and cropping (ANTs)</p> 	<p><code>dwi-preprocessing</code></p> <p>Correction of raw DWI data (FSL, ANTs, MRtrix3)</p> <p>b = 0 s/mm<sup>2</sup>    b = 1000 s/mm<sup>2</sup></p> 	<p><code>pet-linear</code></p> <p>Affine registration to standard space (MNI) via T1 MRI, intensity normalization and cropping (ANTs)</p> 
<p><code>t1-volume*</code></p> <p>Tissue segmentation (GM, WM, CSF), normalization to standard space (MNI) (SPM, CAT12)</p> 	<p><code>dwi-dti</code></p> <p>Extraction of DTI-based measures, normalization to standard space (MNI) (FSL, MRtrix3, ANTs)</p> <p>FA    MD</p> <p>AD    RD</p> 	<p><code>pet-volume</code></p> <p>Registration to T1 MRI, partial volume correction, intensity normalization, spatial normalization to standard space (MNI) (SPM, PETPVC, CAT12)</p> 
<p><code>t1-freesurfer*</code></p> <p>Cortical surface extraction, segmentation of subcortical structures, cortical thickness estimation, spatial normalization to standard space (FreeSurfer)</p> 	<p><code>dwi-connectome</code></p> <p>Tractography &amp; connectome (FSL, FreeSurfer, MRtrix3)</p> 	<p><code>pet-surface*</code></p> <p>Projection of PET signal onto surfaces of the cortex (FreeSurfer, FSL, SPM, PETPVC)</p> 

## Standardized data structures



## Machine Learning

SVM, logistic regression, random forest

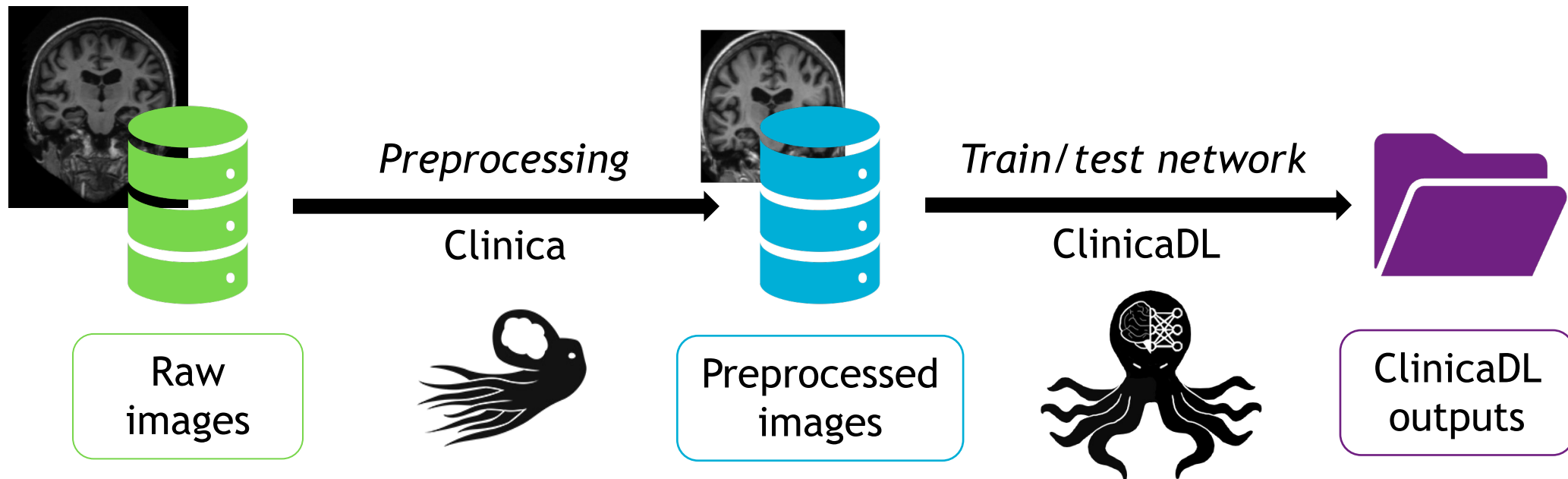


(Repeated) K-fold cross validation, repeated hold-out validation

## Statistical models

Surface-based analysis (`statistics-surface`):

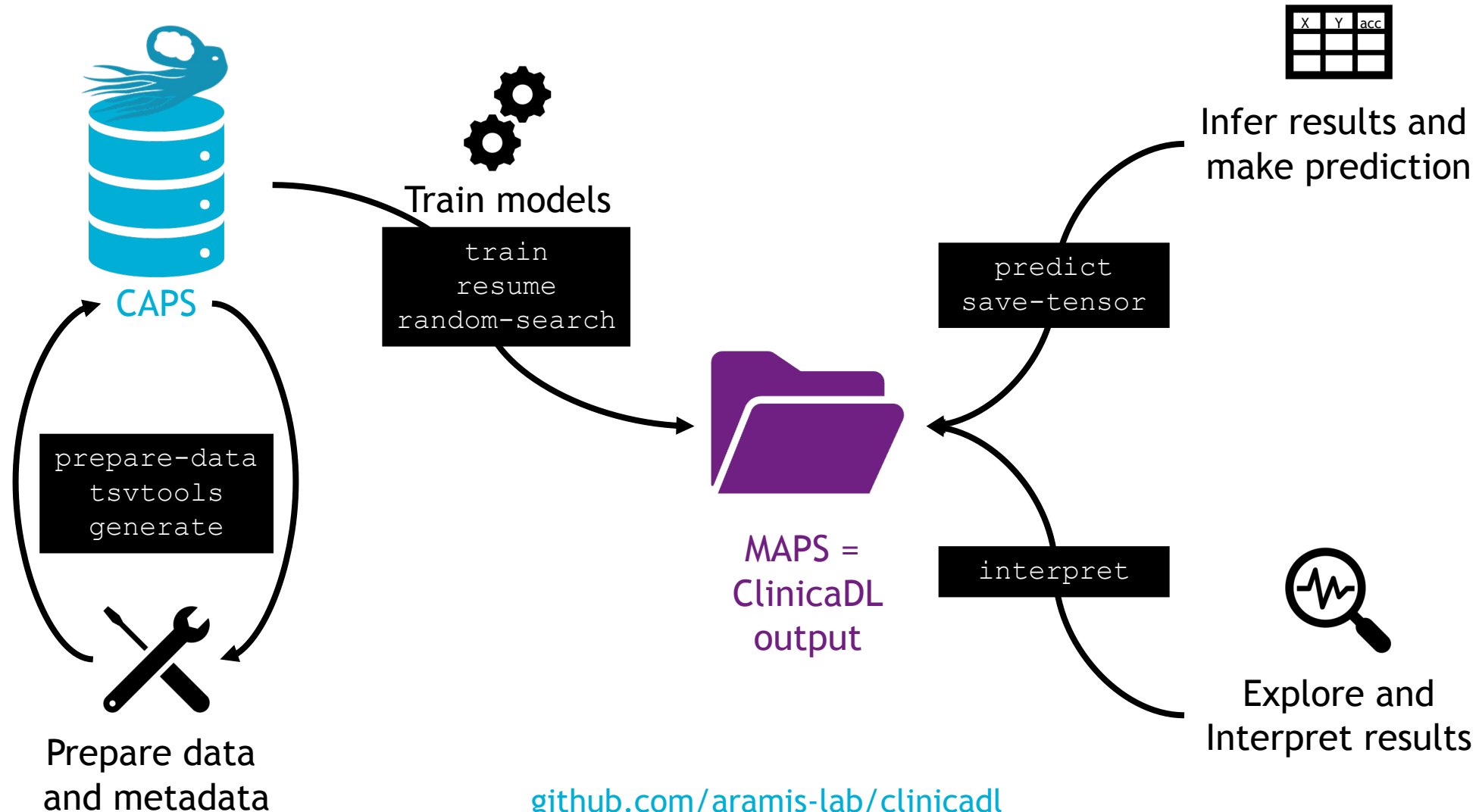
- Group comparison
- Correlation







## Open-source software for reproducible DL in neuroimaging

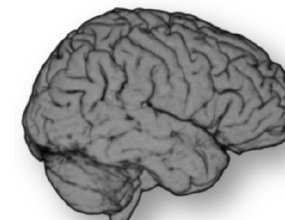
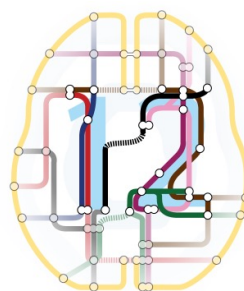


# Clinica's environment

GitHub



Advanced  
Normalization  
Tools



ANTs

FreeSurfer

SPM

FSL

MRTrix

# Who develops Clinica

Simona Bottani

Jérémy Guillon

Stanley Durrleman

You?

Tristan Moreau

Thomas Jacquemont

Ghislain Vaillant

Omar El Rifai

Arnaud Marcoux

Ninon Burgos

Olivier Colliot

Nicolas Gensollen

Alexandre Routier

Pascal Lu

Alexis Guyot

Elina Thibeau--Sutre

Matthieu Joulot

Jorge Samper-Gonzalez

Ravi Hassanaly

Sabrina Fontanella

Junhao Wen

Adam Wild

Michael Bacci

Mauricio Diaz

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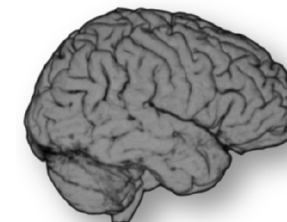
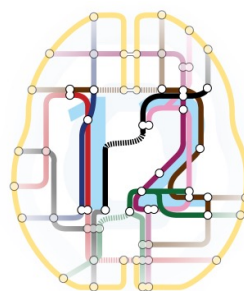
Mauricio Diaz

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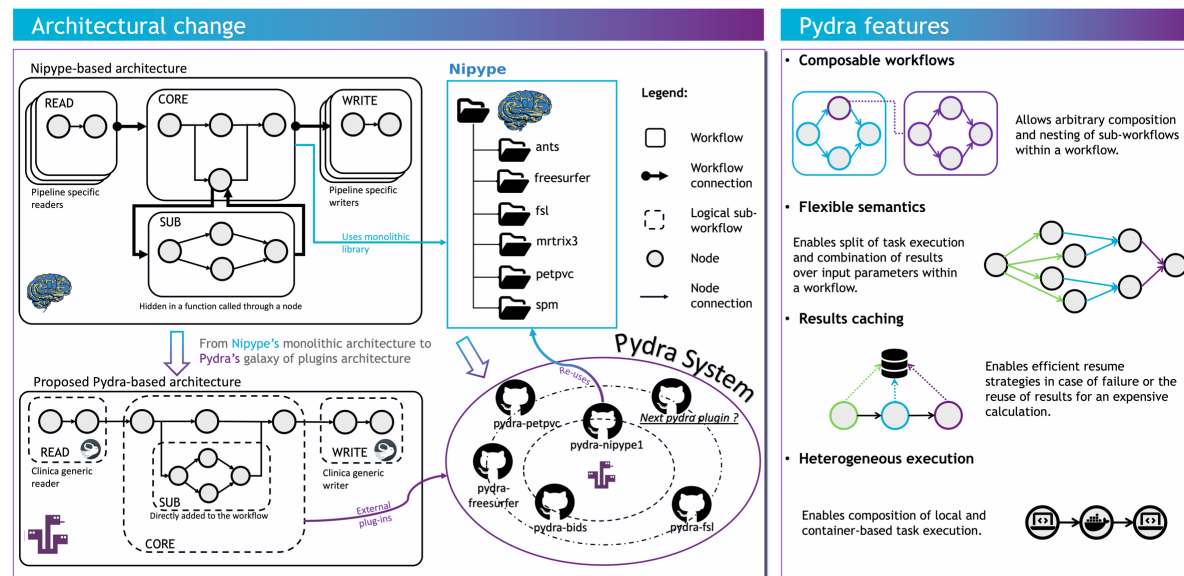
MRTrix



## BIDS

- BIDS Extension Proposal 009: Positron Emission Tomography → v1.6.0
- BIDS Extension Proposal 028: Provenance

## Pydra (aka the new Nipype)



# Was it worth making Clinica open?

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## A few numbers

- 84 citations of Routier et al., *Frontiers in Neuroinformatics*, 2021
  - 60+ actually use the software, 30+ outside the ICM
- ~200 stars on GitHub, ~60 forks, ~200 issues opened (~65 by external users), ~170 Google Group conversations

## What cannot be counted

Olivier Colliot  
Ninon Burgos  
Stanley Durrleman  
Nicolas Gensollen

Michael Bacci  
Simona Bottani  
Mauricio Diaz  
Omar El Rifai  
Sabrina Fontanella  
Jérémy Guillon



[www.clinica.run](http://www.clinica.run)

Alexis Guyot  
Ravi Hassanaly  
Thomas Jacquemont  
Matthieu Joulot  
Pascal Lu  
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Tristan Moreau  
Alexandre Routier  
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Adam Wild

**Software platform for neuroimaging studies**