



**ARAMIS
LAB**
BRAIN DATA SCIENCE



Paris **Brain
Institute**
ICM

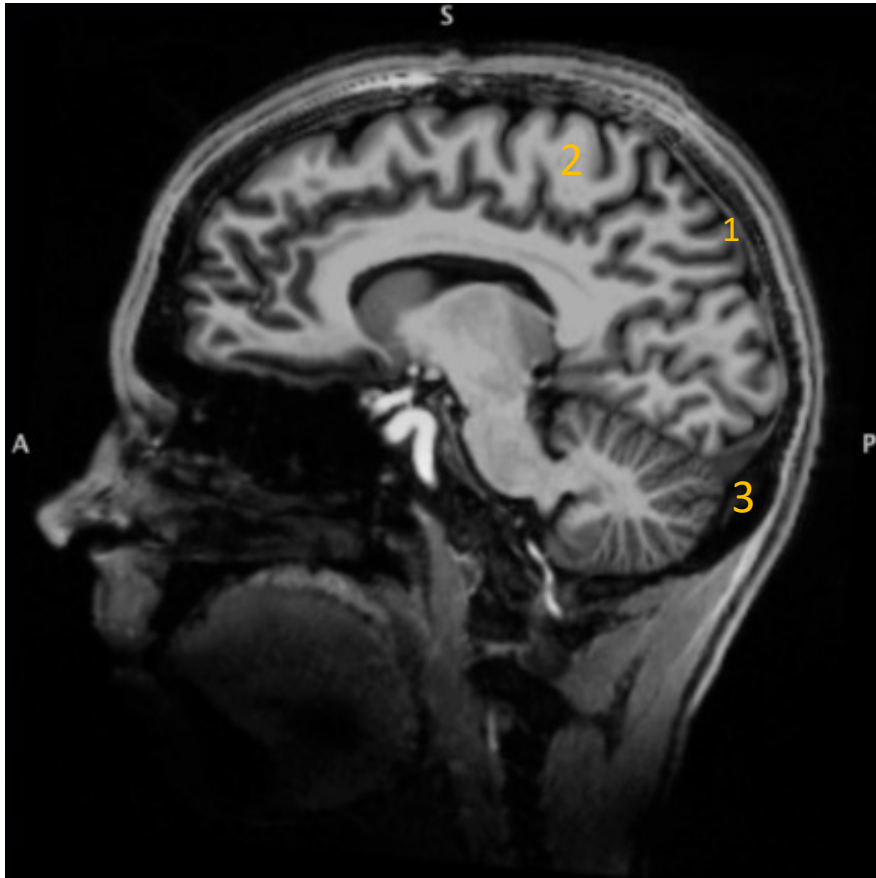
February 6th, 2024

Using UK Biobank Imaging data : Focus on T1w images

Elise Delzant, 2nd -year PhD student

Supervisor : Baptiste Couvy-Duchesne, PhD

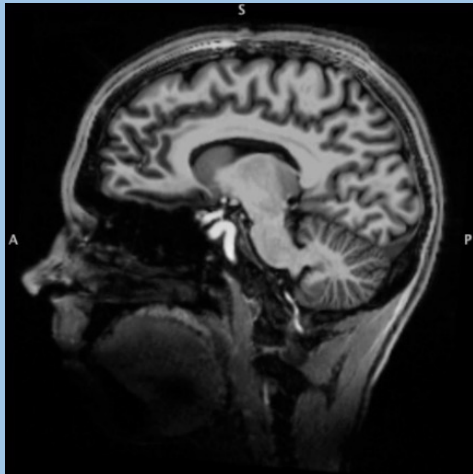
What is a T1w image ?



1. Grey-matter in dark grey
2. White-matter in light grey
3. Cerebro-spinal fluid (CSF) in dark

- Structural MRI
- Gives details of the brain :
 - enhances the signal of fatty tissue
 - suppresses signal of water
- Best depicts anatomy and can accentuate pathology

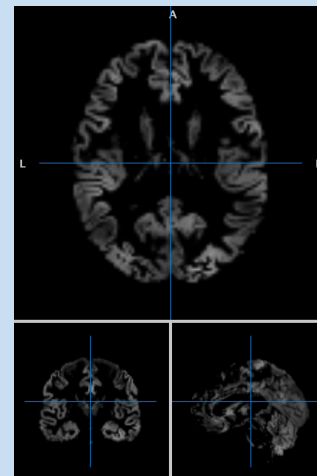
Brain MRI raw
image (T1)



Unstructured data

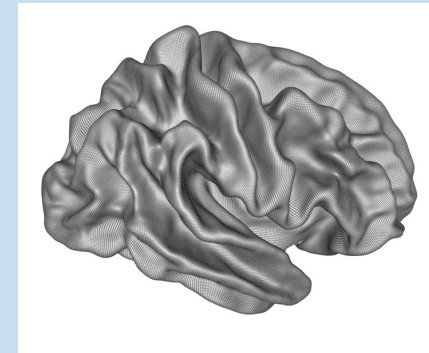
- *Deep-learning*

High dimensional processing



Volume-based

OR

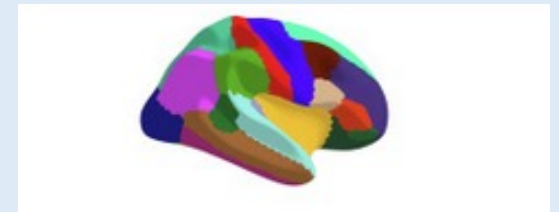


Surface-based

Structured data – high dimensional

- *High performance cluster*
- *Complex statistics, machine learning*

Region of Interest



*Desikian-Killiany atlas
(Desikian et al., 2006). 68 ROI*

Structured data – low dimensional

- *Simple statistics*

- 0 Initial assessment visit (2006-2010) at which participants were recruited and consent given
- 1 First repeat assessment visit (2012-13)
- 2 Imaging visit (2014+)
- 3 First repeat imaging visit (2019+)

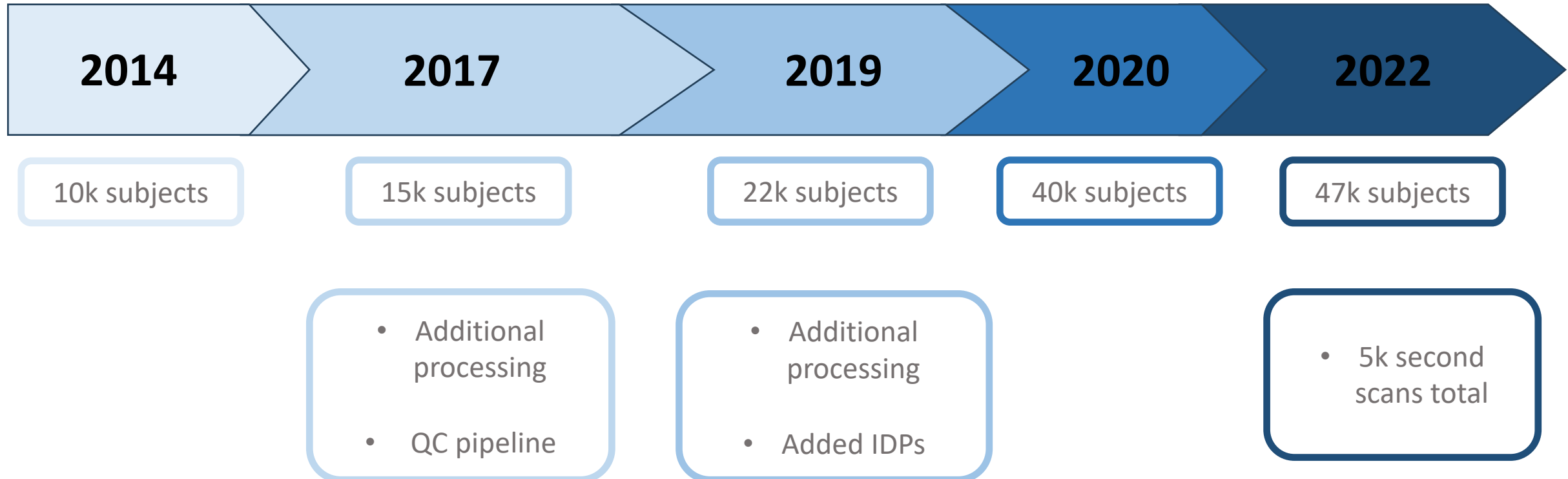


- 2 Imaging visit (2014+)
- 3 First repeat imaging visit (2019+)



5 centers :

- assessment around 4-5 hours
- provides detailed MRI measures of the brain



Field ID Description

Category

20252 T1 structural brain images - NIFTI T1 structural brain MRI ‡

*Raw + volume-based
processed images*

2nd visit : 66,952 participants
3rd visit : 5211 participants

2 types of T1w available :

*Surface-based
processed images*

Field ID Description

Category

20263 T1 surface model files and additional structural segmentations T1 structural brain MRI ‡

2nd visit : 43,139 participants
3rd visit : 4629 participants



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Data-Field 20252

Description: T1 structural brain images - NIFTI

Category: [Assessment centre](#) [Imaging](#) [Brain MRI](#) [T1 structural brain MRI](#)
[Brain MRI scans + XNAT image package + Brain MRI](#)

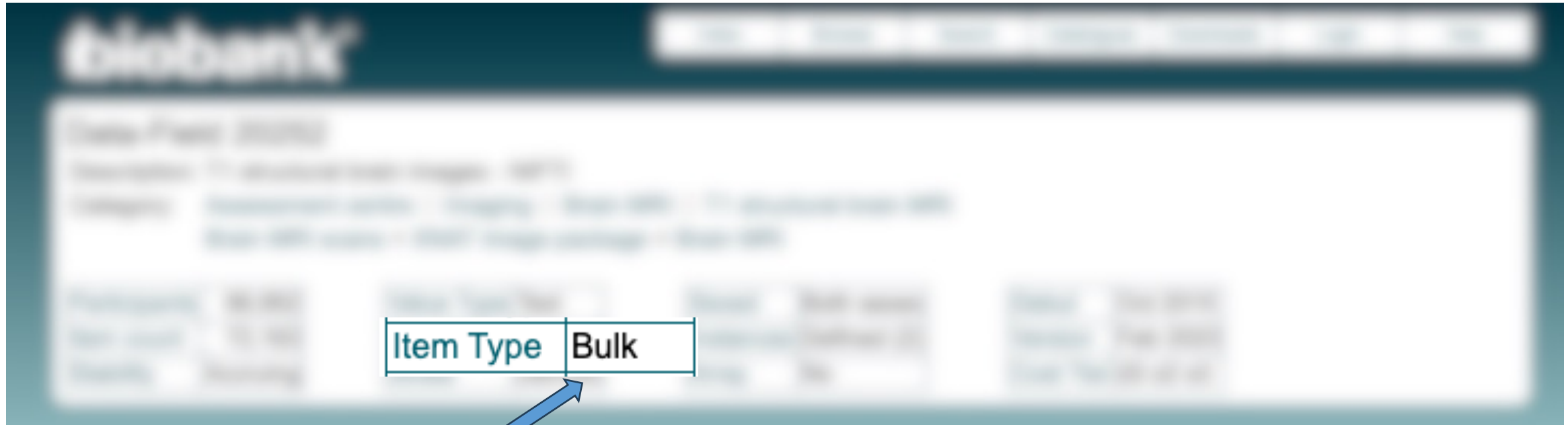
Participants	66,952
Item count	72,163
Stability	Accruing

Value Type	Text
Item Type	Bulk
Strata	Derived

Sexed	Both sexes
Instances	Defined (2)
Array	No

Debut	Oct 2015
Version	Feb 2023
Cost Tier	d3 o2 s3

Type : Bulk



- Large and complex items, such as genomic data, imaging data.

Field ID Description

Category

20252 T1 structural brain images - NIFTI T1 structural brain MRI ‡

- 1 fields marked ‡ are blob/bulk.

Download bulk files

```
ukbconv ukb673035.enc_ukb bulk -s20252
```



File ukb673035.bulk

Participant ID **Field_Instance_Array**

3422567 20252_2_0

5321753 20252_2_0

2457842 20252_3_0

Download bulk files

```
ukbconv ukb673035.enc_ukb bulk -s20252
```



File ukb673035.bulk

Participant ID **Field_Instance_Array**

3422567 20252_2_0

5321753 20252_2_0

2457842 20252_3_0

```
ukbfetch -bukb673035.bulk -ak53185r673035.key
```



Download every file from Field 20252

Download bulk files

```
ukbconv ukb673035.enc_ukb bulk -s20252
```



File ukb673035.bulk

Participant ID **Field_Instance_Array**

3422567 20252_2_0

5321753 20252_2_0

2457842 20252_3_0

```
ukbfetch -bukb673035.bulk -ak53185r673035.key
```



Download every file from Field 20252

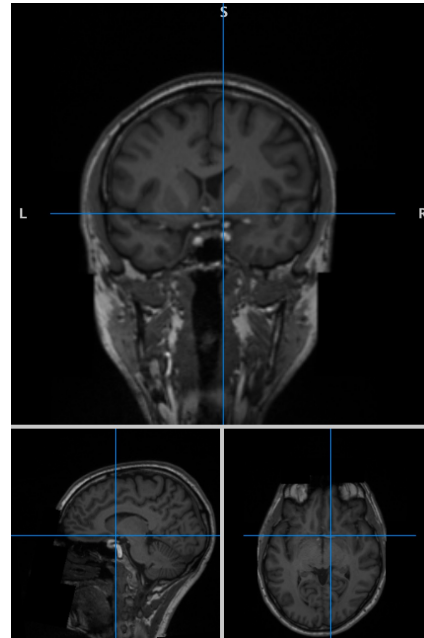
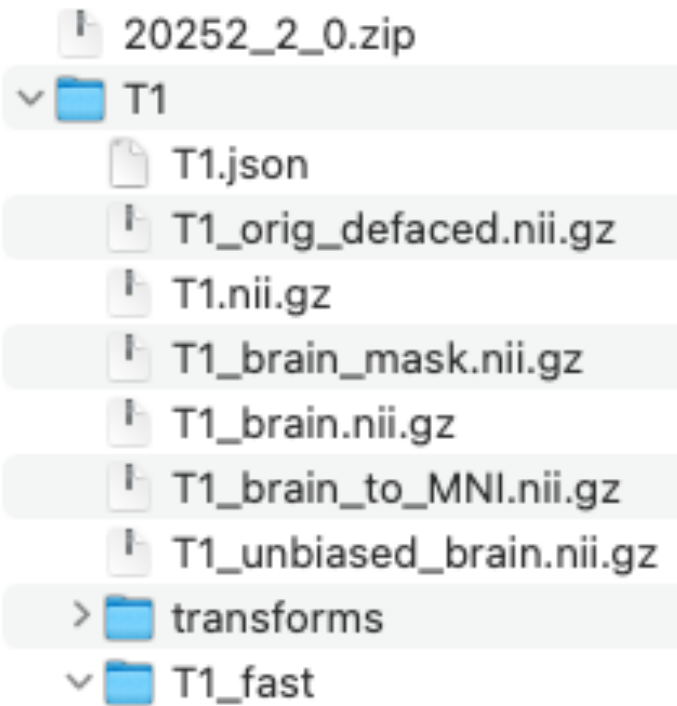
.bulk files can
be edited

Number of files
to download
can be specified

A limit of 1,000
files per
ukbfetch call

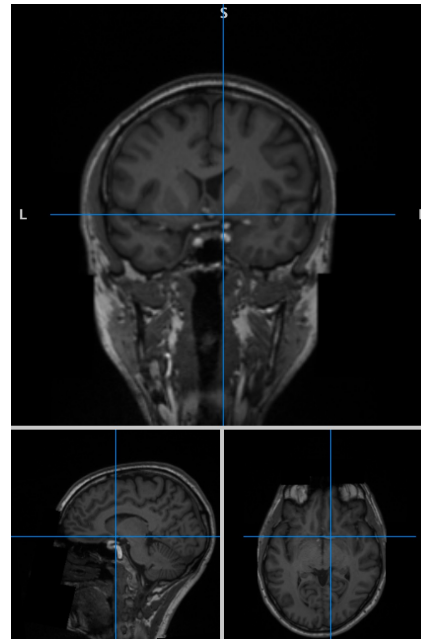
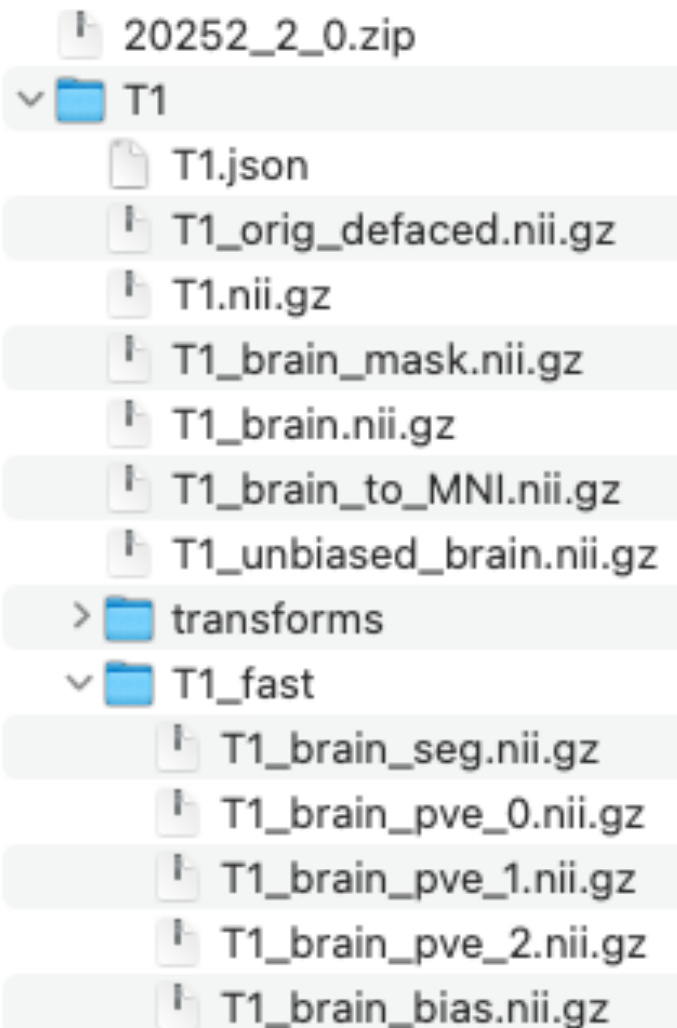
```
ukbfetch -bukb673035.bulk -s1 -m1000  
ukbfetch -bukb673035.bulk -s1001 -m2000  
ukbfetch -bukb673035.bulk -s2001 -m3000
```

<https://biobank.ndph.ox.ac.uk/ukb/field.cgi?id=20252>



T1_orig_defaced.nii.gz:
full raw T1 image after
defacing

<https://biobank.ndph.ox.ac.uk/ukb/field.cgi?id=20252>



T1_orig_defaced.nii.gz:
full raw T1 image after
defacing

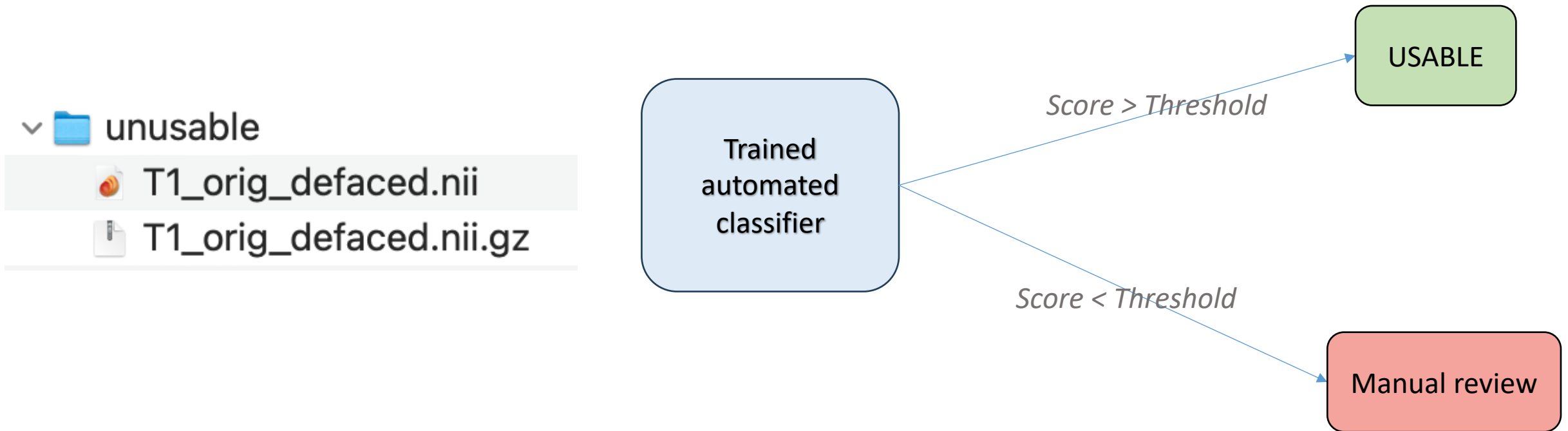


T1_brain_pve_1.nii.gz:
segmentation of
grey-matter



Jenkinson et al., 2012

**50 MB per
participant**

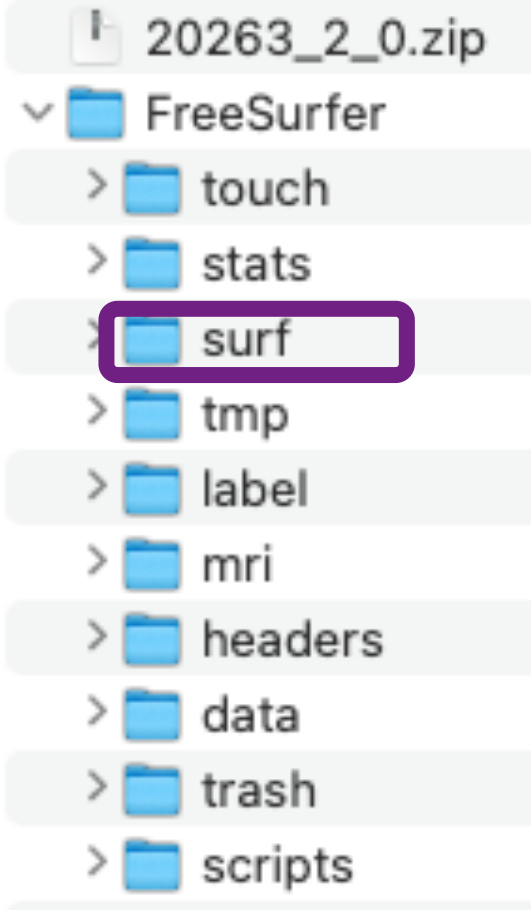


Surface-based processed images-Data Field 20263

<https://biobank.ndph.ox.ac.uk/ukb/field.cgi?id=20263>

- 20263_2_0.zip
- FreeSurfer
 - touch
 - stats
 - surf
 - tmp
 - label
 - mri
 - headers
 - data
 - trash
 - scripts

<https://biobank.ndph.ox.ac.uk/ukb/field.cgi?id=20263>

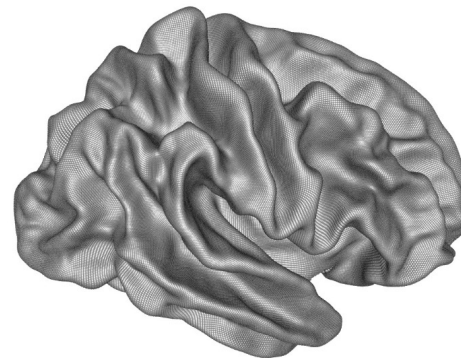


FreeSurfer

~ 6h per subject

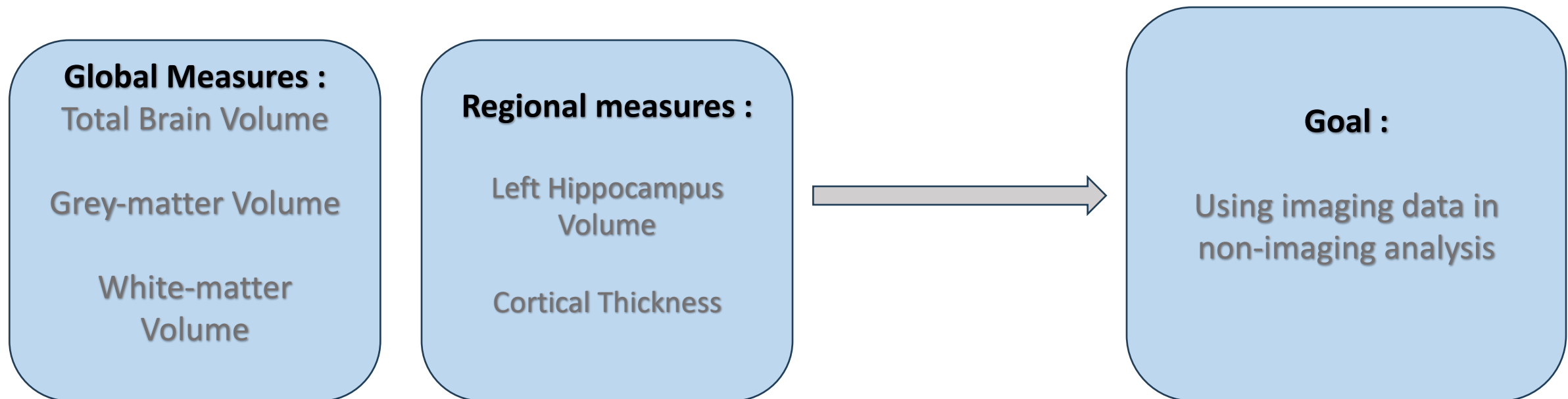
Surface model files :

- Cortical surface : left and right hemisphere area, volume, cortical thickness
- Subcortical data



350 MB per participant

Derived measures of brain structure



- *Association of a wide range of individual chronic diseases and their multimorbidity with brain volumes in the UK Biobank: A cross-sectional study, X. Shang et al., eClinicalMedecine, 2022*
- *Structural brain imaging correlates of general intelligence in UK Biobank, Cox et al., Intelligence, 2019*



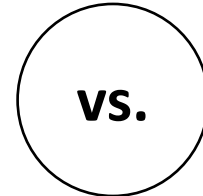
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**Example of an application with
T1w images**

Which of FSL or FreeSurfer capture the most information ?



FreeSurfer



N=42,272

X 350 MB = **15 TeraBytes**

N=42,272

X 50 MB = **2 TeraBytes**

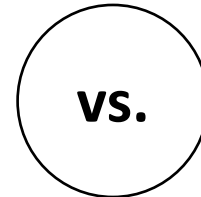
Which of FSL or FreeSurfer capture the most information ?



FreeSurfer

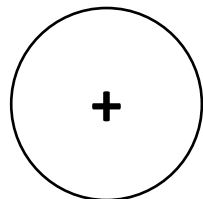
N=42,272

X 350 MB = **15 TeraBytes**

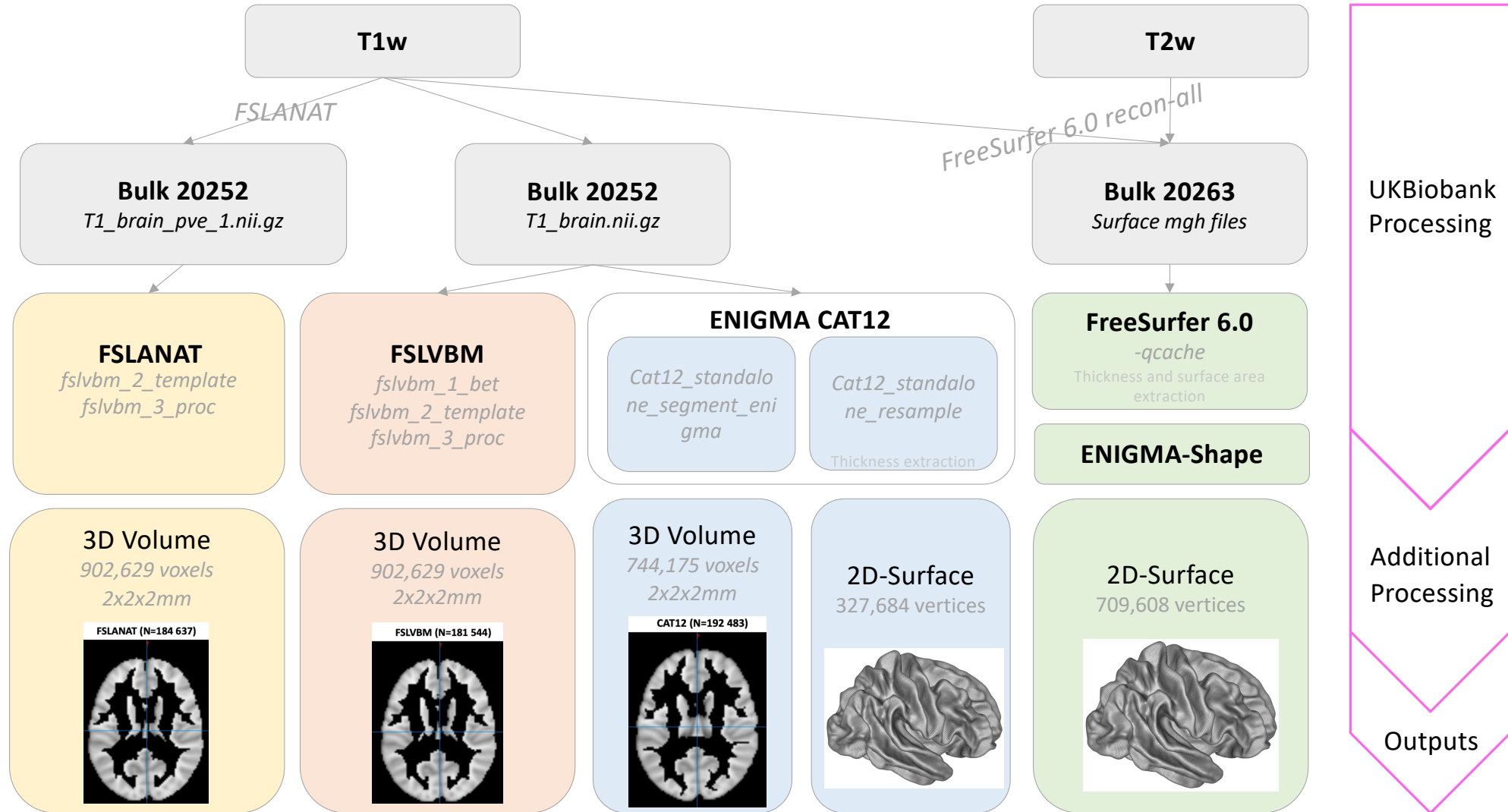


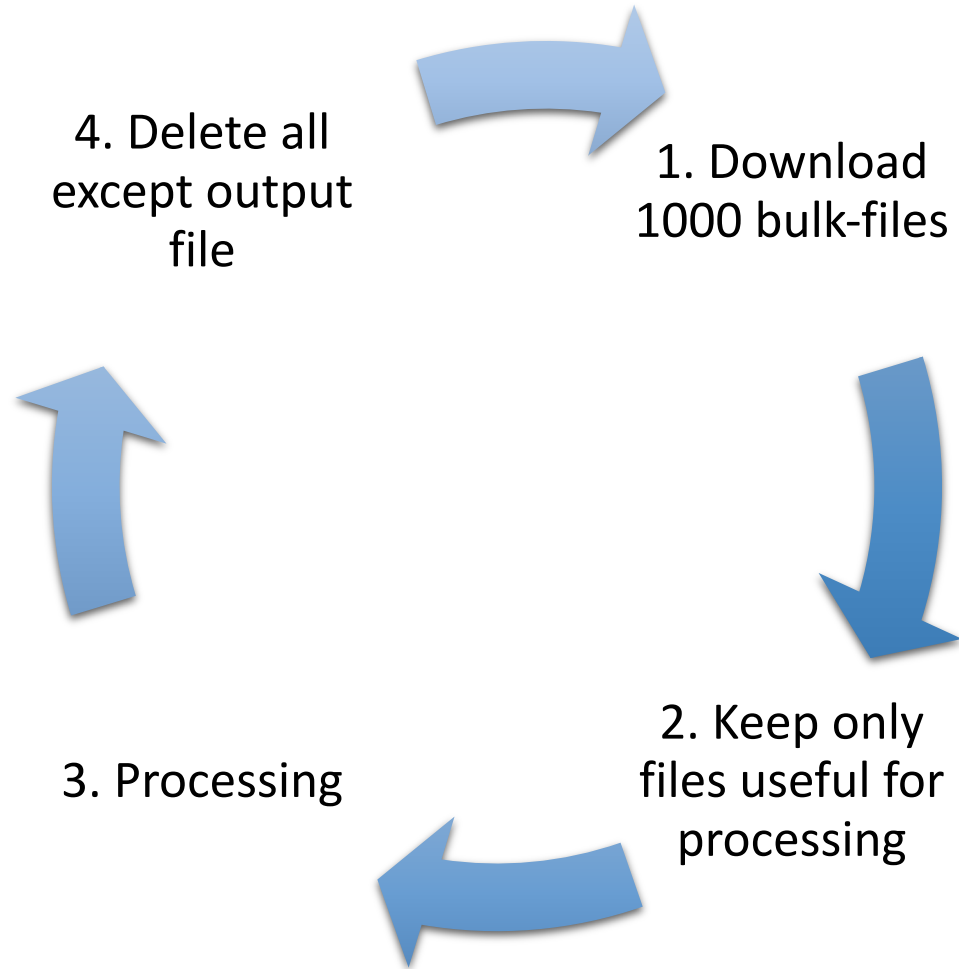
N=42,272

X 50 MB = **2 TeraBytes**



Detailed processing





N=42,272

X 350 MB = **15 TeraBytes**



N=42,272

X 50 MB = **2 TeraBytes**



1 file per subject per processing

FSL : 143G Total

We estimated, for each processing, the percentage of variance of a trait captured by all brain features

Potential confounders

- *Signal to noise ratio*
- *Time difference to first MRI*
- *Discrepancy between T1 and template*
- *Mean head motion*
- *X and Y brain position*

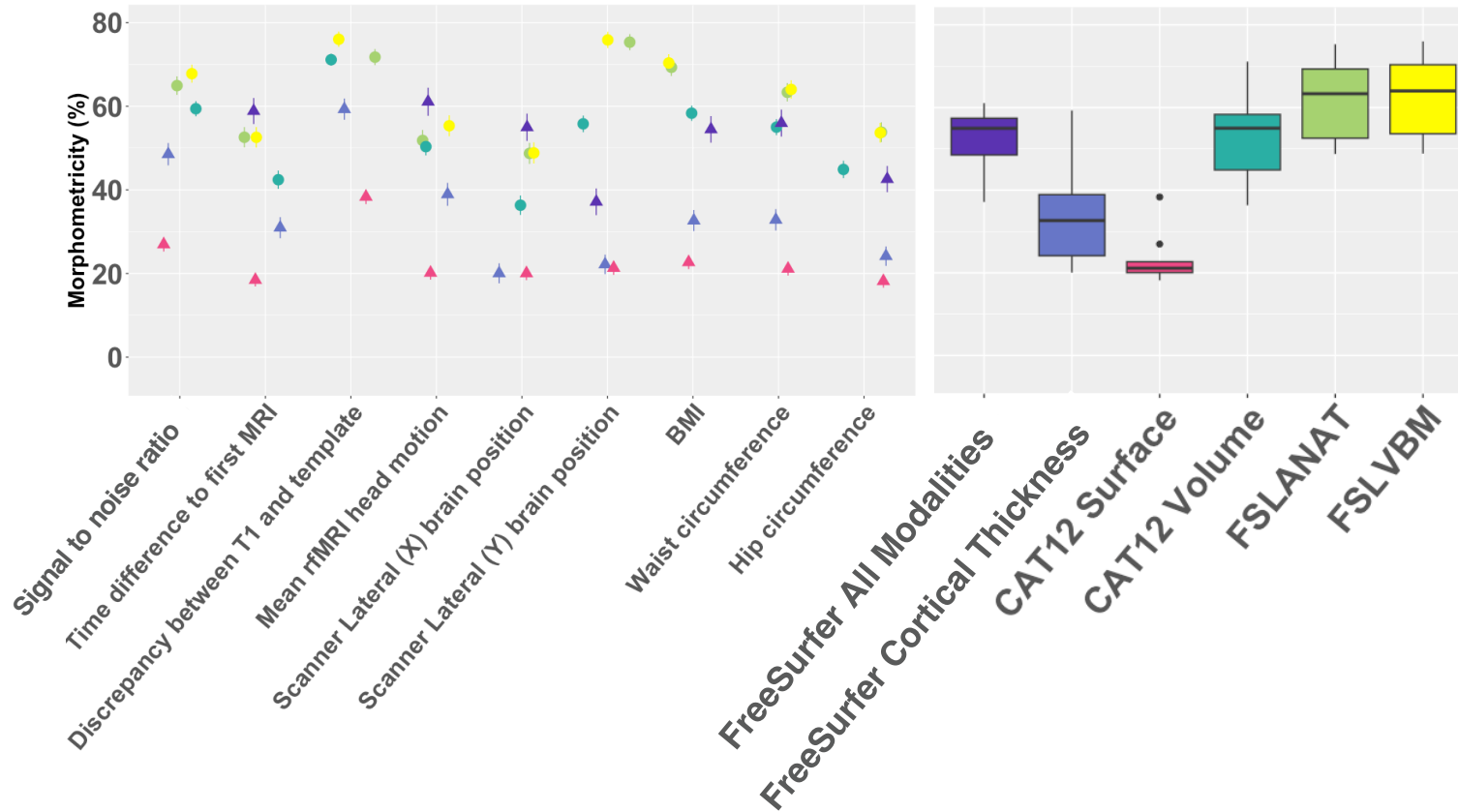
Body-size covariates

- *BMI*
- *Waist circumference*
- *Hip circumference*

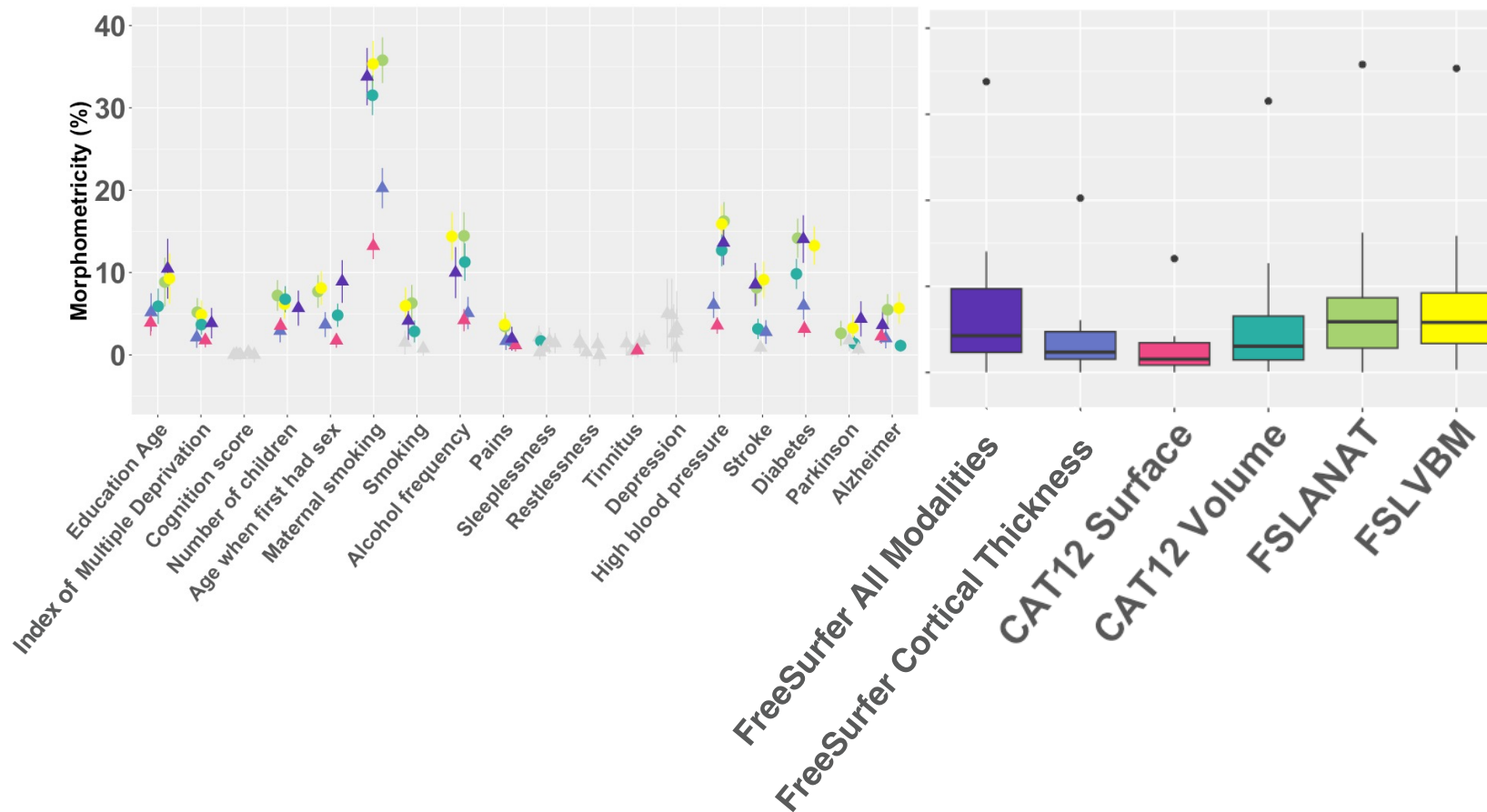
Traits of interest

- *Education*
- *Age*
- *Index of deprivation*
- *Cognition score*
- *Number of children*
- *Age First had sex*
- *Maternal smoking*
- *Smoking*
- *Alcohol frequency*
- *Pains*
- *Sleeplessness*
- *Restlessness*
- *Tinnitus*
- *Depression*
- *High blood pressure*
- *Stroke*
- *Diabetes*
- *Parkinson*
- *Alzheimer*

Results : potential confounders

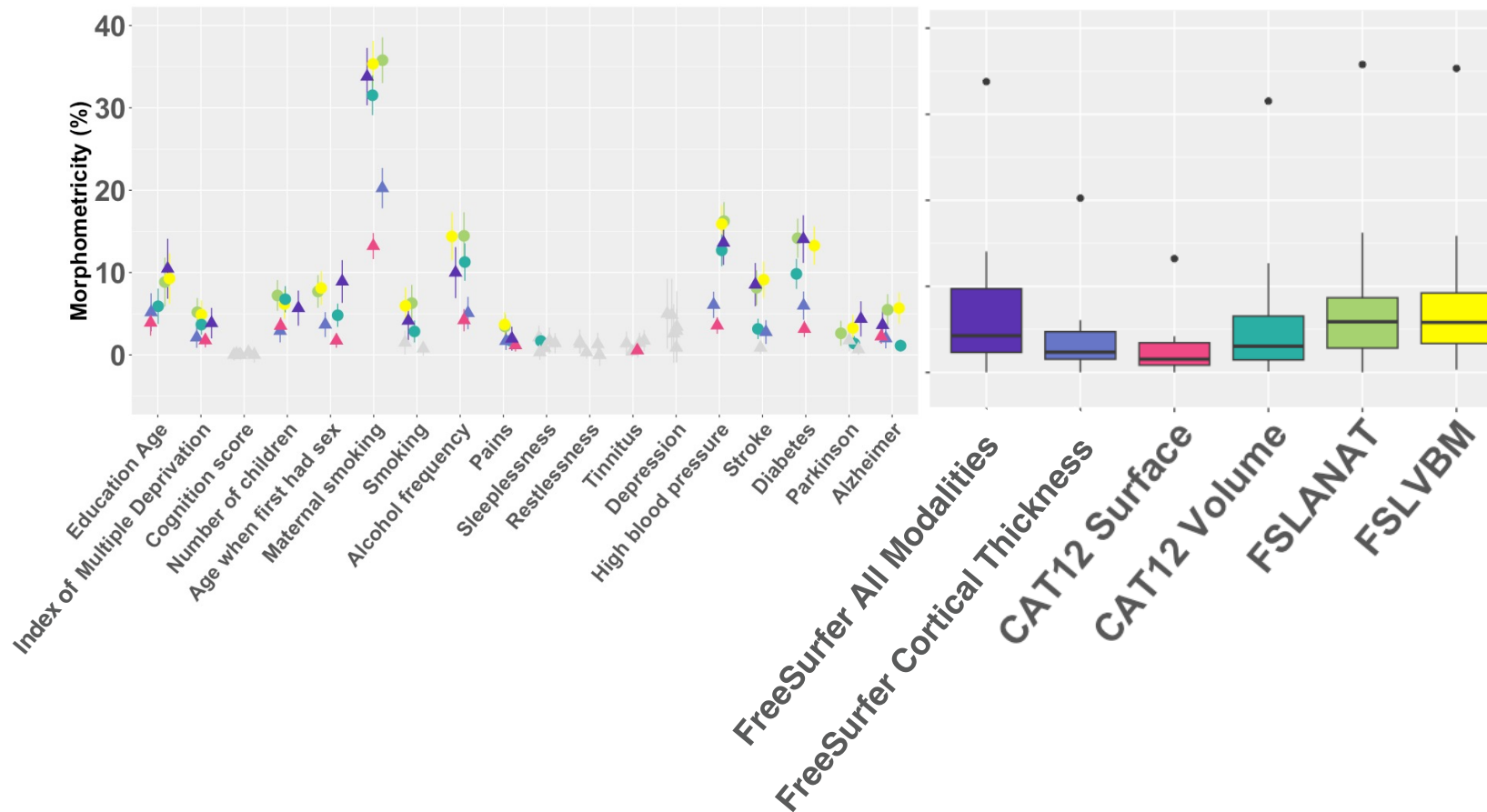


- All confounders should be included in grey-matter analysis of the UK Biobank



- The choice of processing impacts the percentage of variance estimated

Results : traits of interest



- The choice of processing impacts the percentage of variance estimated



UK Biobank Brain Imaging Documentation

Version 1.9
September 2022

primary documentation authors:

Stephen M. Smith, Fidel Alfaró-Almagro and Karla L. Miller

Wellcome Centre for Integrative Neuroimaging (WIN-FMRIB), Oxford University on behalf of UK Biobank

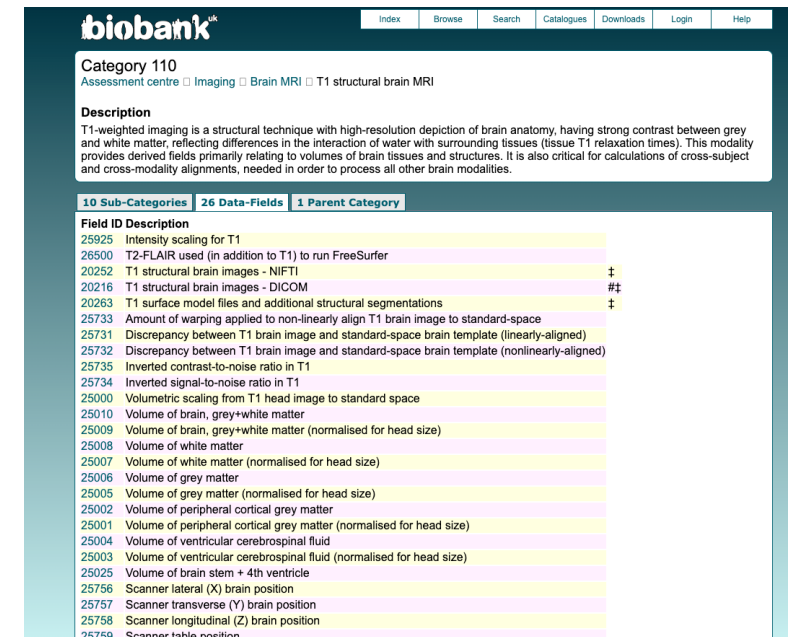
Resource | [Published: 19 September 2016](#)

Multimodal population brain imaging in the UK Biobank prospective epidemiological study

[Karla L Miller](#) , [Fidel Alfaró-Almagro](#), [Neal K Bangerter](#), [David L Thomas](#), [Essa Yacoub](#), [Junqian Xu](#), [Andreas J Bartsch](#), [Saad Jbabdi](#), [Stamatios N Sotiropoulos](#), [Jesper L R Andersson](#), [Ludovica Griffanti](#), [Gwenaëlle Douaud](#), [Thomas W Okell](#), [Peter Weale](#), [Iulius Dragonu](#), [Steve Garratt](#), [Sarah Hudson](#), [Rory Collins](#), [Mark Jenkinson](#), [Paul M Matthews](#) & [Stephen M Smith](#)

Image processing and Quality Control for the first 10,000 brain imaging datasets from UK Biobank

[Fidel Alfaró-Almagro](#),^{a,*} [Mark Jenkinson](#),^a [Neal K. Bangerter](#),^b [Jesper L.R. Andersson](#),^a [Ludovica Griffanti](#),^a [Gwenaëlle Douaud](#),^a [Stamatios N. Sotiropoulos](#),^{a,c} [Saad Jbabdi](#),^a [Moises Hernandez-Fernandez](#),^a [Emmanuel Vallee](#),^a [Diego Vidaurre](#),^d [Matthew Webster](#),^a [Paul McCarthy](#),^a [Christopher Rorden](#),^e [Alessandro Daducci](#),^{f,g} [Daniel C. Alexander](#),^h [Hui Zhang](#),^h [Iulius Dragonu](#),ⁱ [Paul M. Matthews](#),^{j,k} [Karla L. Miller](#),^a and [Stephen M. Smith](#)^a



Field ID	Description
25925	Intensity scaling for T1
26500	T2-FLAIR used (in addition to T1) to run FreeSurfer
20252	T1 structural brain images - NIFTI
20216	T1 structural brain images - DICOM
20263	T1 surface model files and additional structural segmentations
25733	Amount of warping applied to non-linearly align T1 brain image to standard-space
25731	Discrepancy between T1 brain image and standard-space brain template (linearly-aligned)
25732	Discrepancy between T1 brain image and standard-space brain template (nonlinearly-aligned)
25735	Inverted contrast-to-noise ratio in T1
25734	Inverted signal-to-noise ratio in T1
25000	Volumetric scaling from T1 head image to standard space
25010	Volume of brain, grey+white matter
25009	Volume of brain, grey+white matter (normalised for head size)
25008	Volume of white matter
25007	Volume of white matter (normalised for head size)
25006	Volume of grey matter
25005	Volume of grey matter (normalised for head size)
25002	Volume of peripheral cortical grey matter
25001	Volume of peripheral cortical grey matter (normalised for head size)
25004	Volume of ventricular cerebrospinal fluid
25003	Volume of ventricular cerebrospinal fluid (normalised for head size)
25025	Volume of brain stem + 4th ventricle
25756	Scanner lateral (X) brain position
25757	Scanner transverse (Y) brain position
25758	Scanner longitudinal (Z) brain position
25759	Scanner table position

<https://biobank.ndph.ox.ac.uk/ukb/label.cgi?id=110>

Thank you for listening !

Thank you to all UK Biobank staff and participants !



Any questions?

This research has been conducted using the UK Biobank Resource under Application Number 53185