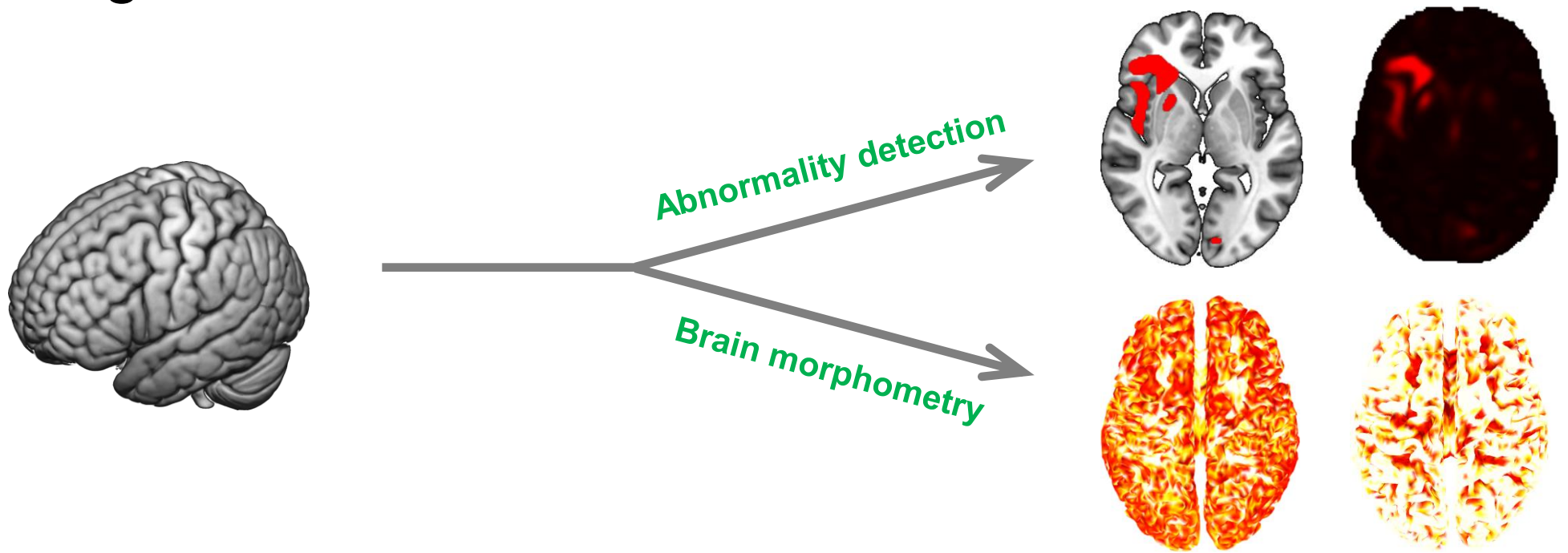


Structural MRI (2): Data Processing

구조 자기공명영상 (2):
데이터 처리 방법

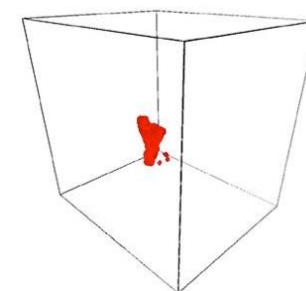
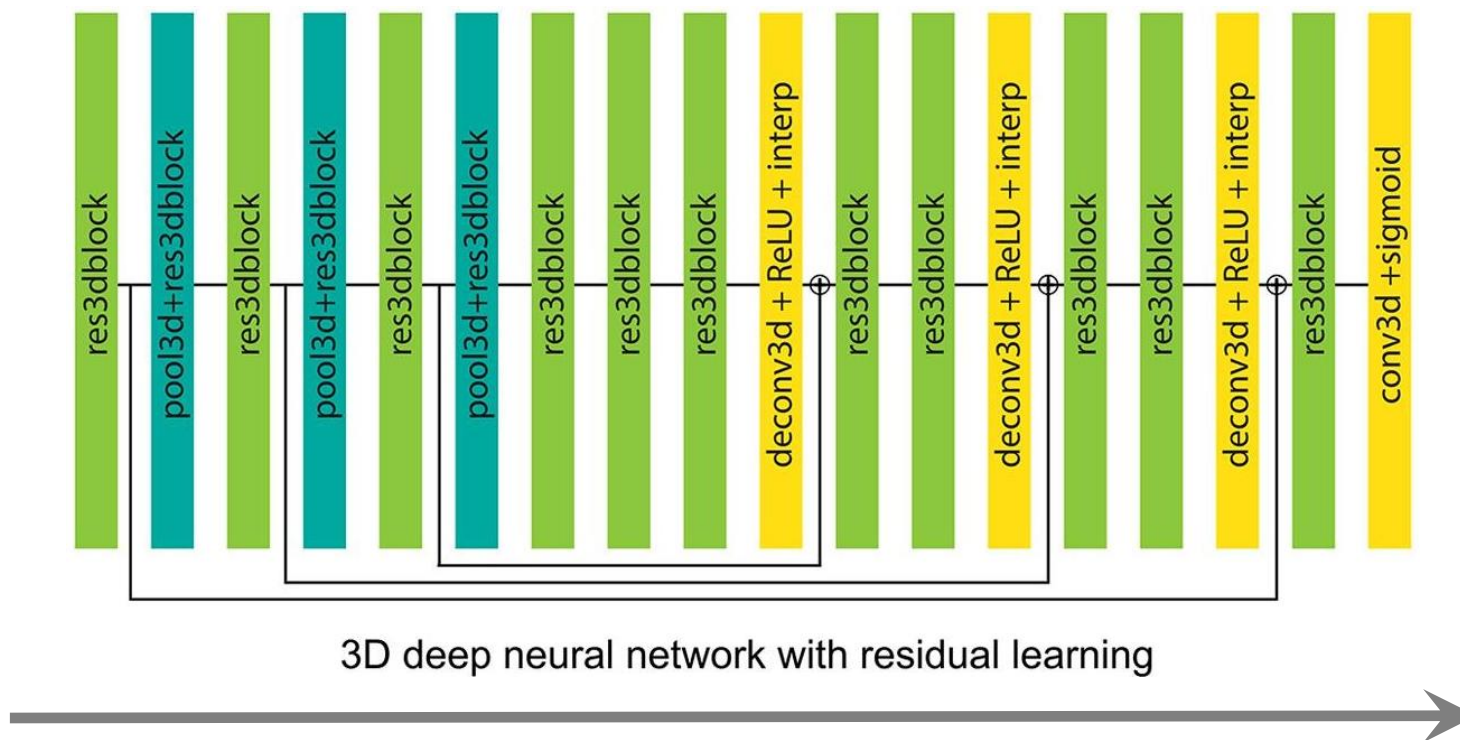
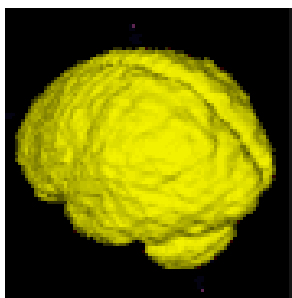
Brain Mapping with Structural MRI (sMRI)

- T1/T2-weighted sMRI



Automatic Abnormality Detection

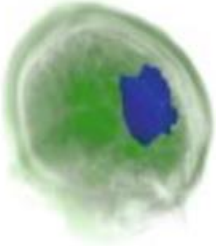

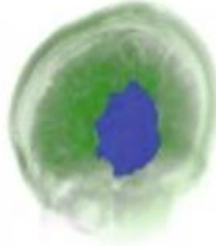
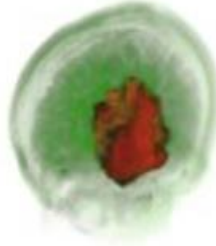


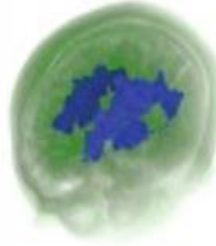





- Segmentation
 - 3D residual convolutional neural network for volumetric segmentation of stroke lesions on a T1-weighted image [\[Tomita et al., 2020\]](#)
- Grading
 - 3D residual convolutional neural network for predicting the severity of enlarged perivascular spaces on a T2-weighted image [\[Williamson et al., 2022\]](#)



[Tomita et al., 2020]

Automatic segmentation of a stroke lesion

$$DSC = \frac{2|X \cap Y|}{|X| + |Y|}$$

| DSC | Reference Standard | Predictions | Reference Standard | Predictions |
|-------|---|--|--|--|
| 0.813 |  |  |  |  |
| 0.788 |  |  |  |  |
| 0.801 |  |  |  |  |

[Tomita et al., 2020]

Evaluation of the performance of stroke lesion segmentation

Preprocessing

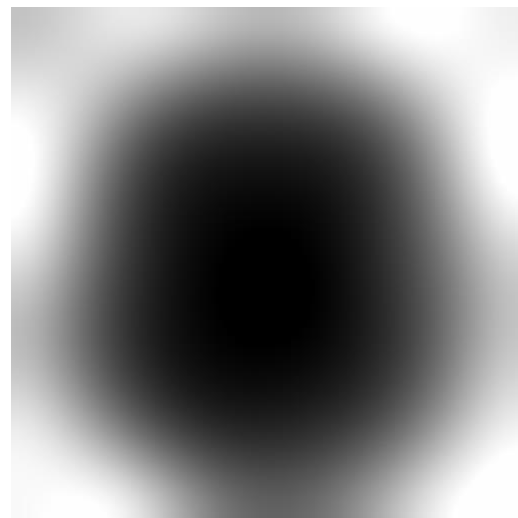
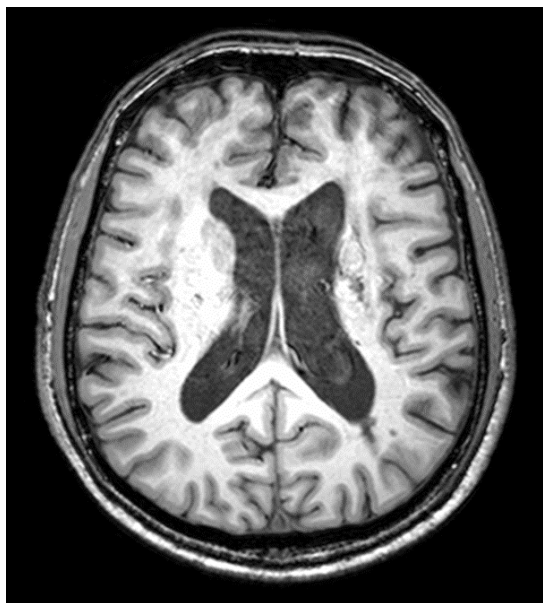
- Numerous steps to clean and standardise sMRI data before brain morphometry
 - Correction for bias fields (intensity non-uniformities)
 - From a broader range of sources, including imperfections in the MRI scanner's main magnetic field, inhomogeneities in the radiofrequency coil performance, and magnetic susceptibility-induced field inhomogeneities
 - Often characterized by a smooth variation in image brightness

– Segmentation

- Classifies an image into the non-brain and brain and, furthermore, the brain into different tissues usually including grey matter, white matter, and cerebrospinal fluid

– Normalisation

- Transforms an image from a native space to the standard space usually in the Montreal Neurological Institute (MNI) coordinate system



Intensity non-uniformity



Correction for intensity non-uniformity

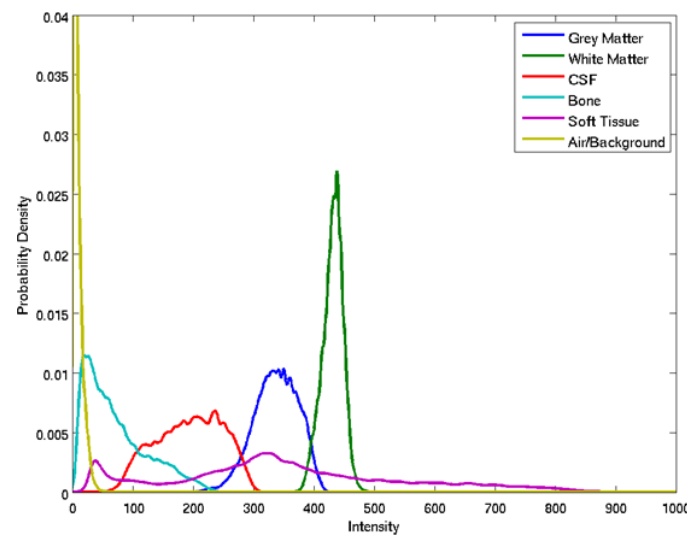
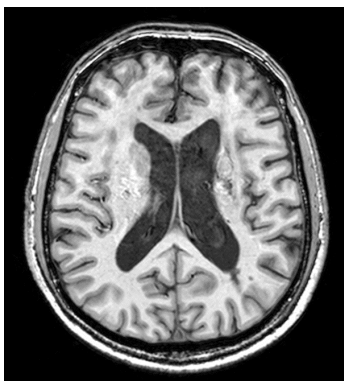
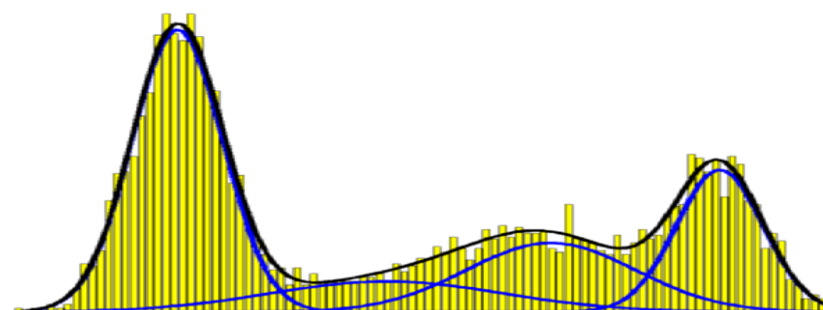


Image intensity distribution



Mixture of Gaussians model



Grey matter



White matter



Cerebrospinal fluid

Segmentation into different tissues

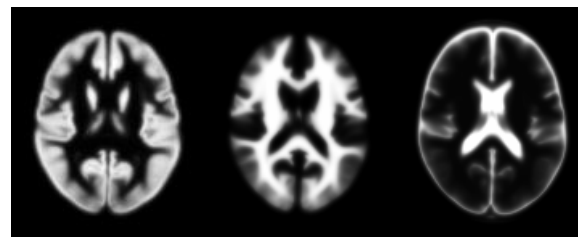
Grey matter



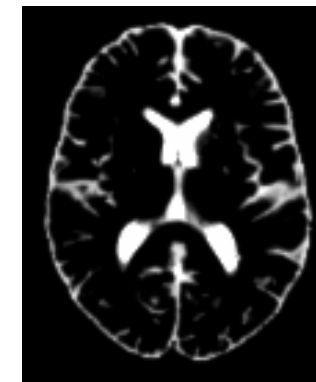
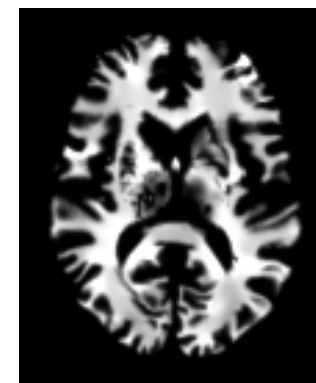
White matter



Cerebrospinal
fluid



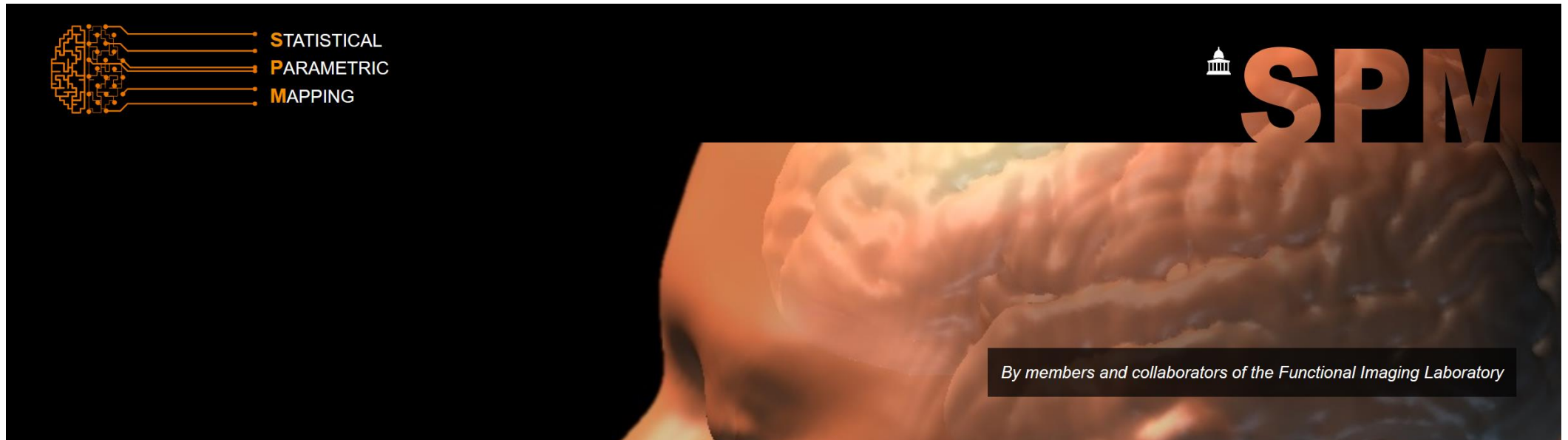
Template tissue probability maps

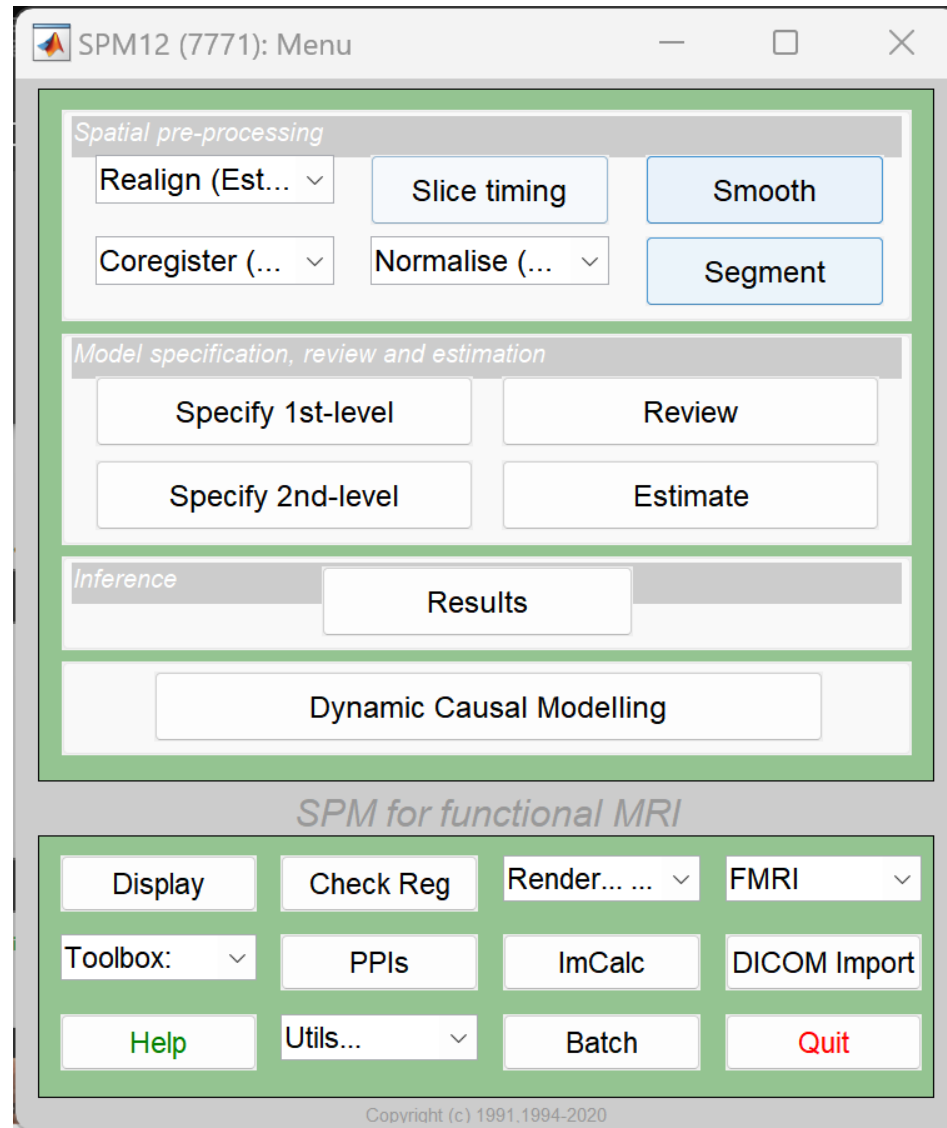


Normalisation

[Preprocessing of sMRI]

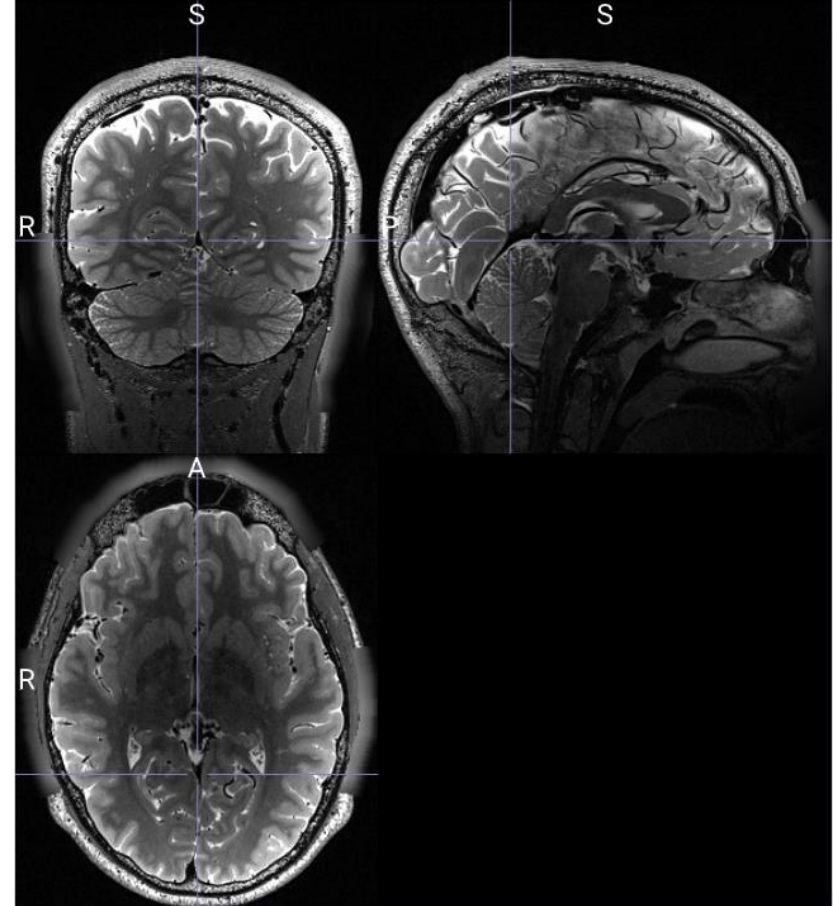
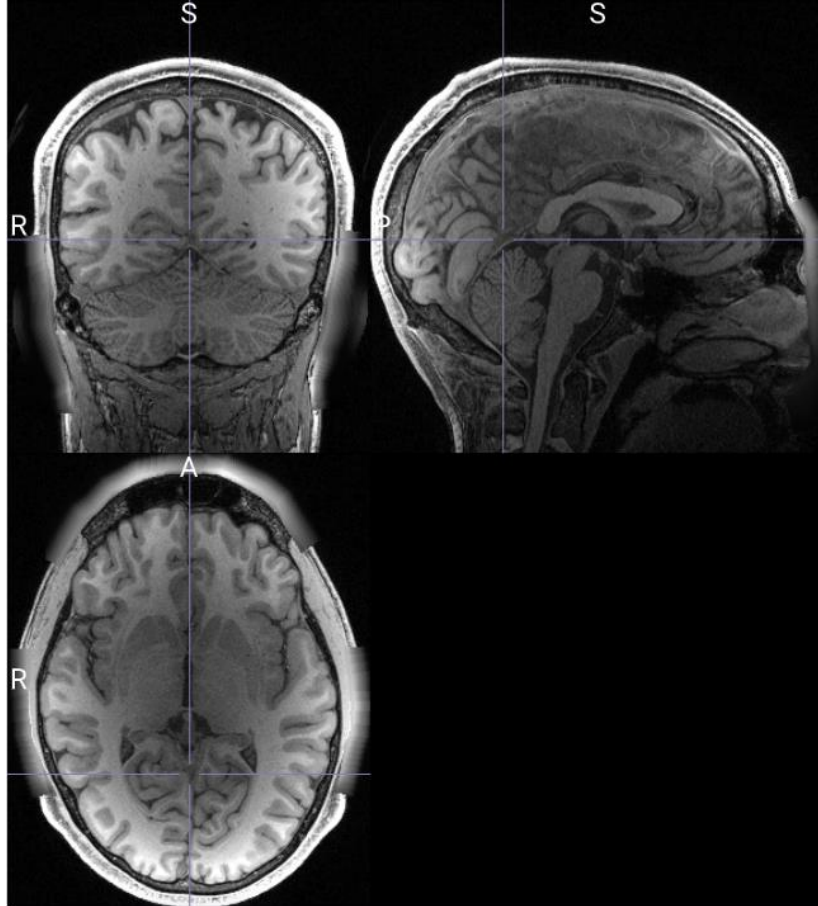
- SPM12 [<https://www.fil.ion.ucl.ac.uk/spm/software/spm12/>]





GUI of the SPM toolbox

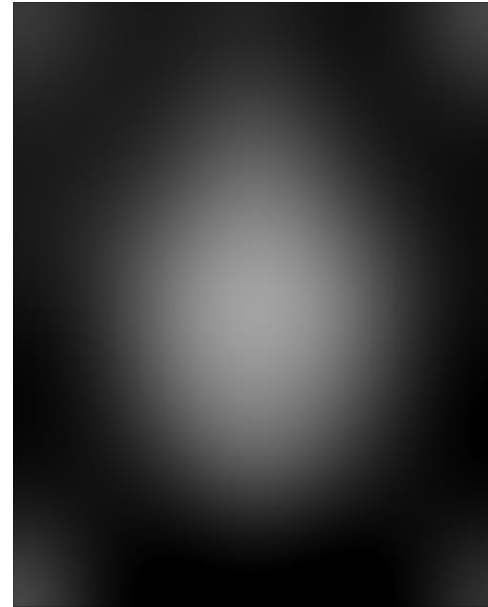
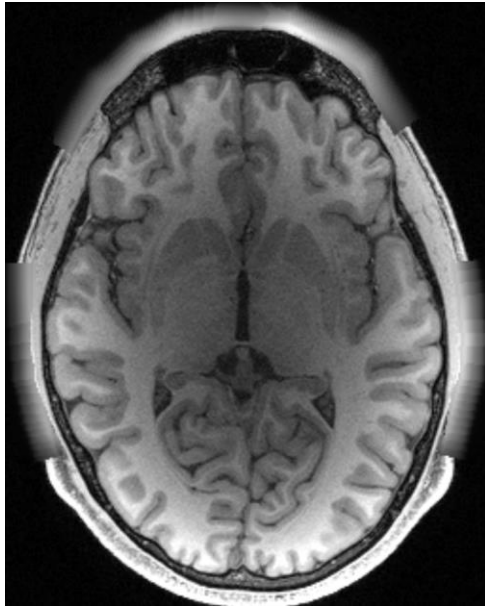
Input



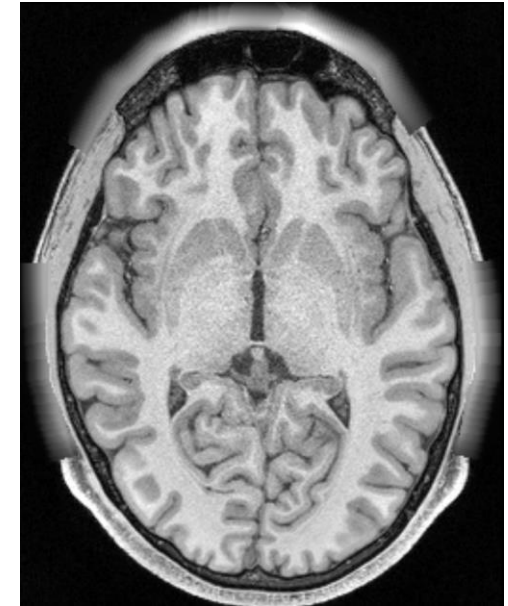
T1-weighted and T2-weighted sMRI

Output

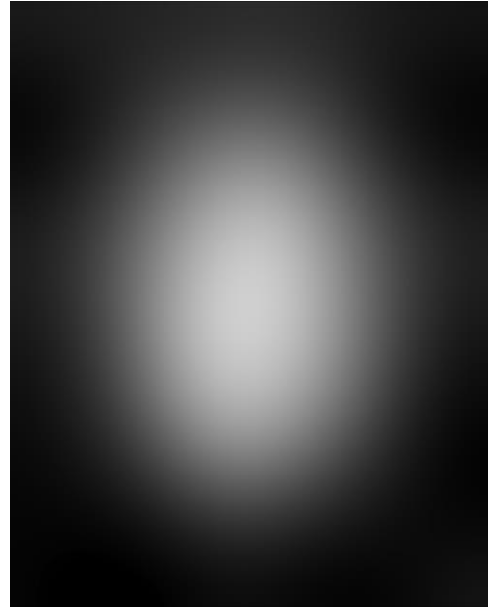
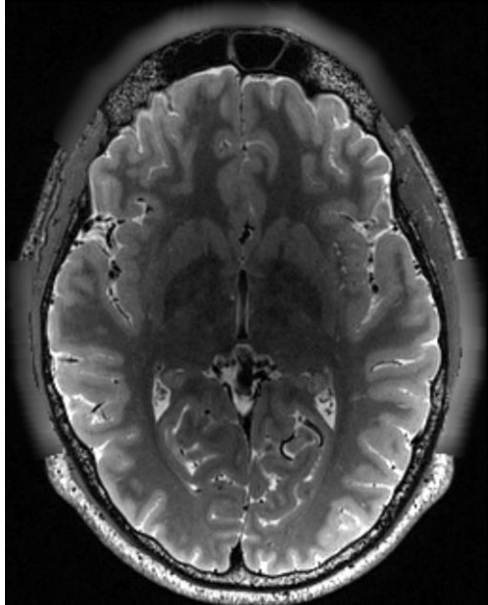
Correction for intensity non-uniformity



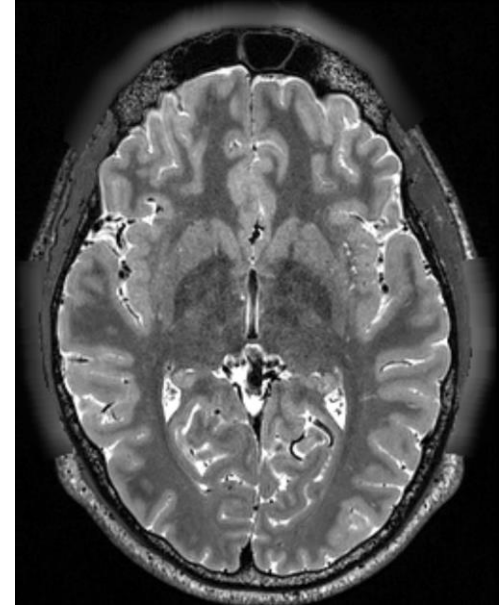
Intensity non-uniformity



Correction of the T1-weighted image for intensity non-uniformity



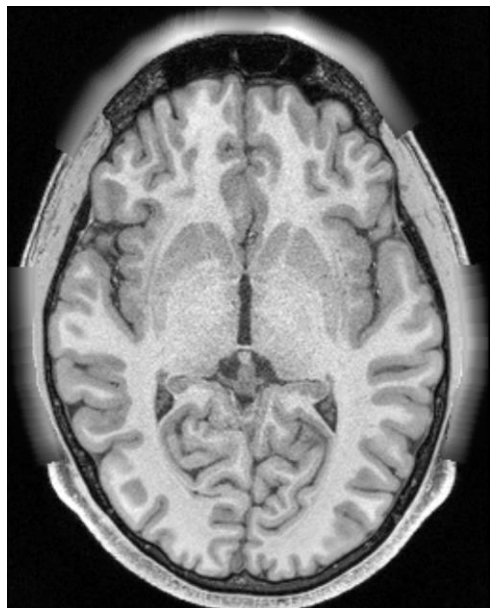
Intensity non-uniformity



Correction of the T2-weighted image for intensity non-uniformity

Output

Segmentation



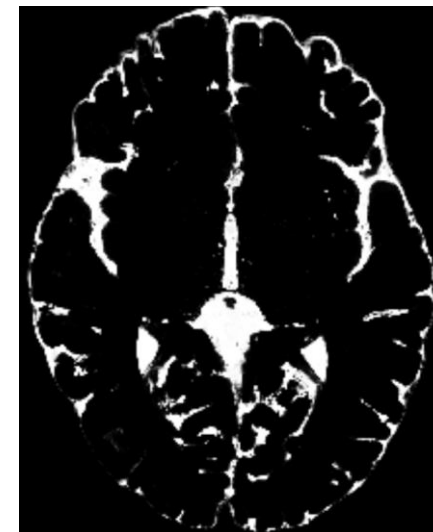
Segmentation



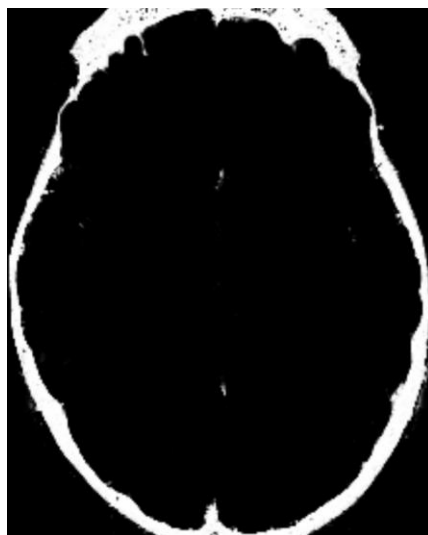
Grey matter



White matter



Cerebrospinal fluid



Bone



Soft tissue



Air/background

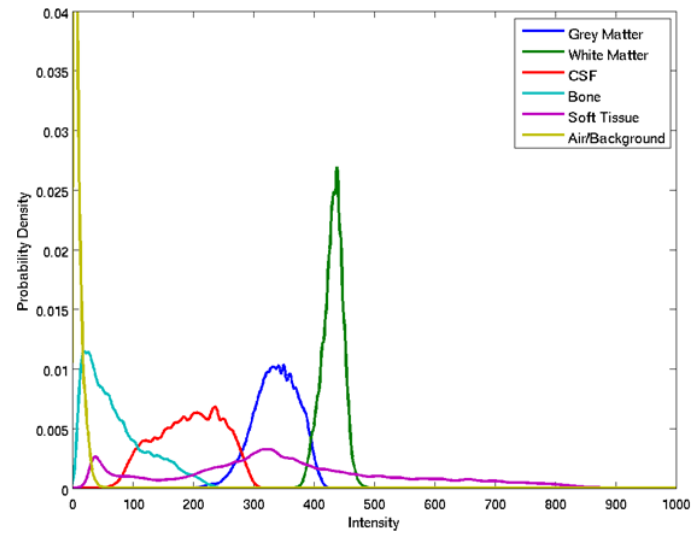
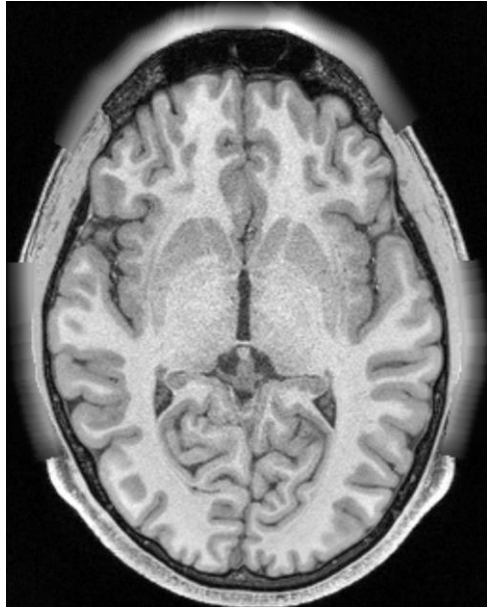
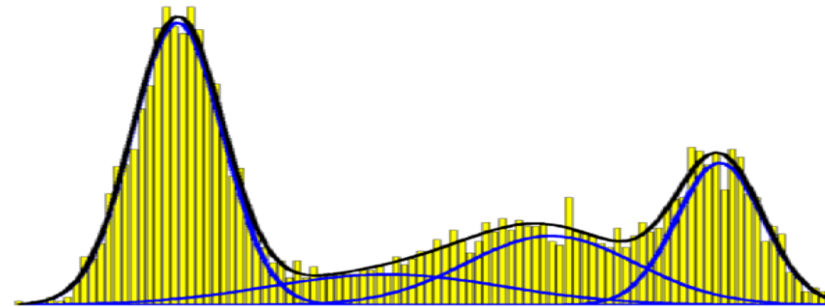
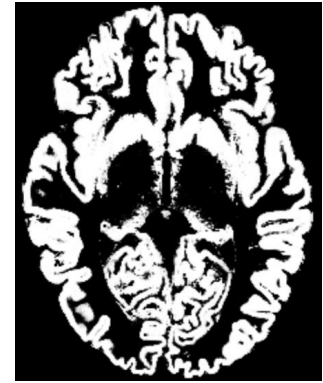


Image intensity distribution



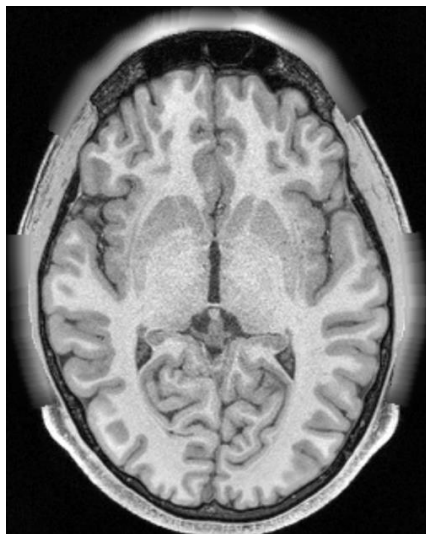
Mixture of Gaussians model



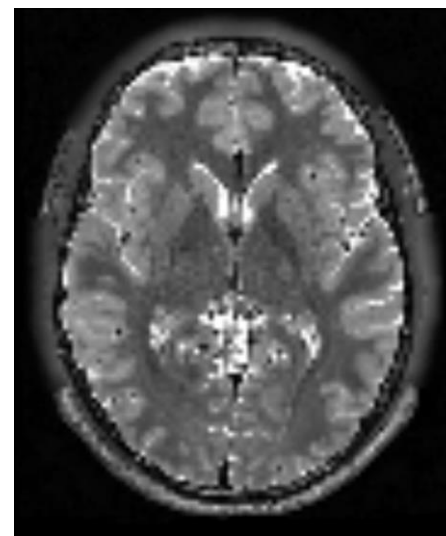
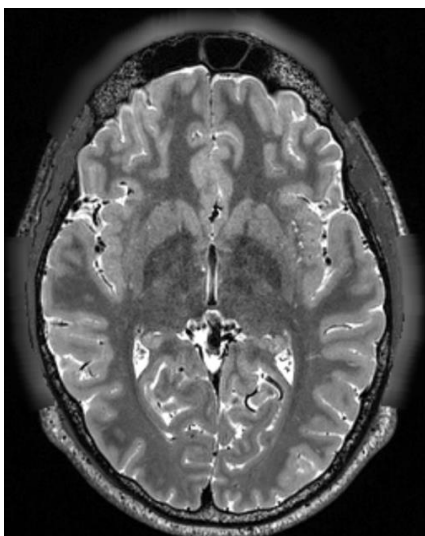
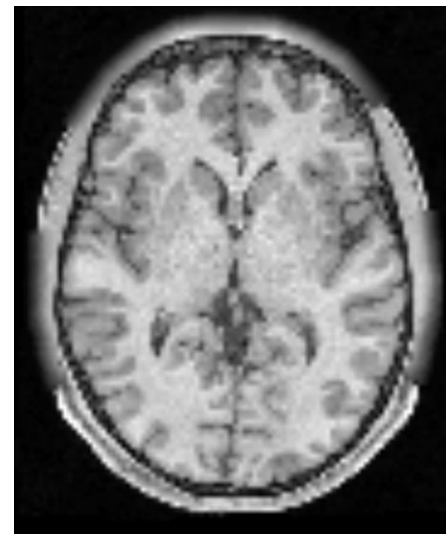
Tissue classification based on a mixture of Gaussians

Output

Normalisation



Normalisation



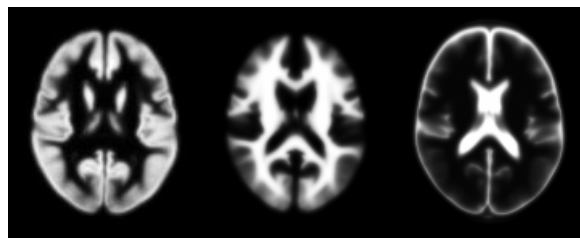
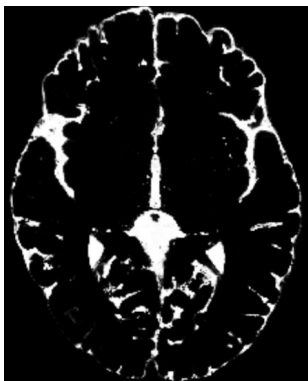
Grey matter



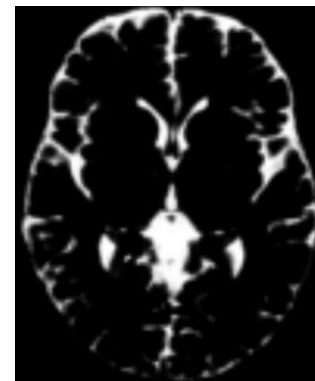
White matter



Cerebrospinal
fluid



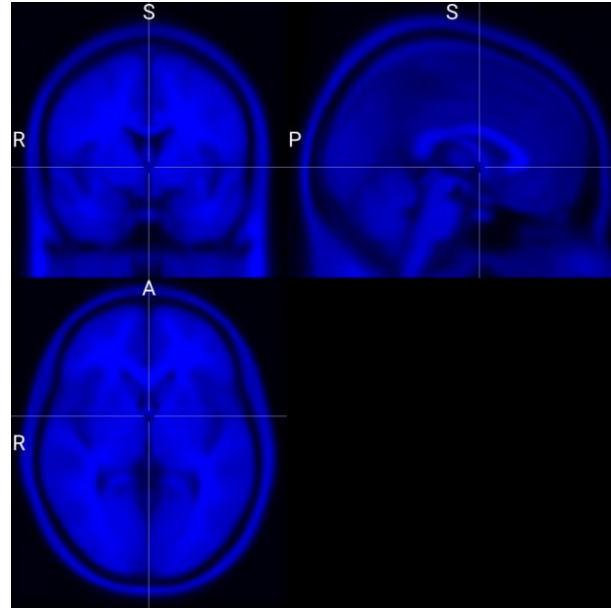
Template tissue probability maps



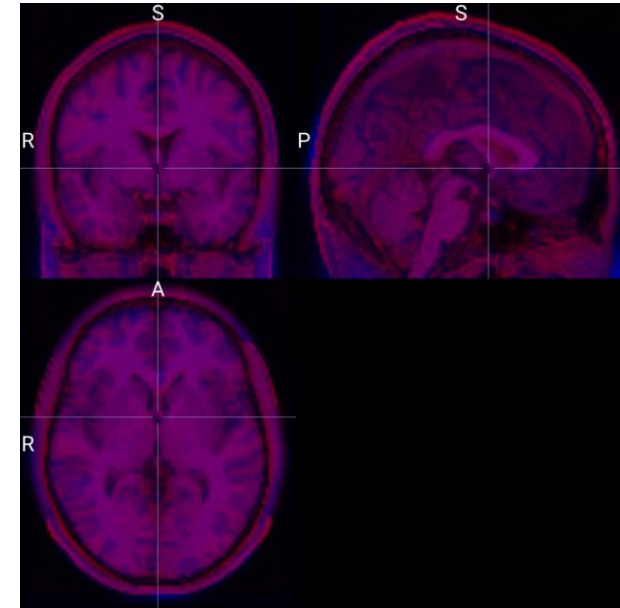
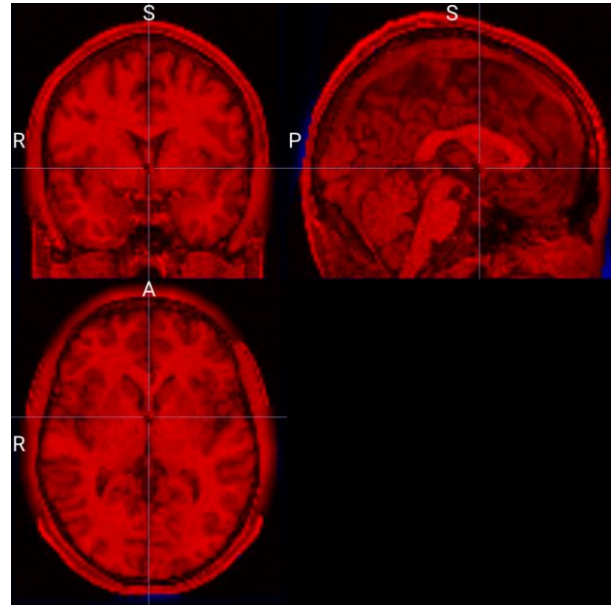
Unified segmentation and normalisation

Confirmation

MNI152 template brain

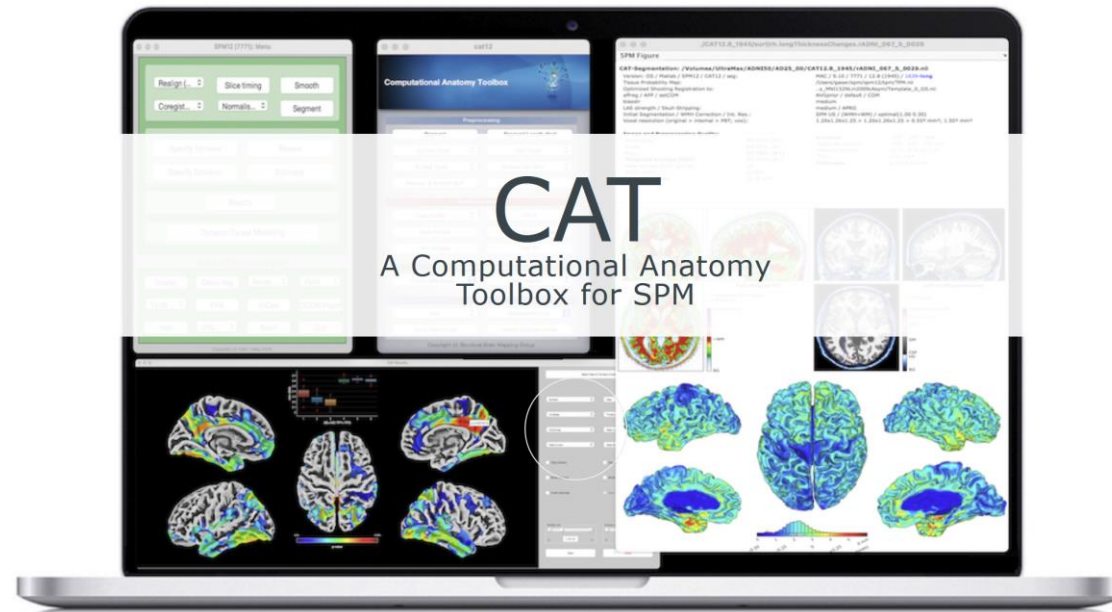


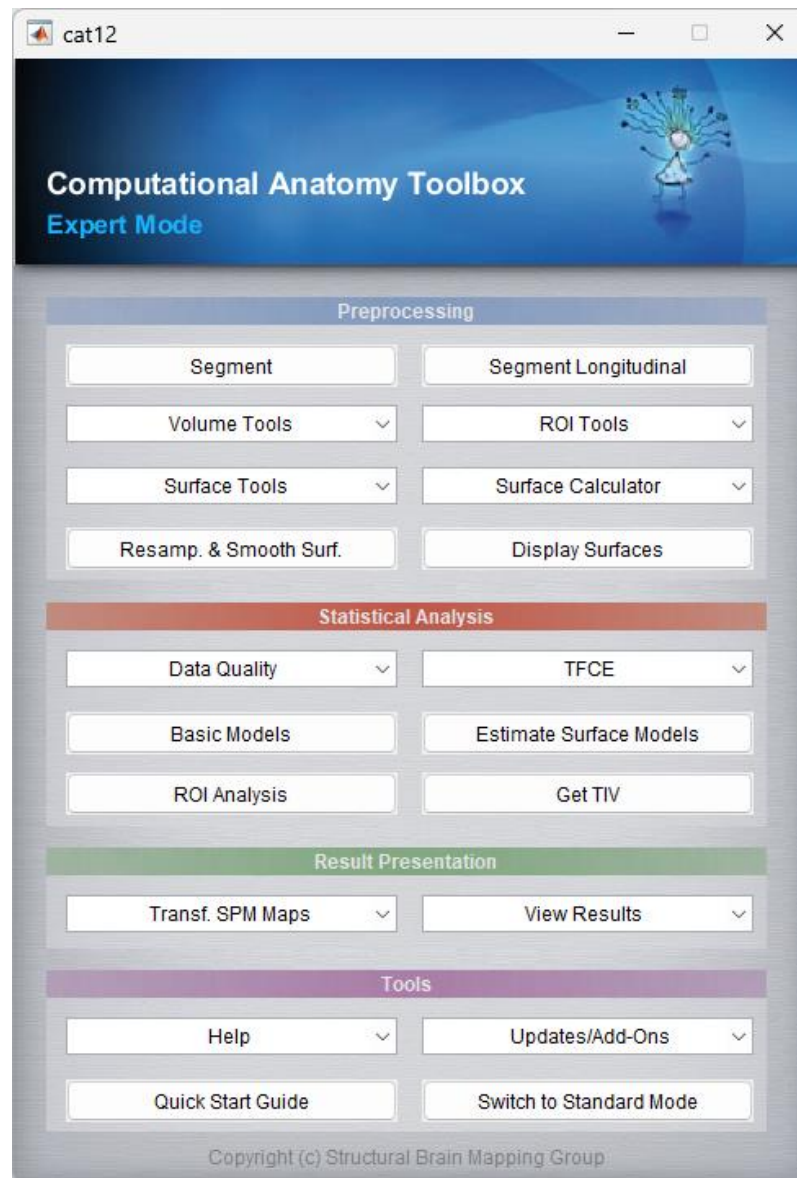
Individual's normalized brain



Voxel-based Morphometry

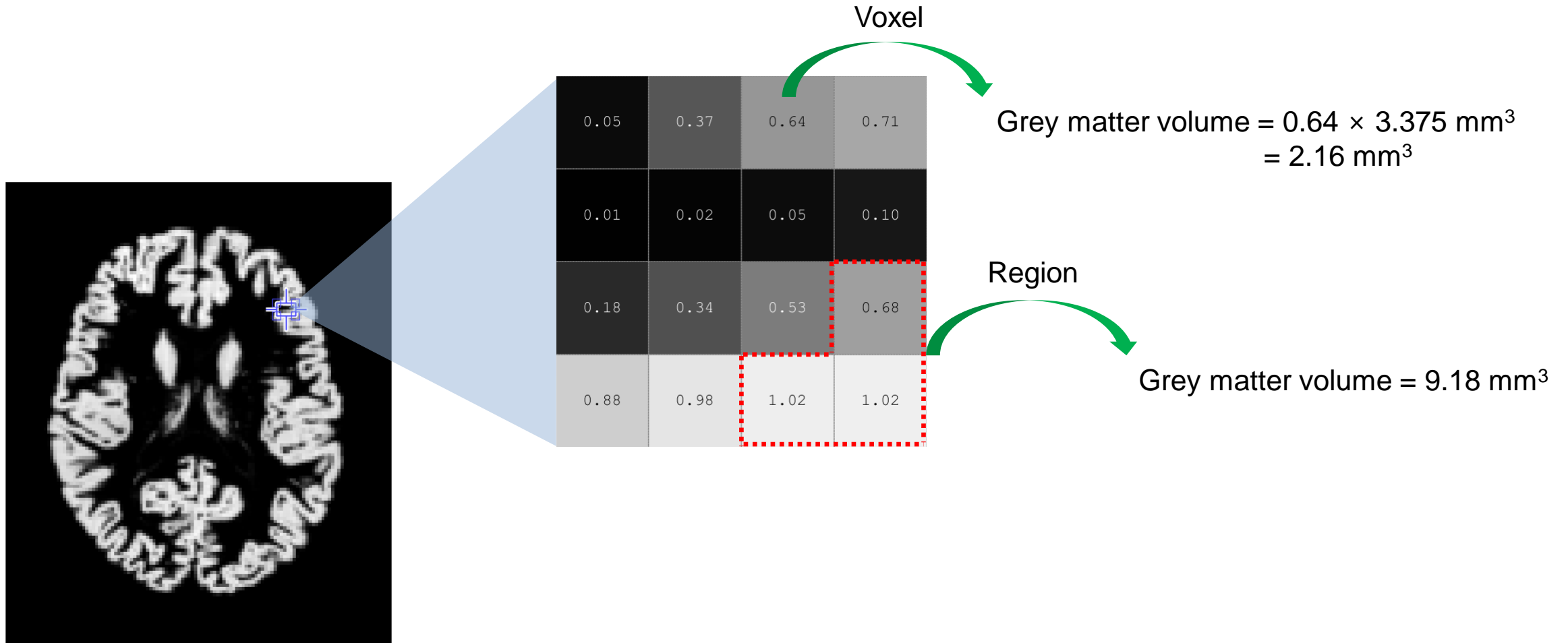
- Without defining boundaries and modelling cortical surfaces
- CAT12 [\[https://neuro-jena.github.io/cat/\]](https://neuro-jena.github.io/cat/)





CAT12 toolbox

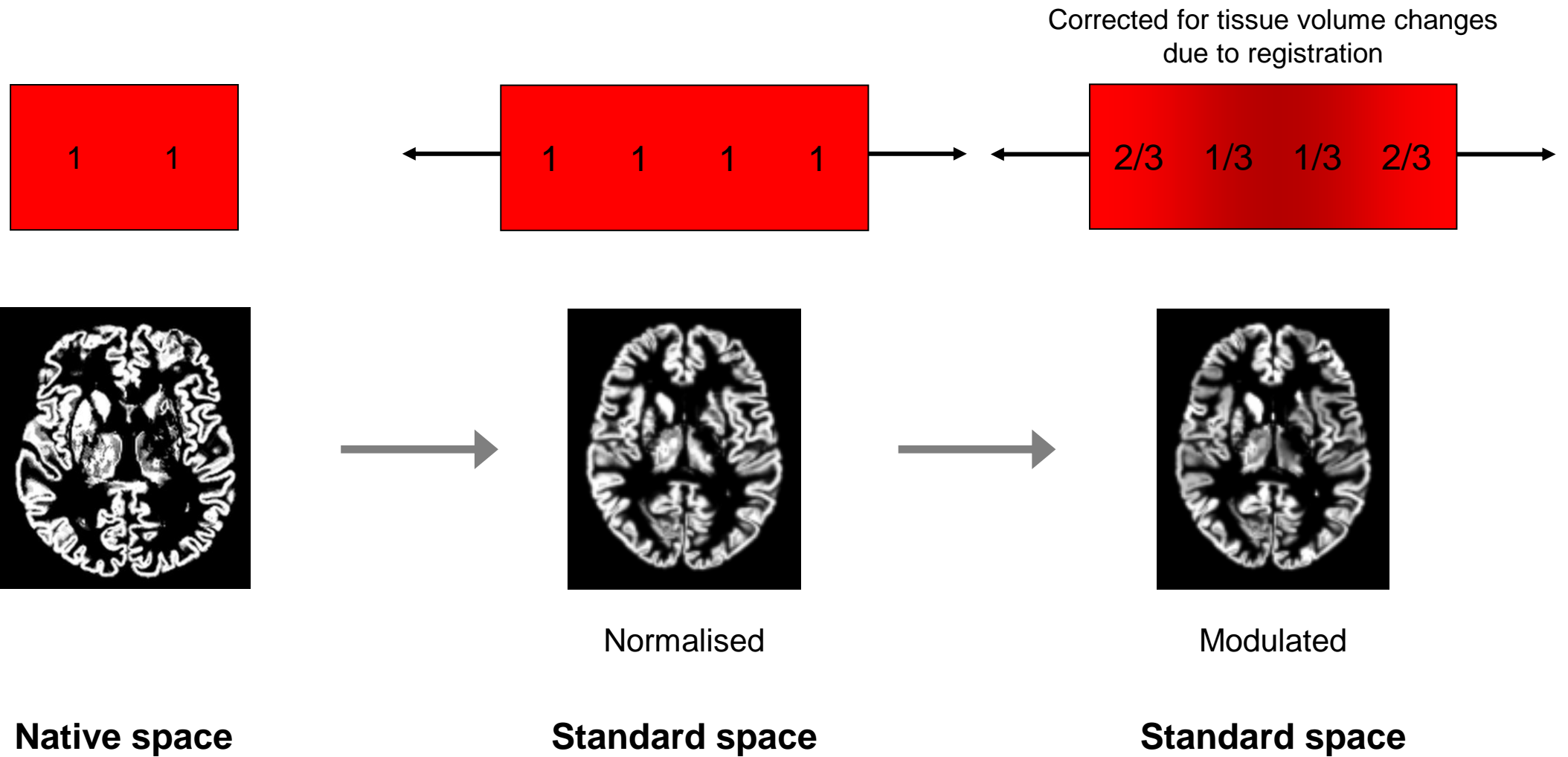
- Grey matter volume
 - Computed by multiplying voxel-wise grey matter probability by voxel volume
 - For a grey matter probability map in the native space or its modulated one in the standard space



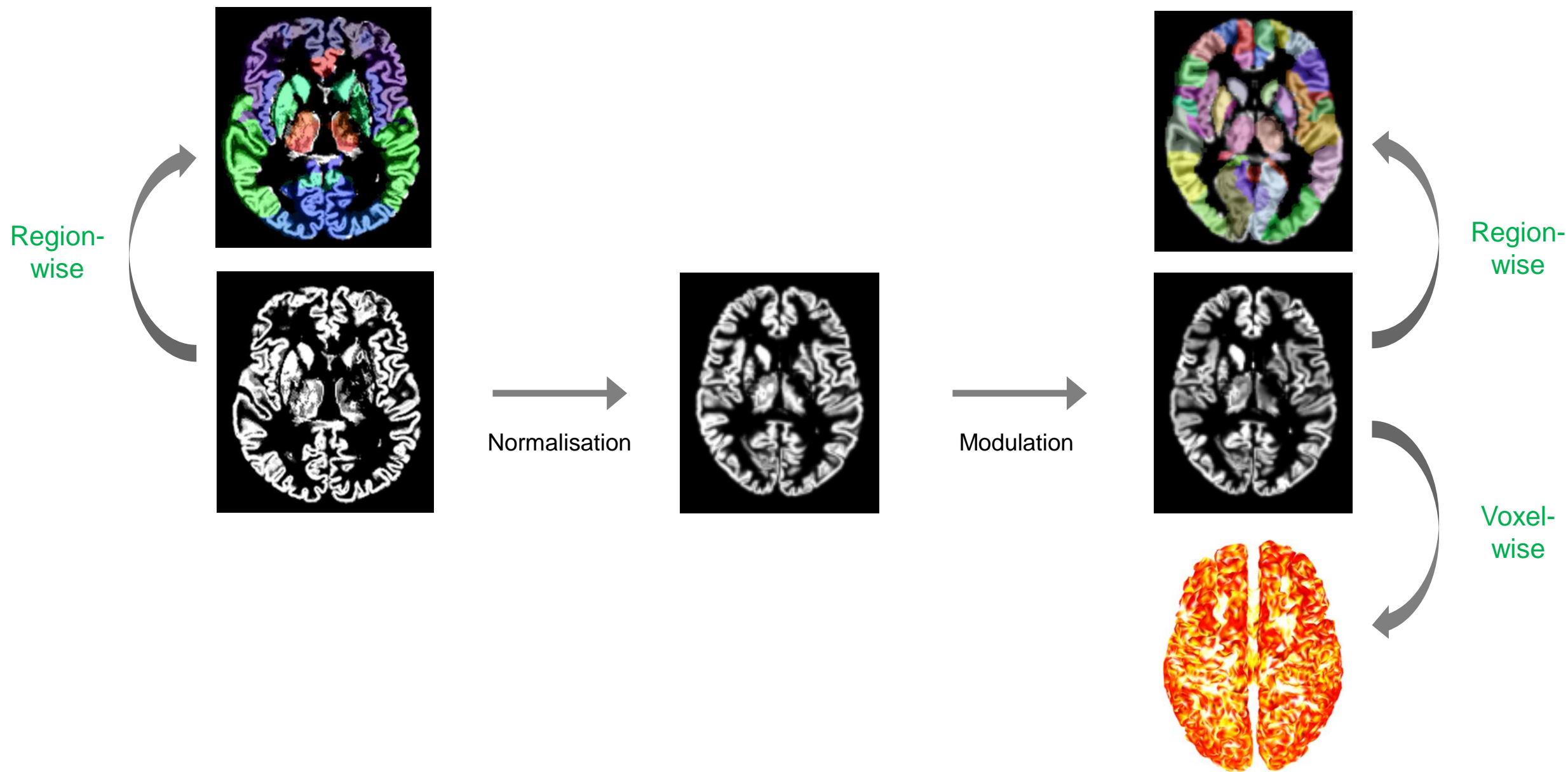
Voxel size: $1.5 \text{ mm} \times 1.5 \text{ mm} \times 1.5 \text{ mm}$

Voxel volume: 3.375 mm^3

Computation of grey matter volume for a voxel or a region



Normalisation and modulation

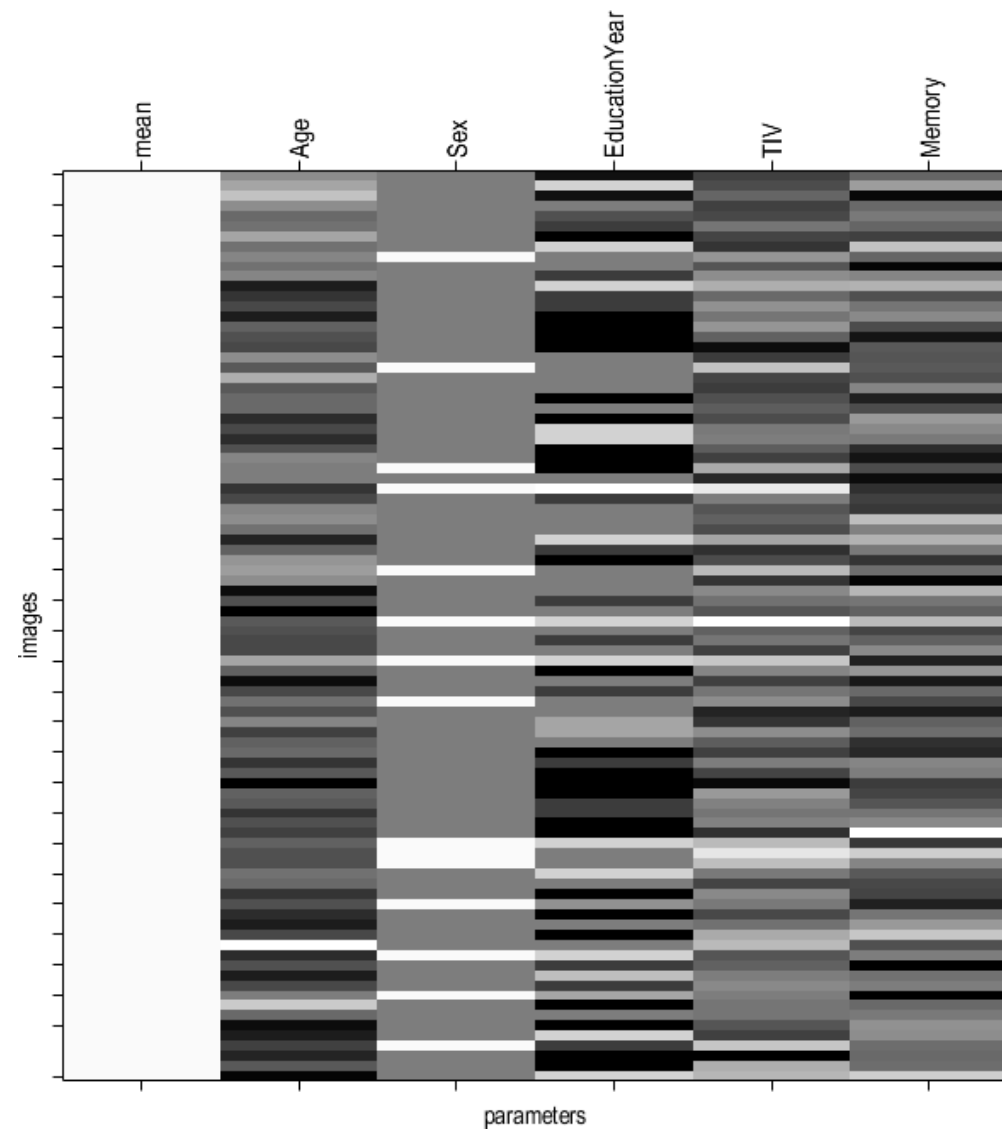


Features of grey matter volume

[Statistical Analysis of sMRI]

- Grey matter volume \sim
Age +
Sex +
Education year +
Total intracranial volume (TIV) +
Memory performance

Design matrix

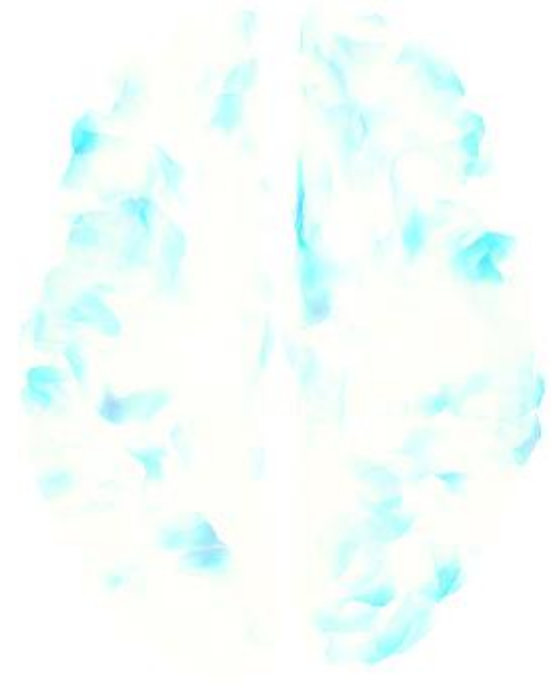


Output

Regression

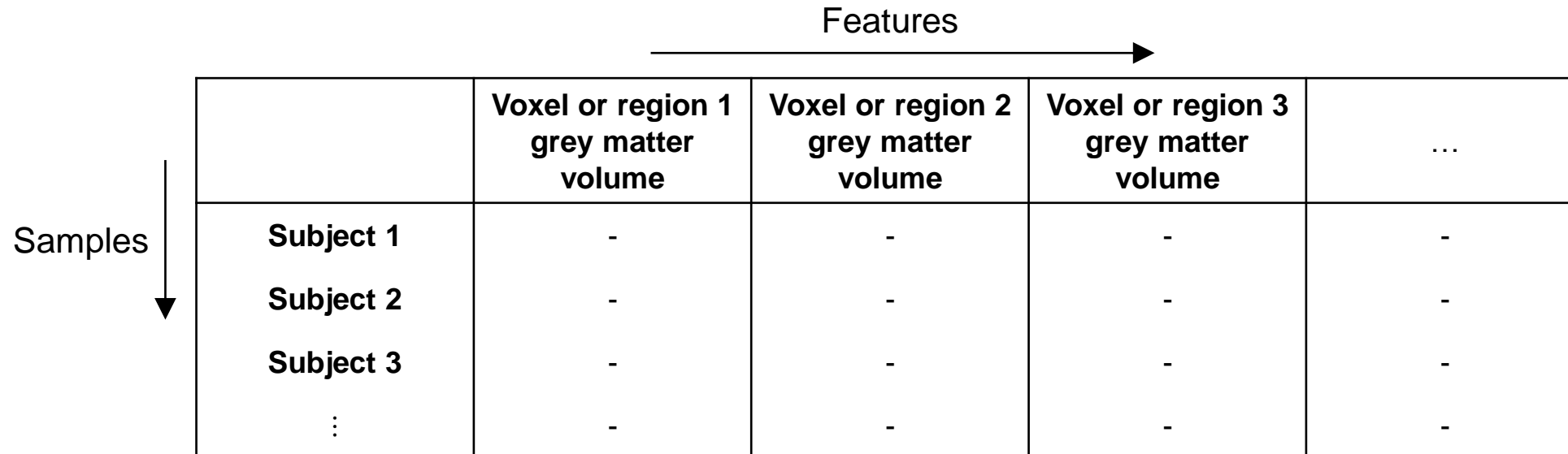


Positive correlaton



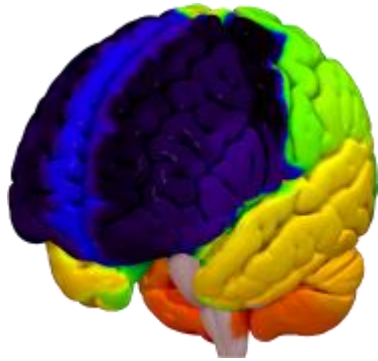
Negative correlation

- Input to machine learning models
 - Table of voxel-wise or region-wise grey matter volume values

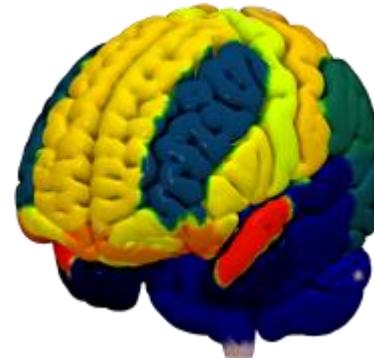


| | | Features → | | | |
|-----------|-----------|--|--|--|-----|
| | | Voxel or region 1 grey matter volume | Voxel or region 2 grey matter volume | Voxel or region 3 grey matter volume | ... |
| Samples ↓ | Subject 1 | - | - | - | - |
| | Subject 2 | - | - | - | - |
| | Subject 3 | - | - | - | - |
| | ⋮ | - | - | - | - |

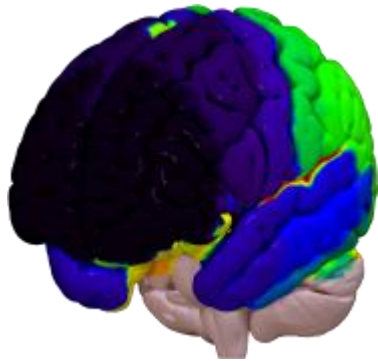
- Grey matter volume map



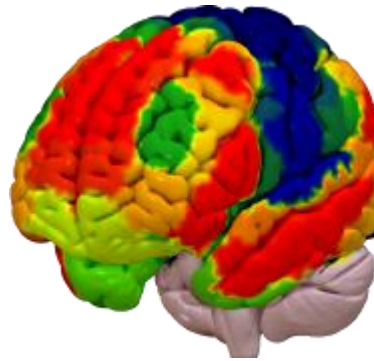
AAL atlas



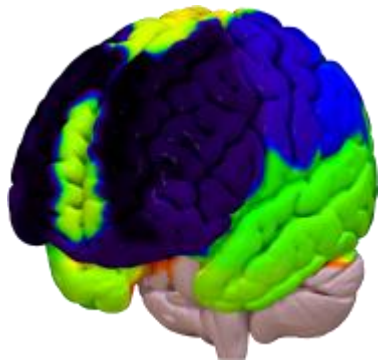
Hammers atlas



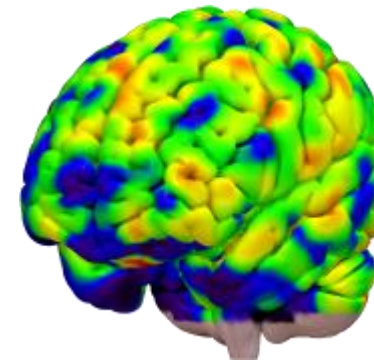
Harvard-Oxford atlas



Yeo atlas

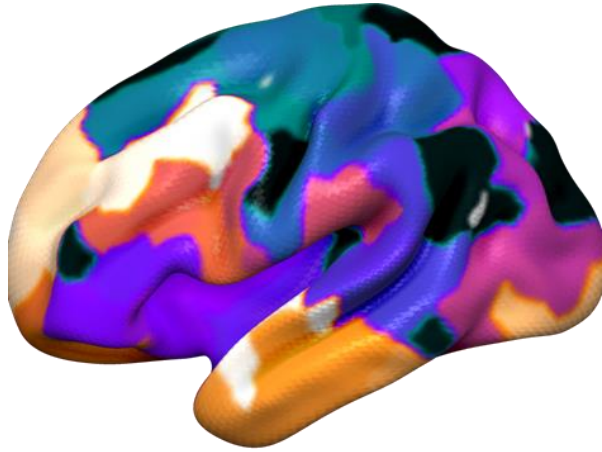


AICHA atlas

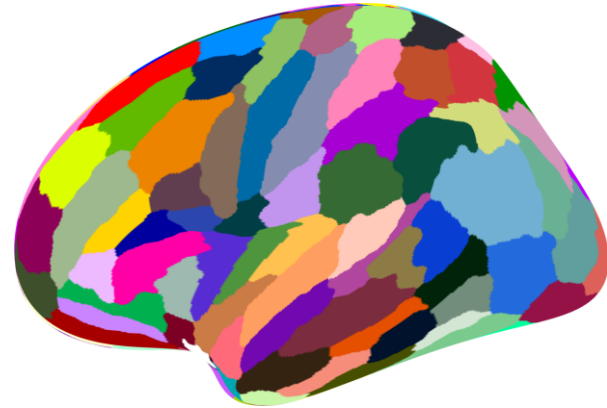


Craddock atlas

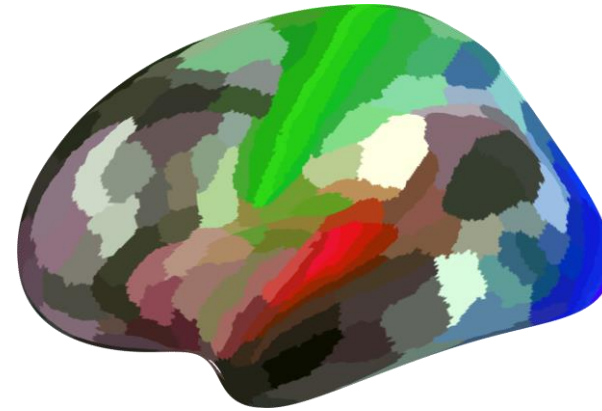
Brain atlases



333 areas
Resting-State Correlations atlas



246 areas
Brainnetome atlas



360 areas
HCP MMP 1.0 atlas

Higher-resolution brain atlases

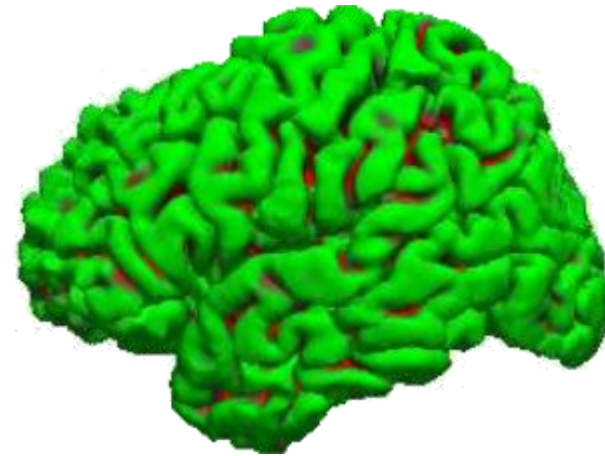
Surface-based morphometry

- Independent of registration and modulation
- Not applicable to subcortical regions
- FreeSurfer [\[https://surfer.nmr.mgh.harvard.edu/\]](https://surfer.nmr.mgh.harvard.edu/)
 - sMRI analysis software of choice for the Human Connectome Project

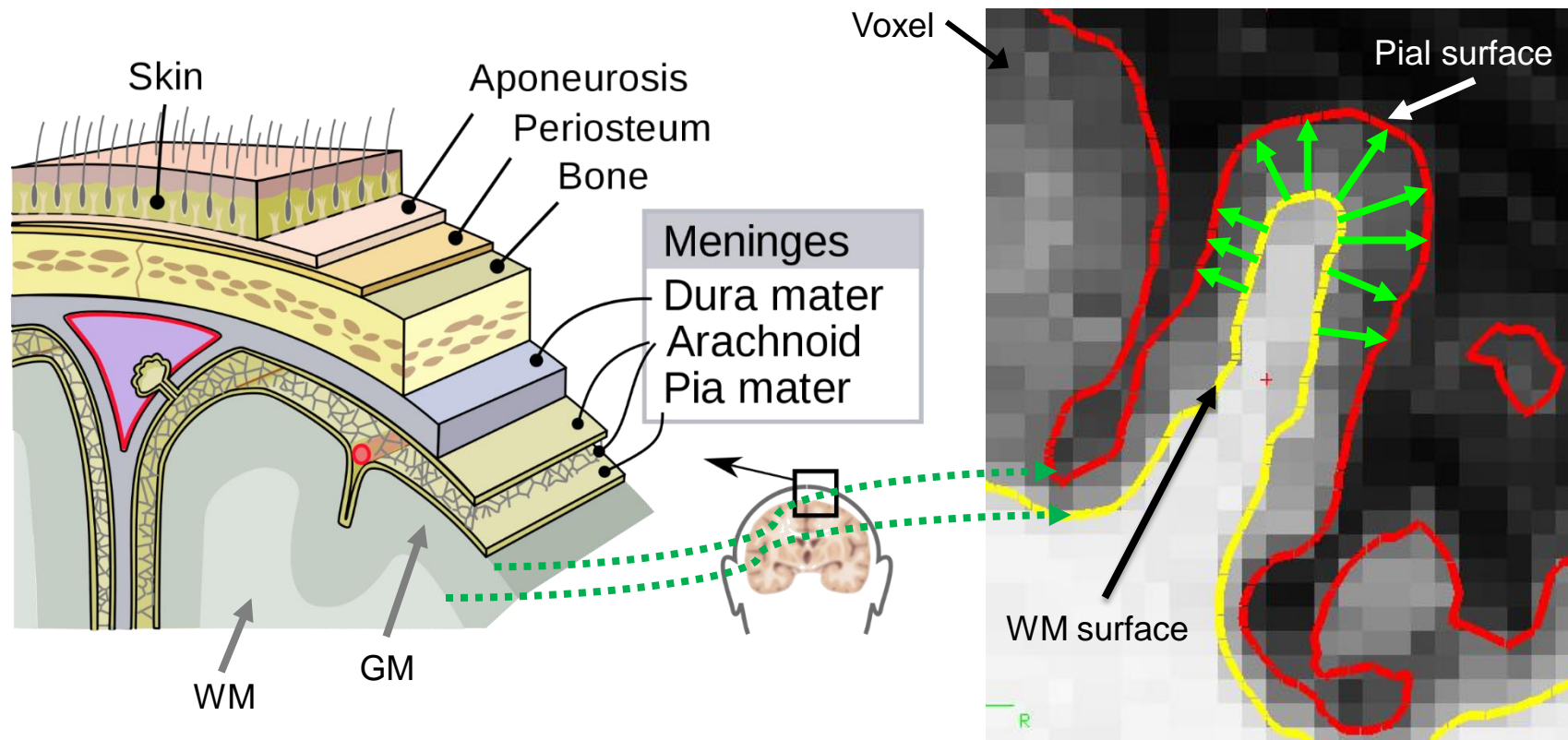
- Surface reconstruction
 - White matter surface: inner cortical boundary between the grey matter and white matter
 - Pial surface: outer cortical boundary between the grey matter and pia mater



White matter surface

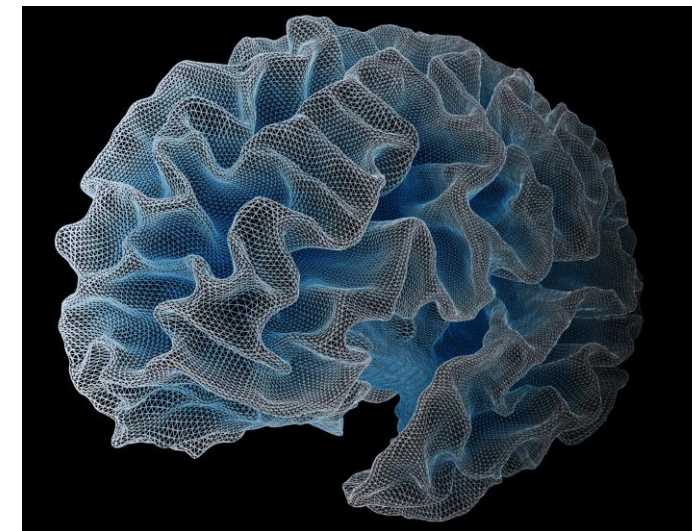
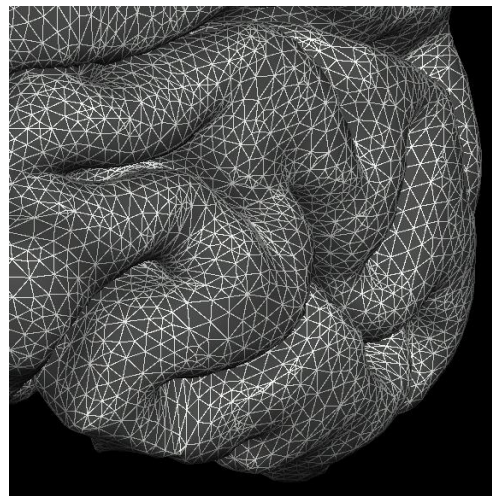
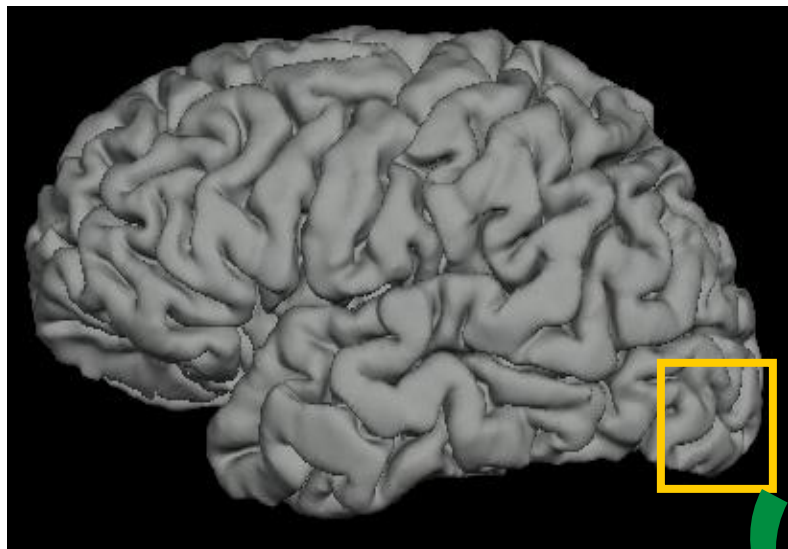


Pial surface



[\[https://www.physio-pedia.com/Meninges\]](https://www.physio-pedia.com/Meninges)

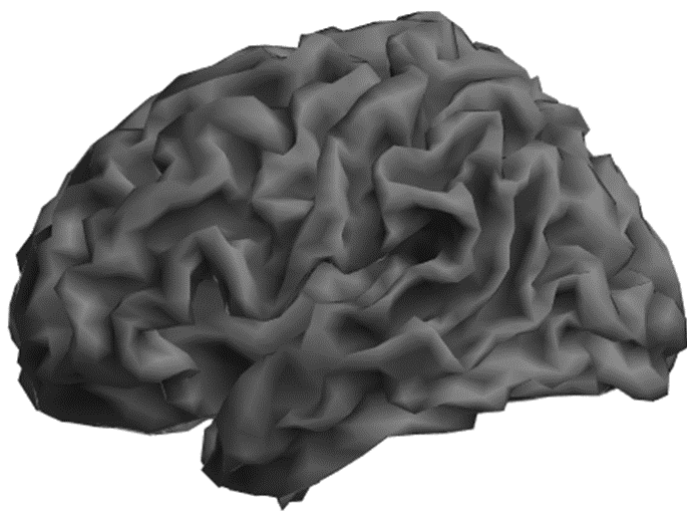
Cortical surfaces beneath cranial meninges



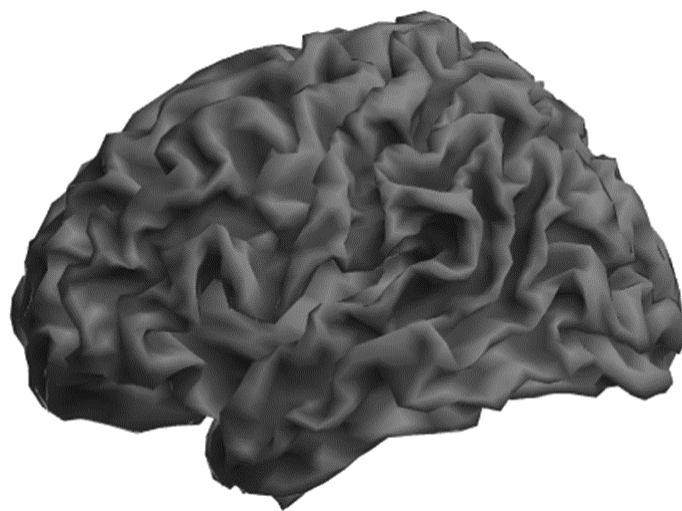
Vertices and faces

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

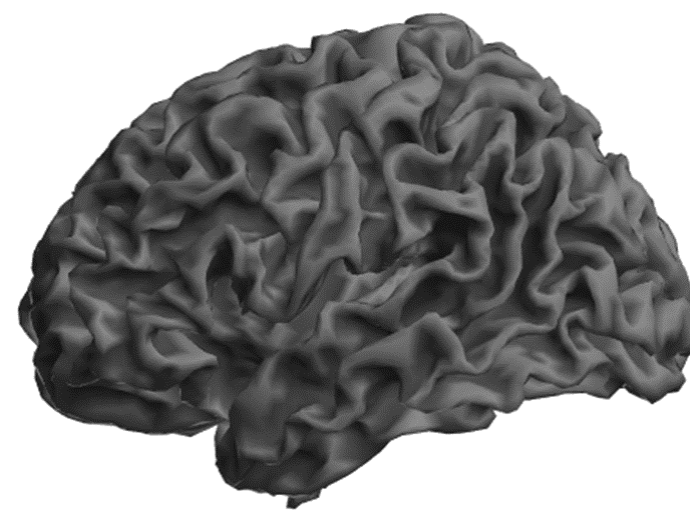
Surface representation of the cerebral cortex



5124 vertices



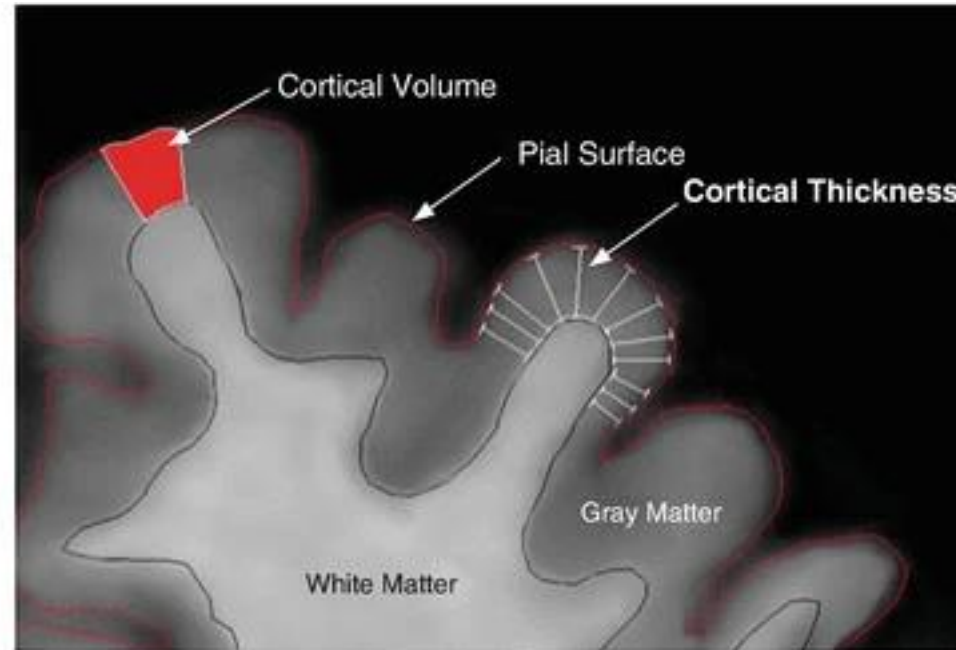
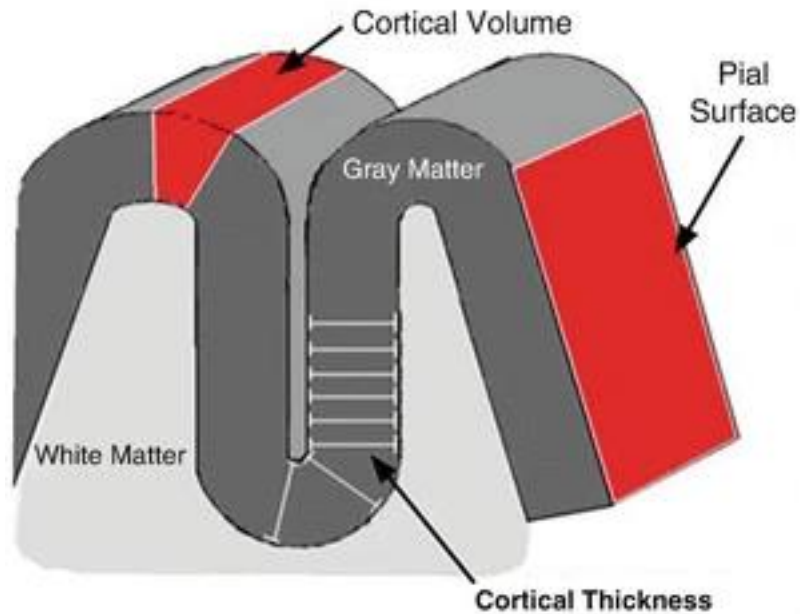
8196 vertices

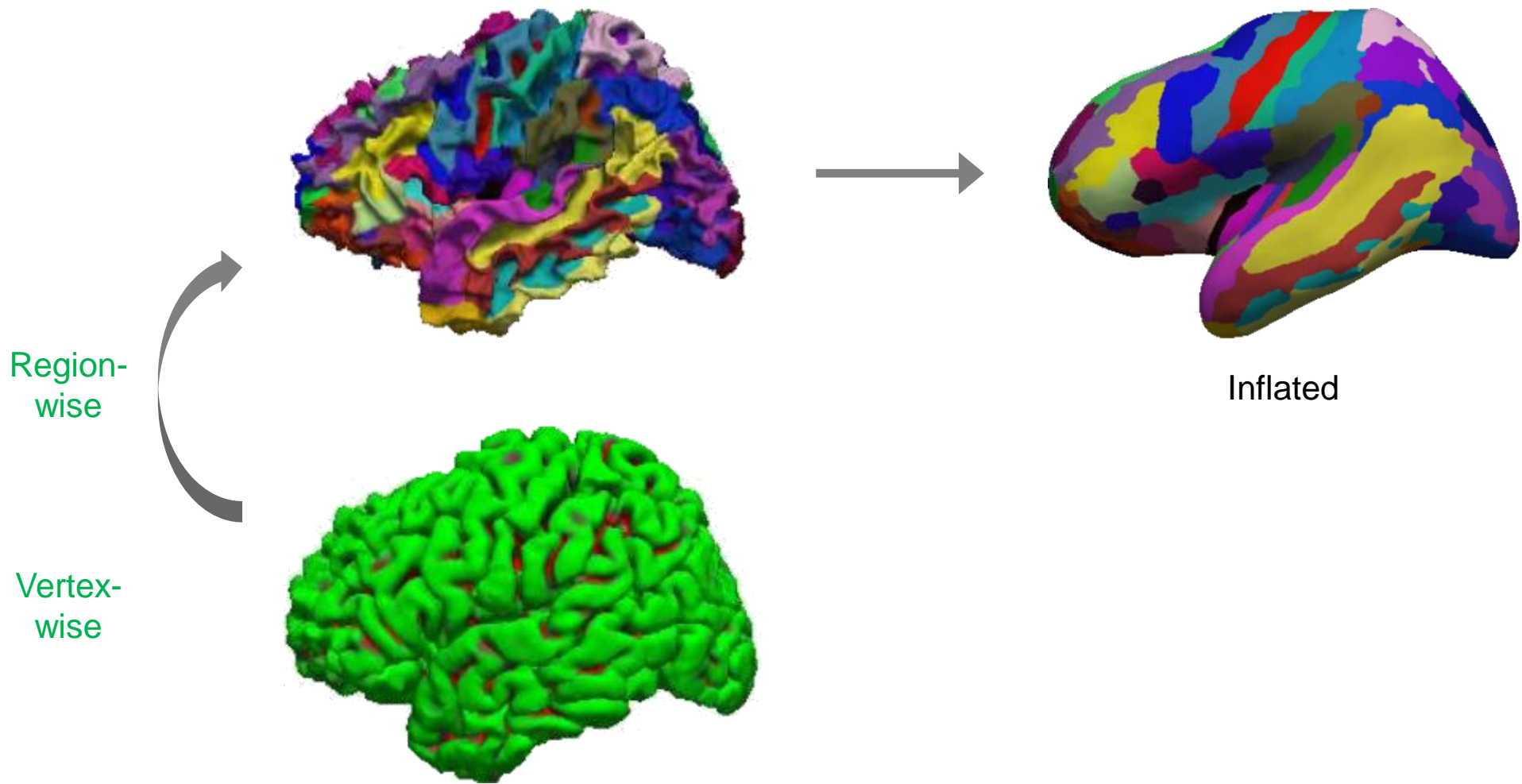


20484 vertices

Surface representation with different numbers of vertices

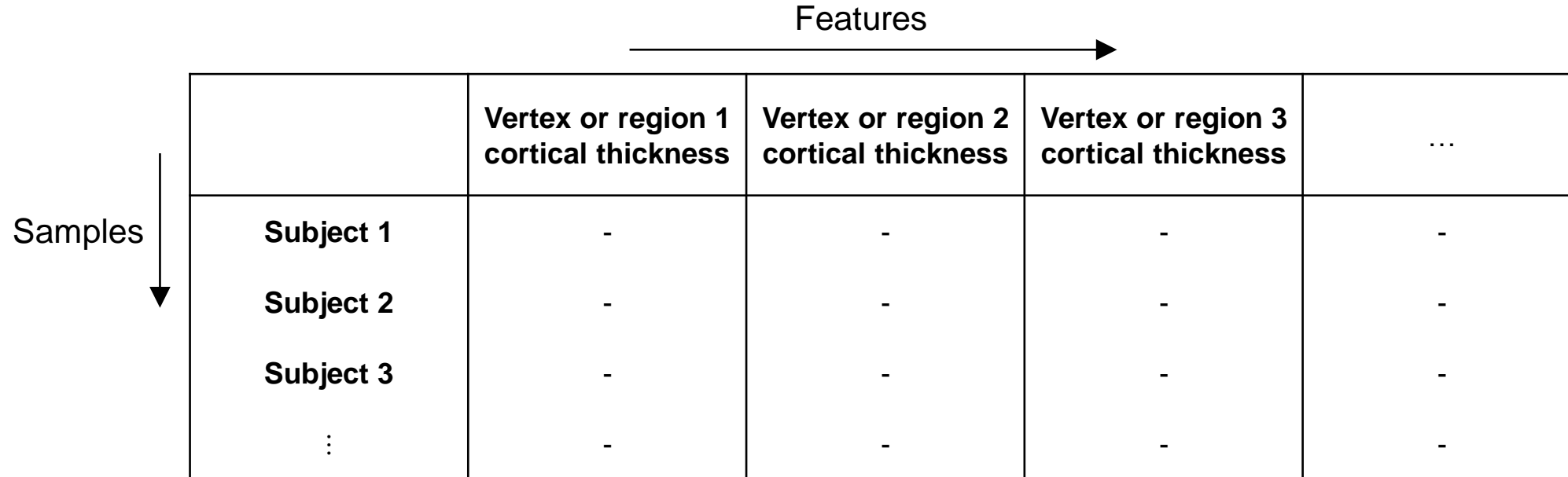
- Cortical thickness
 - Distance between the inner (white matter surface) and outer (pial surface) cortical boundaries





Features of cortical thickness

- Input to machine learning models
 - Table of vertex-wise or region-wise cortical thickness values

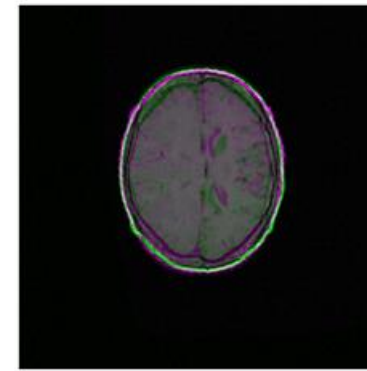
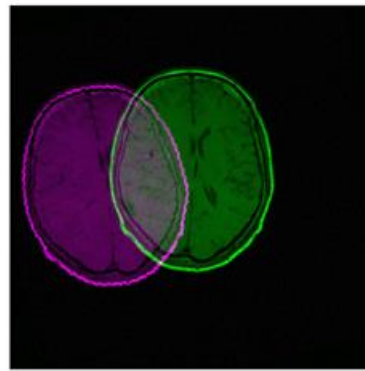


| Features → | | | | |
|------------|--|--|--|-----|
| | Vertex or region 1 cortical thickness | Vertex or region 2 cortical thickness | Vertex or region 3 cortical thickness | ... |
| Samples ↓ | Subject 1 | - | - | - |
| | Subject 2 | - | - | - |
| | Subject 3 | - | - | - |
| | ⋮ | - | - | - |

sMRI as an Individual's Spatial Reference

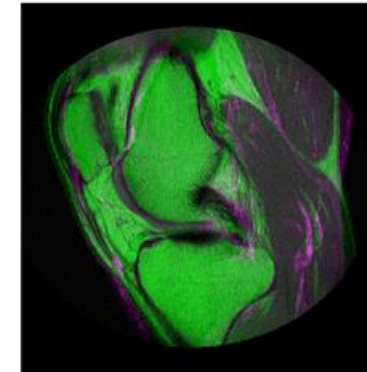
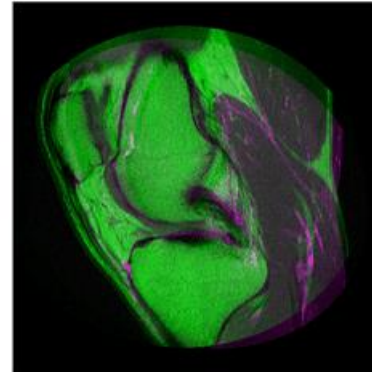
- Anatomical localization of other modalities of MRI
 - Within-subject between-modality registration

Rigid registration
(global shift and rotation)



Within-subject within-modality

Affine registration
(global shift, rotation, scale, and shear)



Within-subject between-modality

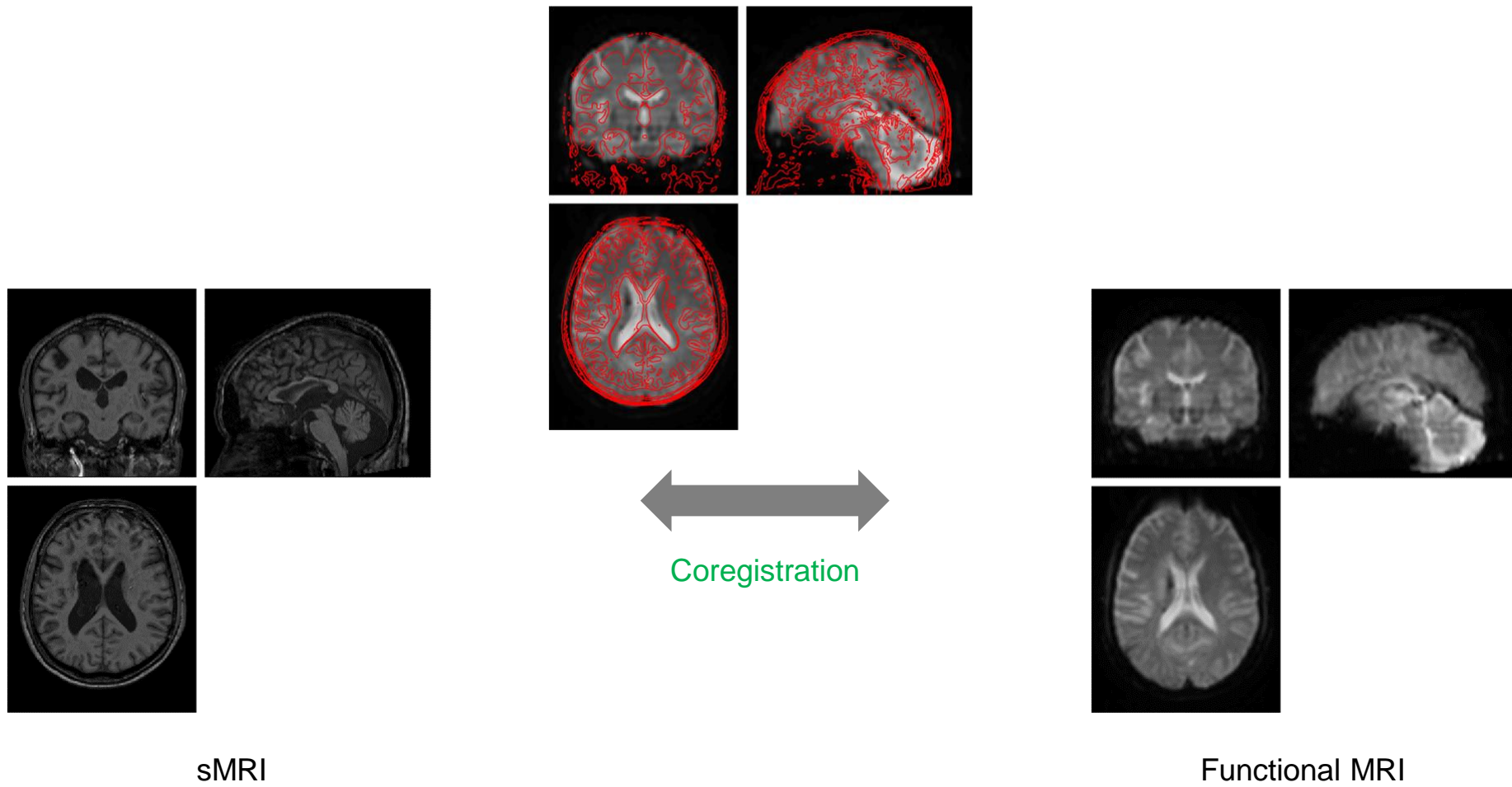
Deformable registration
(local transformations)



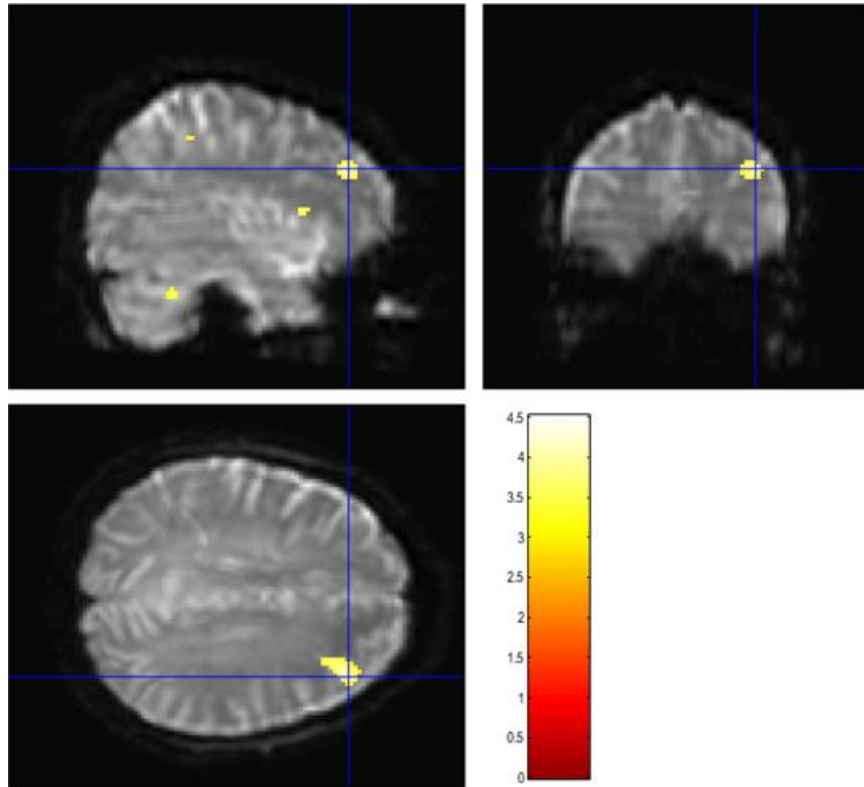
Between-subject

[\[https://kr.mathworks.com/help/medical-imaging/ug/medical-image-registration.html\]](https://kr.mathworks.com/help/medical-imaging/ug/medical-image-registration.html)

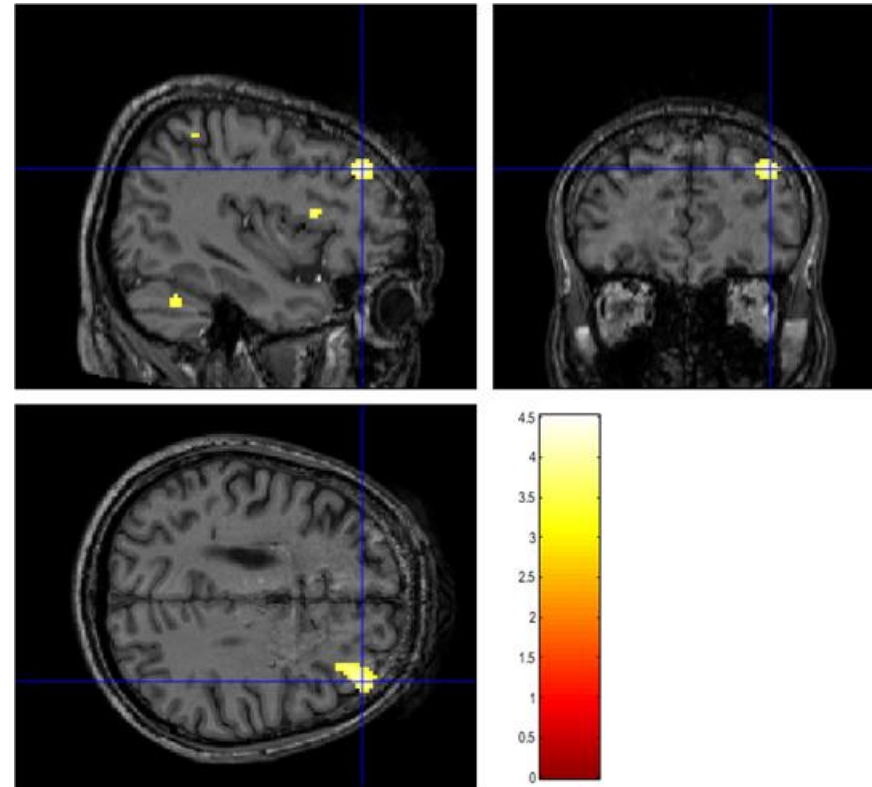
Image registration



Coregistration between sMRI and functional MRI



Brain activity on a functional image



Brain activity on a structural image

Anatomical localization of brain activity