

ALZHEIMER'S ASSOCIATION

AAIC > 23

ALZHEIMER'S ASSOCIATION INTERNATIONAL CONFERENCE®

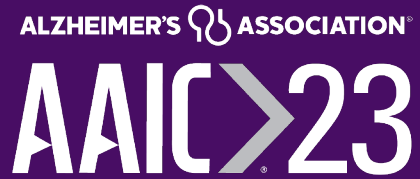
JULY 16-20 > AMSTERDAM, NETHERLANDS, AND ONLINE

ISTAART Neuroimaging PIA THE BASICS OF NEUROIMAGING SEMINAR SERIES



ISTAART Neuroimaging PIA

The Basics of Neuroimaging Series



BASICS OF NEUROIMAGING

DATA STRUCTURE AND FORMATS

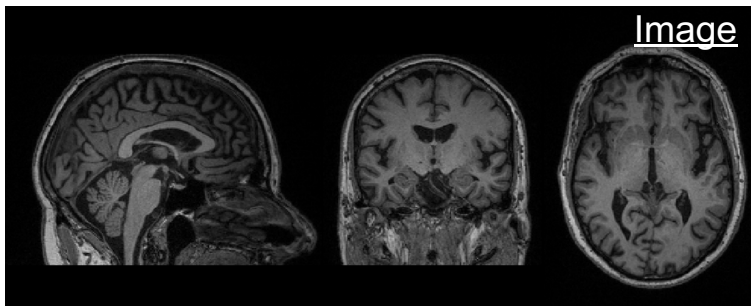
DR LUDOVCA GRIFFANTI
UNIVERSITY OF OXFORD, UK



@ludogriffanti

By the end of this session you should be able to:

- Describe the main **properties of medical images**
- Identify the main **steps of a neuroimaging study**
- Understand **how neuroimaging data are visualized** at different steps of the analysis pipeline



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Image formats



“Raw” Scanner File Format
Example: DICOM

DICOM to NIFTI conversion (e.g. dcm2nii)



Analysis File Format
Example: NIFTI (Neuroimaging Informatics Technology Initiative)

Image = cube of numbers

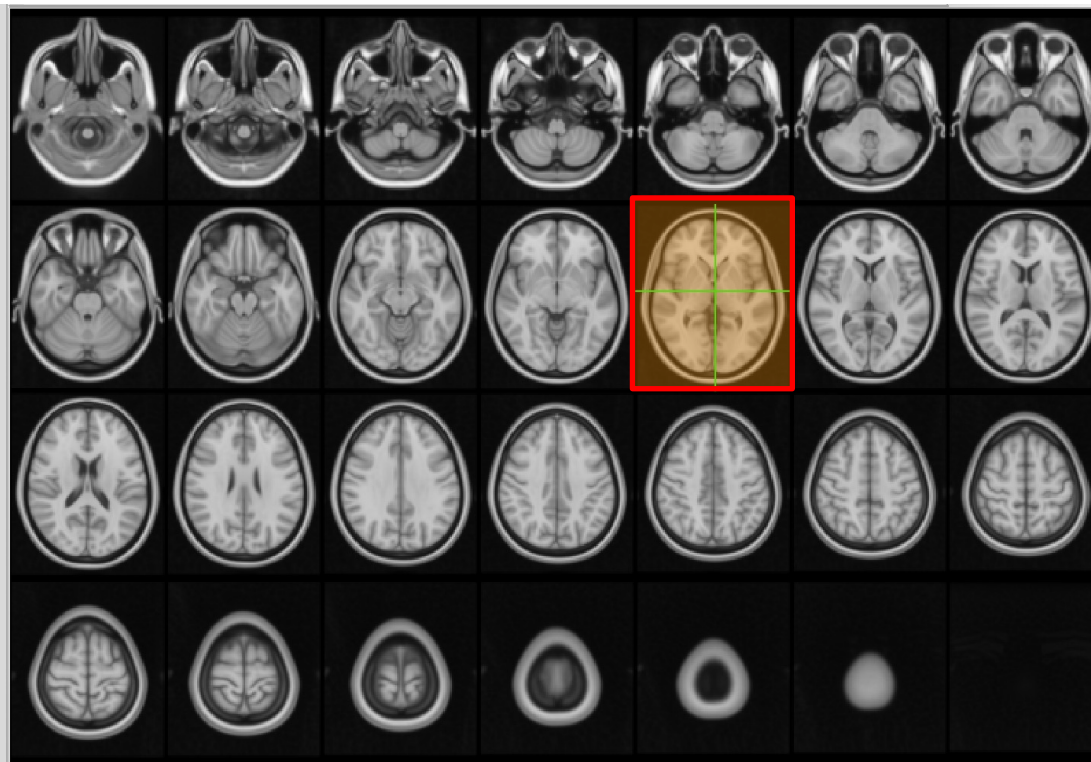
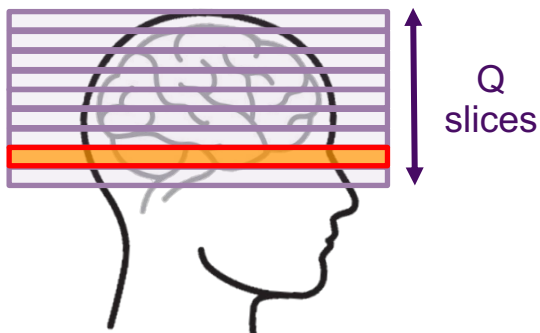
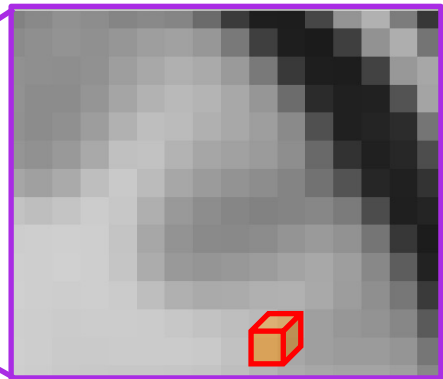
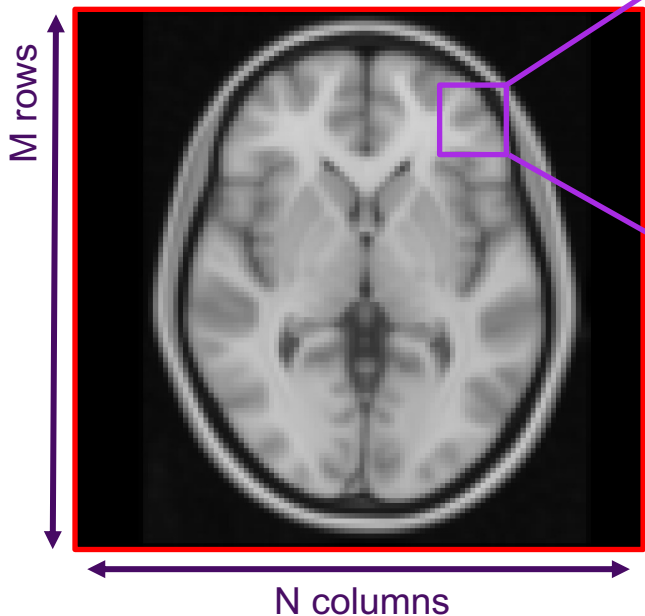
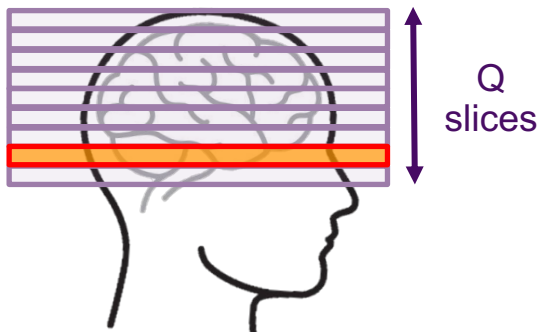


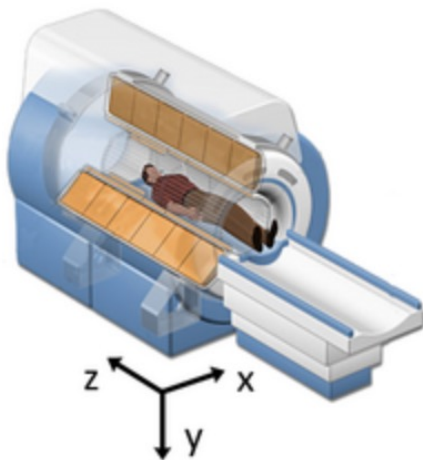
Image = cube of numbers



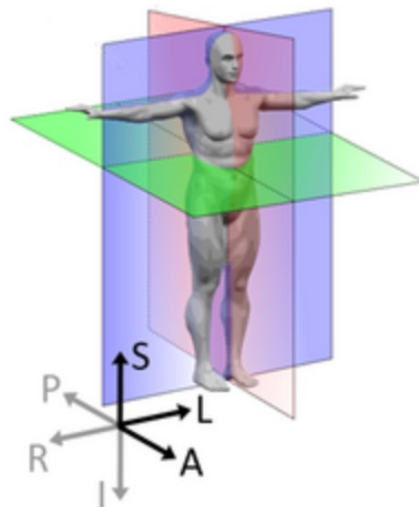
VOXEL (volumetric pixel)

Image resolution =
voxel size in mm

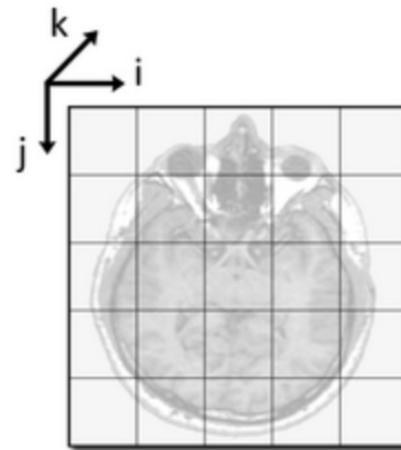
Coordinate systems



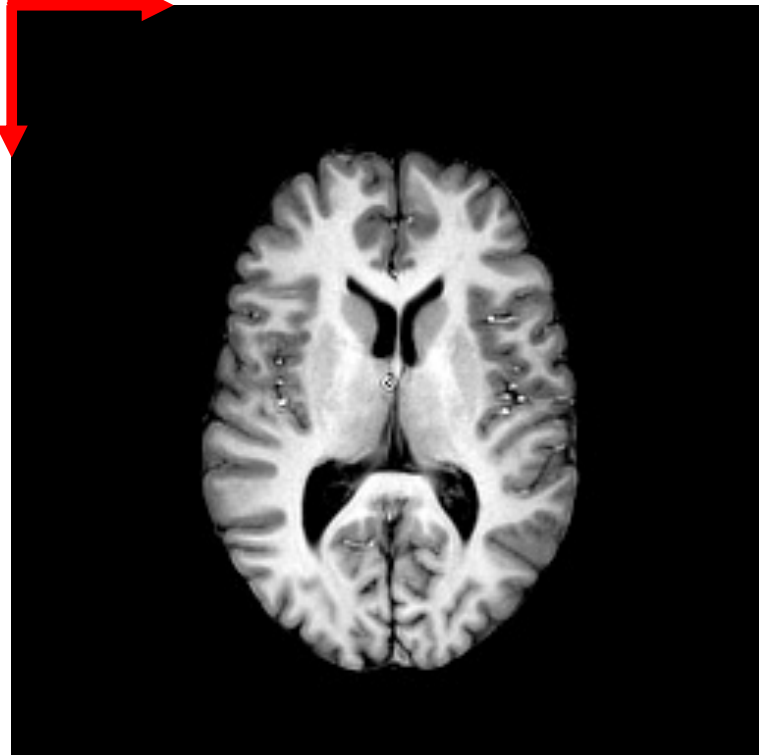
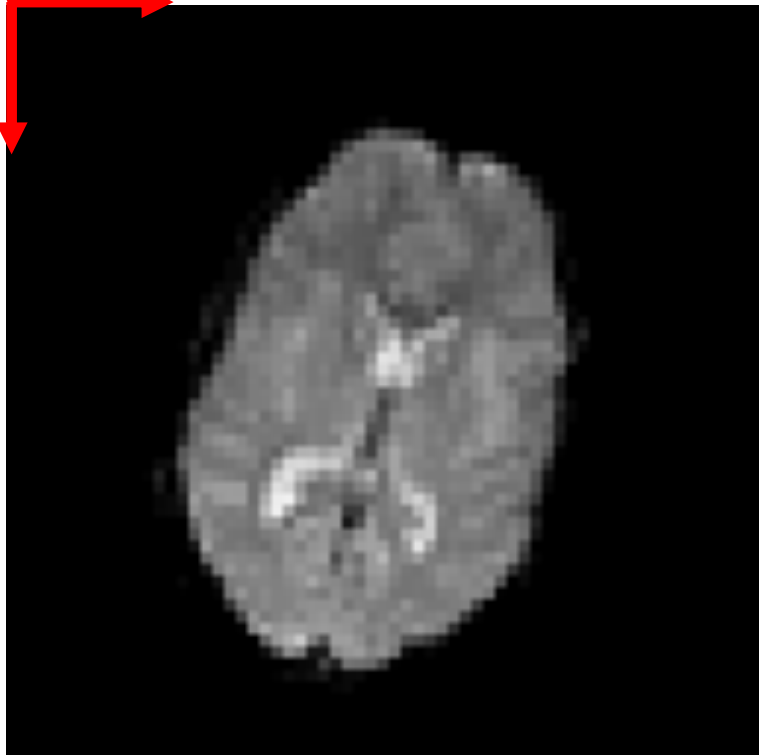
Scanner coordinates

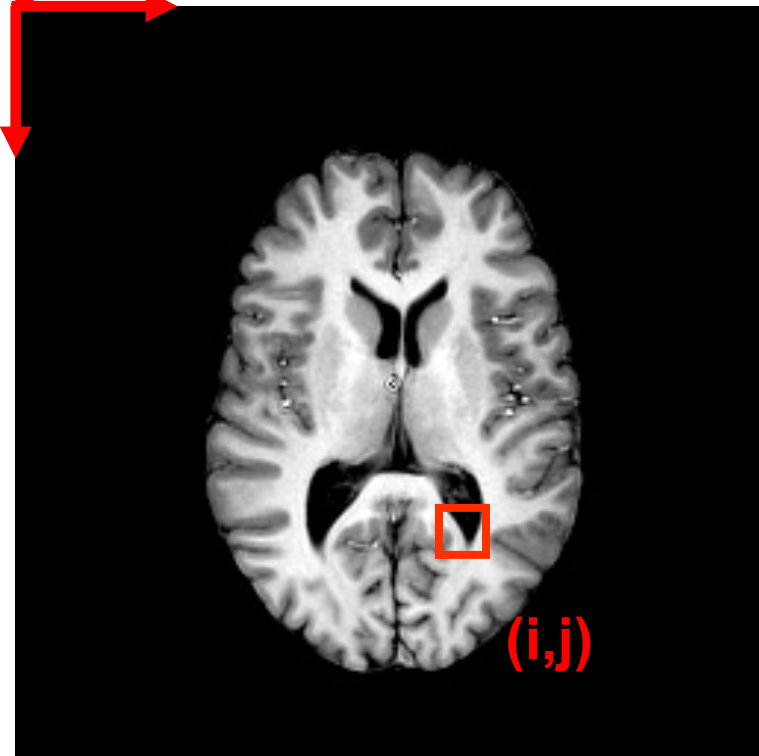
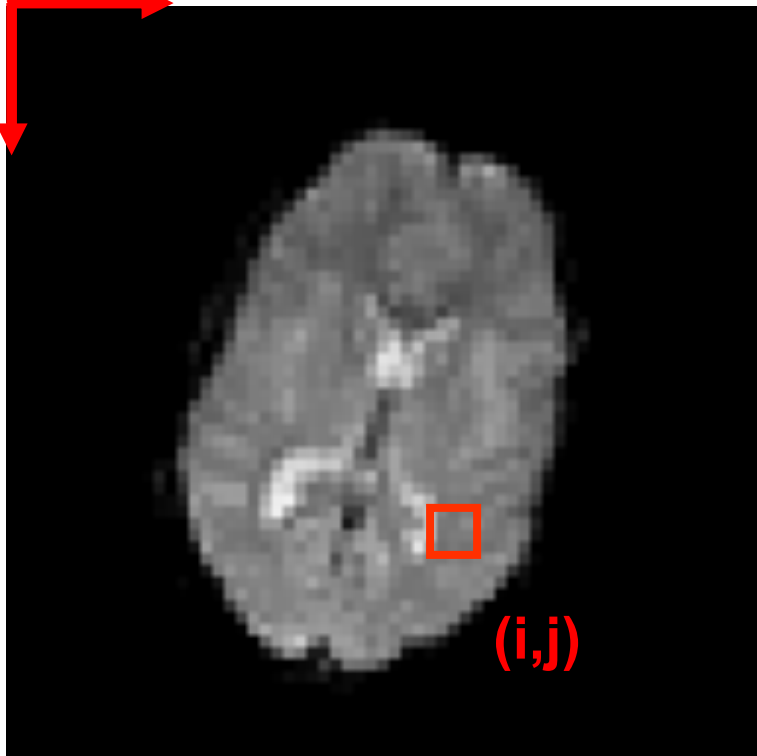


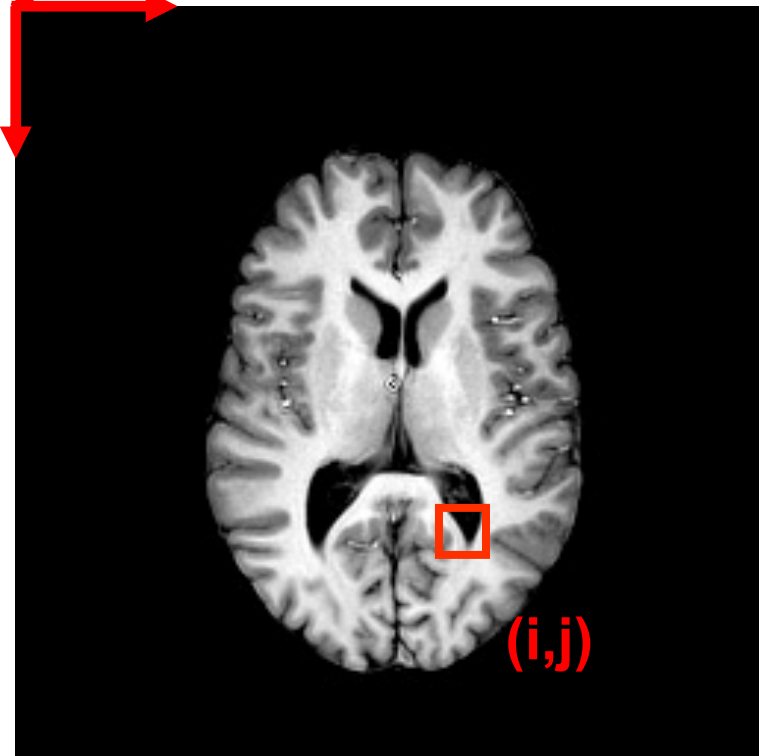
Subject coordinates



Voxel coordinates







- **Registration:** the process of aligning images so that the same voxel in the image corresponds to the same anatomical location in the brain. (i.e. finding a one-to-one map between all points in one image and another)
- **Terminology** varies depending on software and type of transformation applied to the images (Synonyms: coregistration, alignment, normalization)



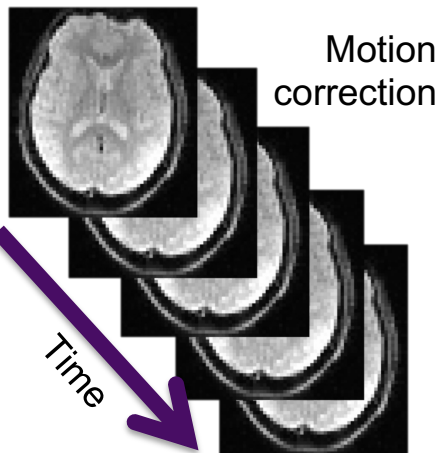
- Input Image
- Moving Image
- Source Image
- Deformed Image

- Reference image
- Stationary Image
- Target Image
- Fixed Image

Within-subject & session

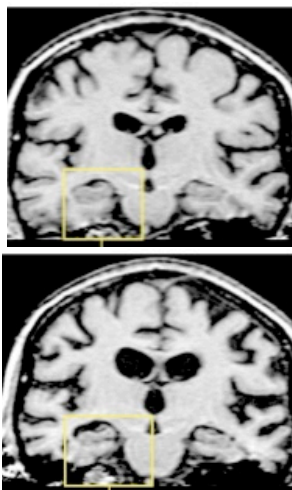


Between-modalities



Motion correction

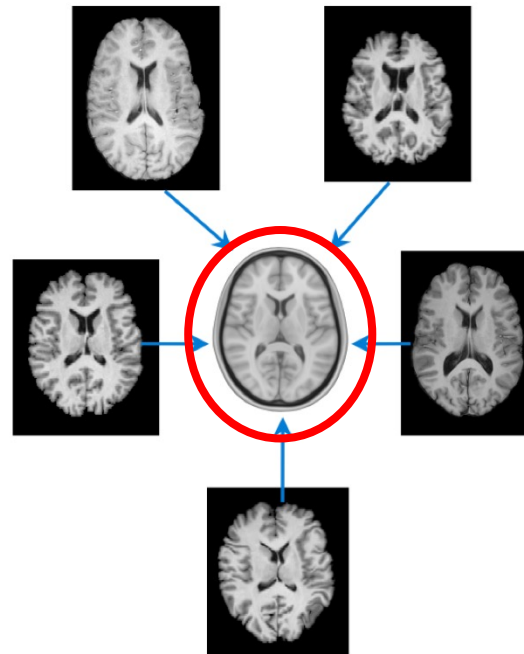
Within-subject, between sessions



Scheltens et al., 2002

Longitudinal data, change over time

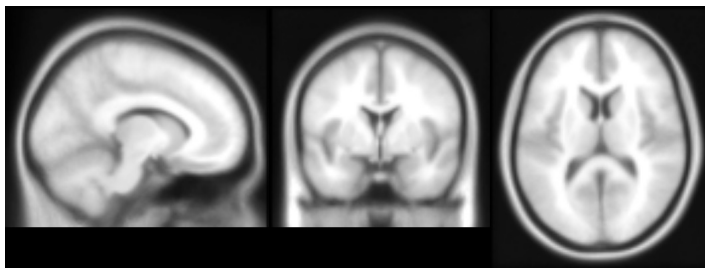
Between-subjects



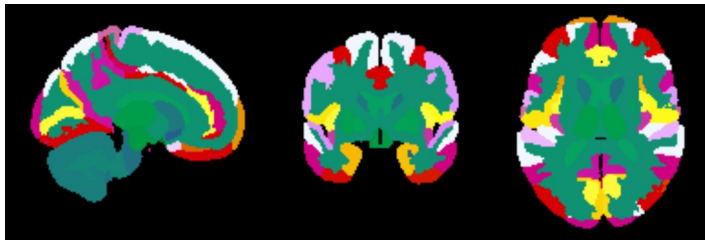
Template / Standard space = "average brain" used as reference

MNI152 Template Space

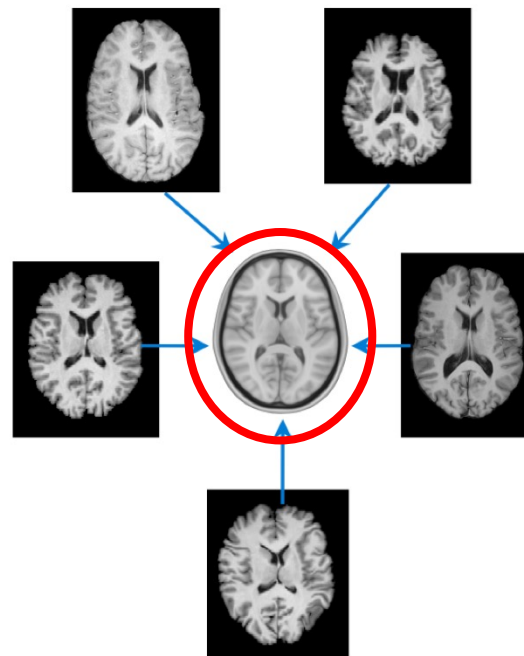
“Average brain”



Atlases



Between-subjects



Template / Standard space =
“average brain” used as reference

NEUROIMAGING DATA ANALYSIS:
A GENERIC BLUEPRINT

1. Data acquisition

Aim: obtain good quality and consistent data

Trade-offs often necessary (e.g. time vs resolution). Optimize protocol for research aim.

2. Data preprocessing

Aim: Reduce noise and prepare data for further analyses

Some steps are common across modalities (e.g. brain extraction, registration), others are modality-specific (e.g. motion correction, distortion correction). Requires careful checking.

3. Single-subject analysis

Aim: Obtain measure of interest for each subject (often an image)

Modality-specific. (Examples: tissue-type segmentation, fractional anisotropy map)

4. Group-level analysis

Aim: Compare single-subject results across groups

Common step across modality (the difference is the input). Usually happens after all images have been aligned (*registered*) to a reference image (*template*)

5. Statistical inference

Aim: test reliability of results and generalizability to the general population

Common step across modalities

1. Data acquisition

1.b Data organization

```
my_dataset/
├── participants.tsv
├── sub-01/
│   ├── anat/
│   │   └── sub-01_T1w.nii.gz
│   ├── func/
│   │   ├── sub-01_task-rest_bold.nii.gz
│   │   └── sub-01_task-rest_bold.json
│   └── dwi/
│       ├── sub-01_dwi.nii.gz
│       ├── sub-01_dwi.json
│       ├── sub-01_dwi.bval
│       └── sub-01_dwi.bvec
├── sub-02/
└── sub-03/
```

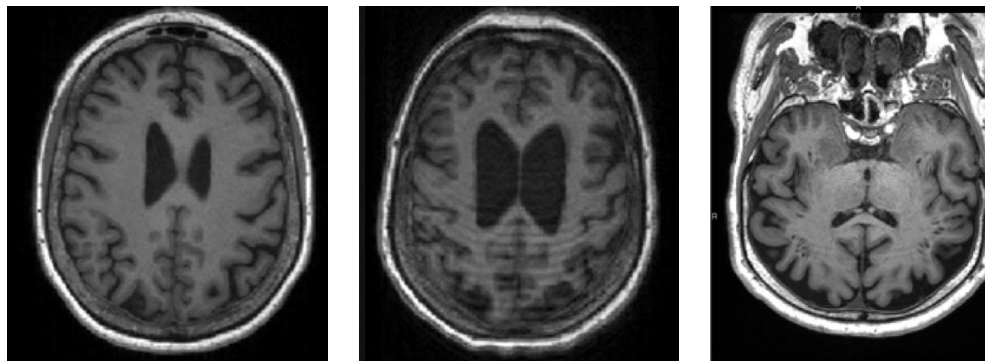
- Neuroimaging experiments usually generate multiple **images and non-imaging data**.
- So far there is **no consensus how to organize and share data** obtained in neuroimaging experiments
- **BIDS** is a framework for organizing data. Standardizes **file names and folders hierarchy** organization and dataset description.



<https://bids.neuroimaging.io>

1. Data acquisition

2. Data preprocessing



MRIQC: group anatomical report

Summary

- Date and time: 2017-02-05, 12:27.
- MRIQC version: 0.9.0.rvc2.

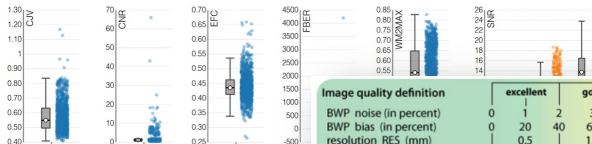


Image quality definition

	excellent	good	satisfactory	sufficient	critical	unacceptable /
BWP noise (in percent)	0	1	2	3	4	5
BWP bias (in percent)	0	20	40	60	80	100
resolution RES (mm)	0.5	1.0	1.5	2.0	2.5	3.0
Quality ratings	100	95	90	85	80	75
percentaged rating points (rps)	100	95	90	85	80	75
linear rating scale	0.5	1	1.5	2	2.5	3
nominal numbers	1+	1-	2+	2-	3+	3-
nominal letters	A+	A-	B+	B-	C+	C-
description	excellent	good	satisfactory	sufficient	critical	unacceptable /

Volume-to-volume motion

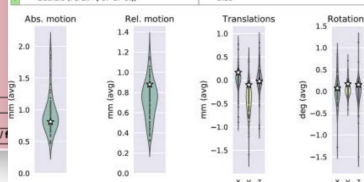
Average abs. motion (mm)	0.81
Average rel. motion (mm)	0.88
Average x translation (mm)	0.17
Average y translation (mm)	-0.10
Average z translation (mm)	-0.02
Average x rotation (deg)	0.07
Average y rotation (deg)	0.17
Average z rotation (deg)	0.15

Within-volume motion

Avg std x translation (mm)	0.02
Avg std y translation (mm)	0.11
Avg std z translation (mm)	0.04
Avg std x rotation (deg)	0.05
Avg std y rotation (deg)	0.05
Avg std z rotation (deg)	0.06

Outliers

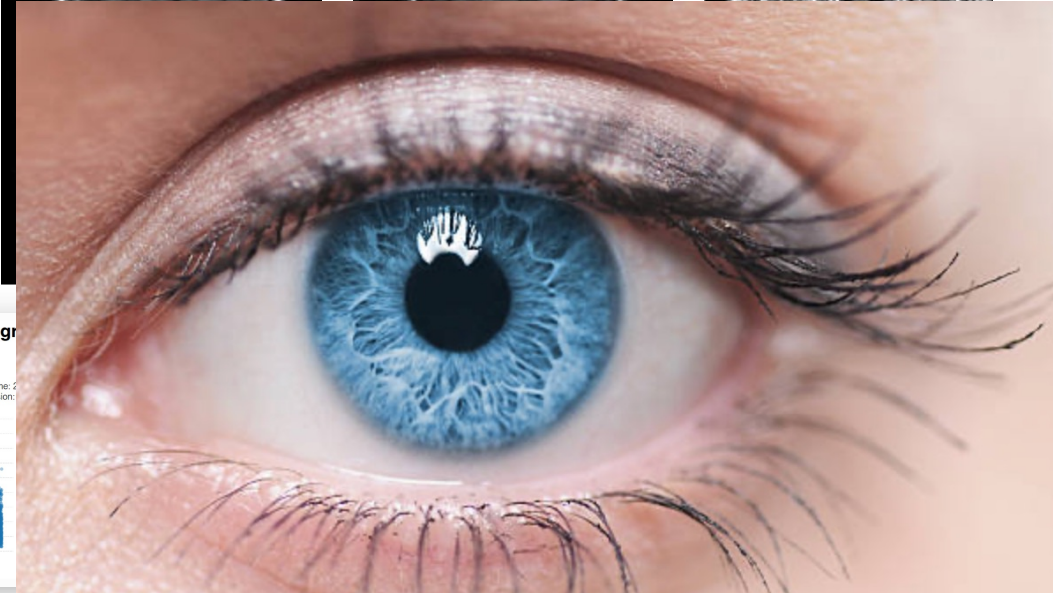
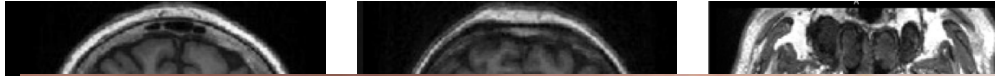
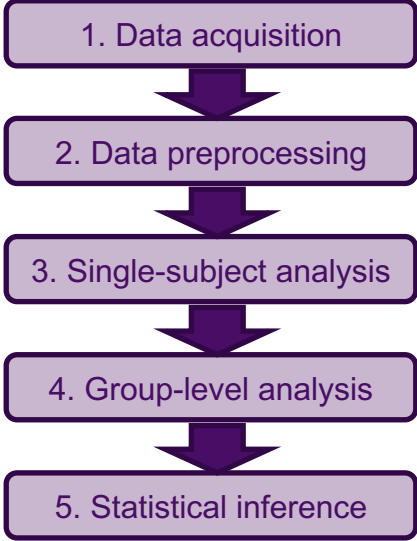
Total outliers (%)	0.11
Outliers (b=1000 mm^2)	0.22
Outliers (b=2000 mm^2)	0.00
Outliers (PE dir=[0, 1, 0, 0])	0.00
Outliers (PE dir=[0, -1, 0, 0])	0.11



MRIQC - <https://mriqc.readthedocs.io/en/latest/about.html>

CAT12 - <https://neuro-jena.github.io/cat/index.html#QC>

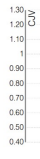
EDDY-QC (FSL) - <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/eddyqc/UsersGuide>



MRIQC: gr

Summary

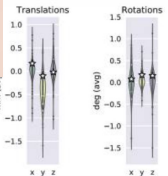
- Date and time: 2
- MRIQC version: 1



Within-volume motion

Avg std x translation (mm)	0.02
Avg std y translation (mm)	0.11
Avg std z translation (mm)	0.04
Avg std x rotation (deg)	0.05
Avg std y rotation (deg)	0.05
Avg std z rotation (deg)	0.06

0.11
0.22
0.00
0.00
0.11

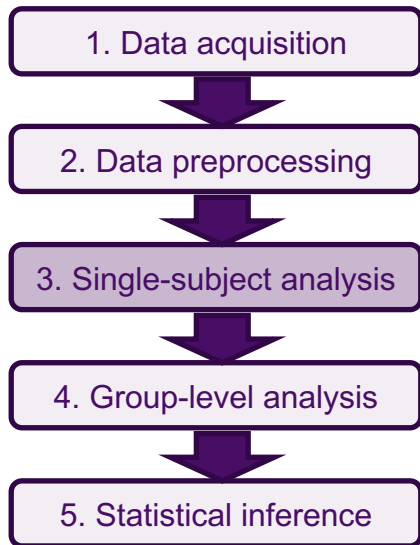


linear rating scale
nominal numbers
nominal letters
description

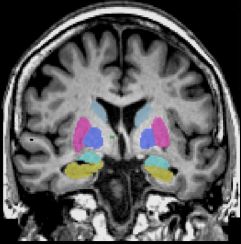
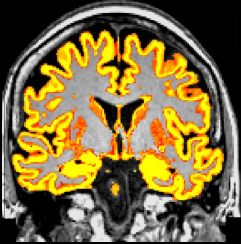
0.5	1	1.5	2	2.5	3	3.5	4	4.5	5	5.5	6	7	8		
1+	1	1-	2+	2-	3+	3-	4+	4-	5+	5-	6	7	8		
A+	A	A-	B+	B	B-	C+	C	C-	D+	D	D-	E+	E	E-	F
excellent	good	satisfactory	sufficient	critical	unacceptable /										

MRIQC - <https://mriqc.readthedocs.io/en/latest/about.html>
 CAT12 - <https://neuro-jena.github.io/cat/index.html#QC>
 EDDY-QC (FSL) - <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/eddyqc/UsersGuide>

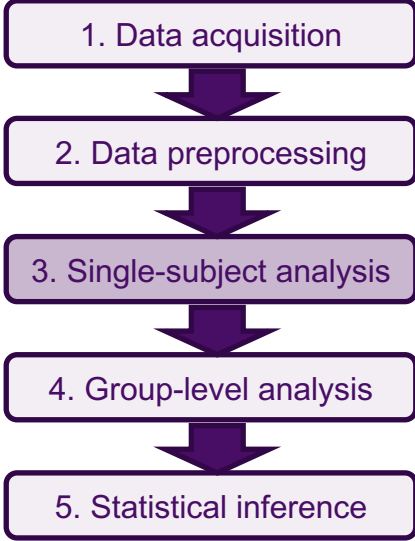
NEUROIMAGING DATA VISUALIZATION: SINGLE SUBJECT OUTPUT



Images
or
Numbers

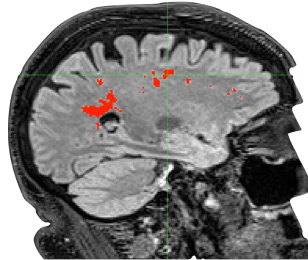
Images	Label/Region of interest (ROI)/ Parcel	Continuous measure
Volumetric		

NEUROIMAGING DATA VISUALIZATION: SINGLE SUBJECT OUTPUT

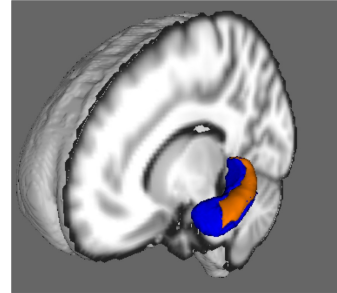


Images
or
Numbers

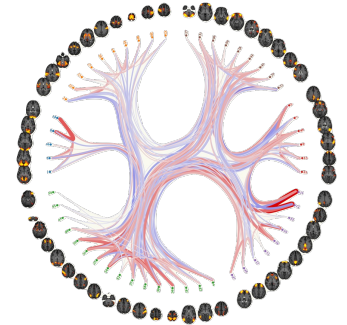
Numbers a.k.a. Imaging Derived Phenotypes (IDPs)



White matter
lesion load



Brain structure
volume



Connection
strength

NEUROIMAGING DATA VISUALIZATION: GROUP-LEVEL STATISTICAL OUTPUT

1. Data acquisition

2. Data preprocessing

3. Single-subject analysis

4. Group-level analysis

5. Statistical inference

- **Numbers** fed into 'classic' stats software (R, SPSS, STATA, python...)
- **Images** require specific stats (usually within imaging software tools)
- Input = single subject output, registered to a template
- Statistical maps in pseudocolours shows **significant voxels** (volumetric) or vertices (surface), overlaid on template.
- **Atlases** can help interpreting results

NEUROIMAGING DATA VISUALIZATION:
GROUP-LEVEL STATISTICAL OUTPUT

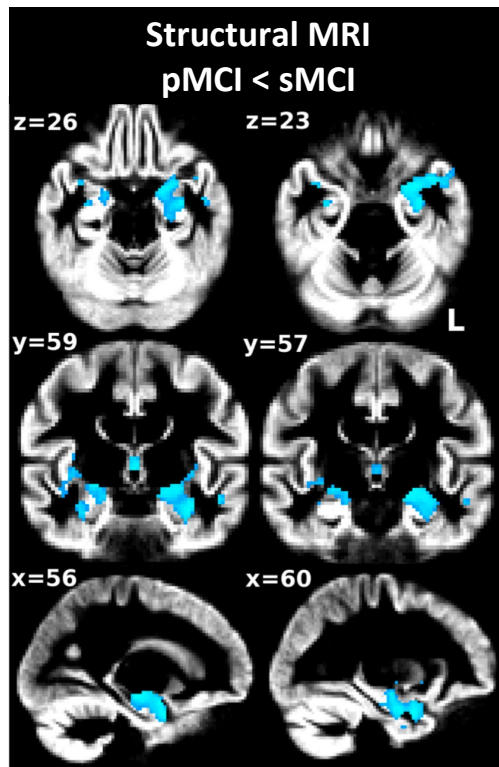
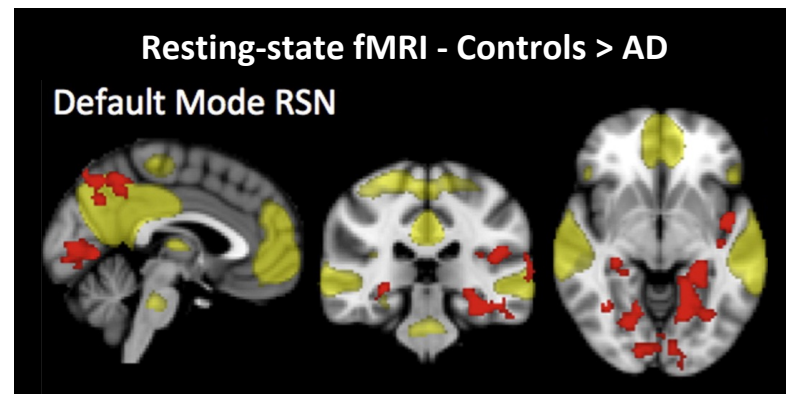
1. Data acquisition

2. Data preprocessing

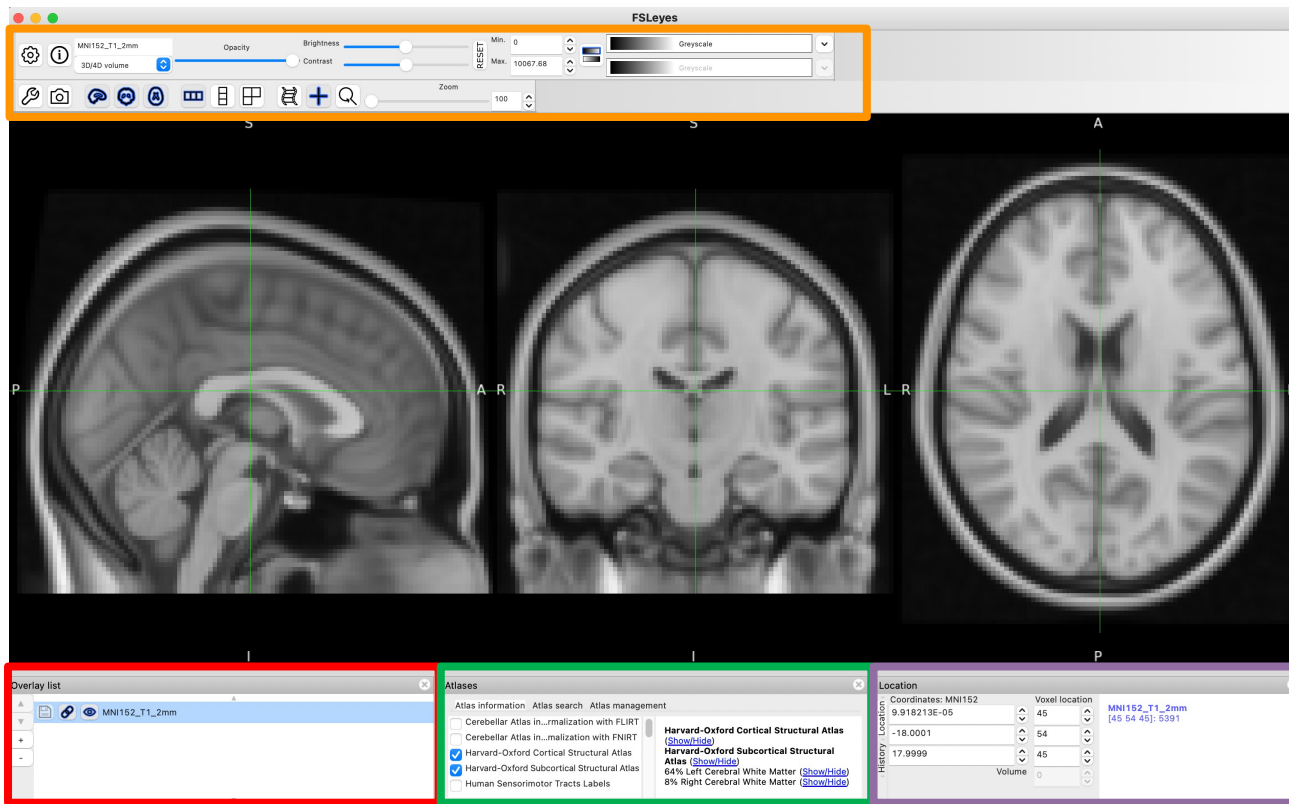
3. Single-subject analysis

4. Group-level analysis

5. Statistical inference

*Douaud et al., JNeurosci 2013**Zamboni et al., Biol Psych. 2013*

1. Getting started



2. Viewing multiple images

4. Viewing Atlases

3. Image information

THANK YOU!



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Next up:

Basics of Neuroimaging: Structural Magnetic Resonance Imaging (MRI) by David Cash

14 April, 2023; 9AM – 10AM CT

Basics of Neuroimaging: Positron emission tomography (PET) by Tobey Betthausen

19 April, 2023; 12PM - 1PM CT

Basics of Neuroimaging: Diffusion-Weighted Imaging (DWI) by Alexa Pichet Binette

21 April, 2023; 9AM – 10AM CT

Basics of Neuroimaging: Functional Magnetic Resonance Imaging (fMRI) by Luigi Lorenzini

26 April, 2023; 10AM – 11AM CT