

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

ALZHEIMER'S  ASSOCIATION®

AAIC > 23

BASICS OF NEUROIMAGING

DATA STRUCTURE AND FORMATS

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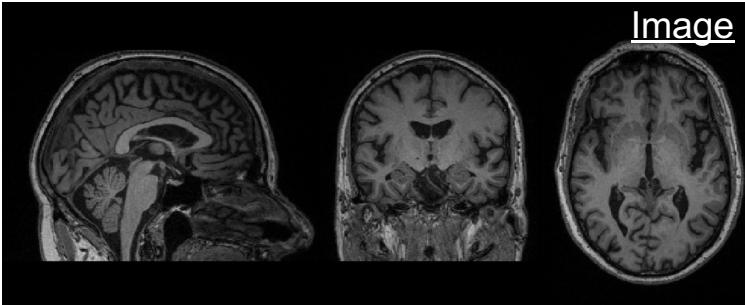


@ludogriffanti

AAIC > 23 GOALS

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

Image

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time_units	s	intent_p2	0.000000
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nhyper	4	qform_name	Scanner Anat
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pixdim2	1.000000	ato_xyz:3	0.032771 0.017677 0.999307 94.454788
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		aux_file	OAS30015_MR_d2004

HeaderImage formats

“Raw” Scanner File Format
Example: DICOM



DICOM to NIFTI conversion (e.g. dcm2niix)

Analysis File Format
Example: **NIfTI** (Neuroimaging
Informatics Technology
Initiative)

Image = cube of numbers

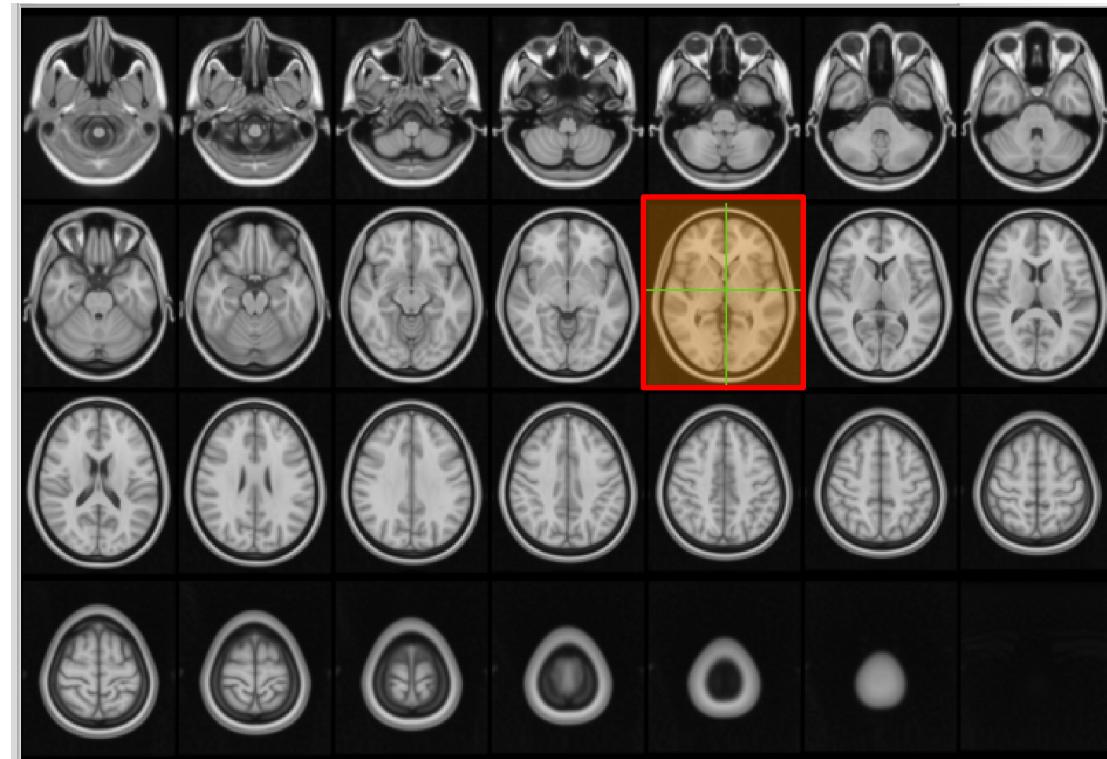
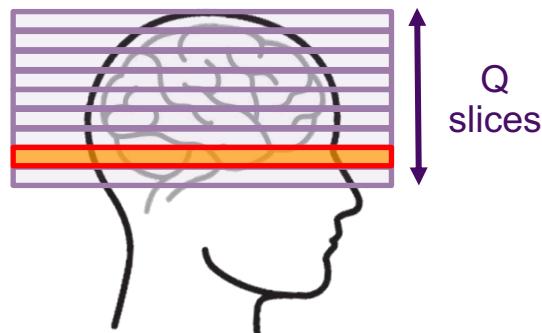


Image = cube of numbers

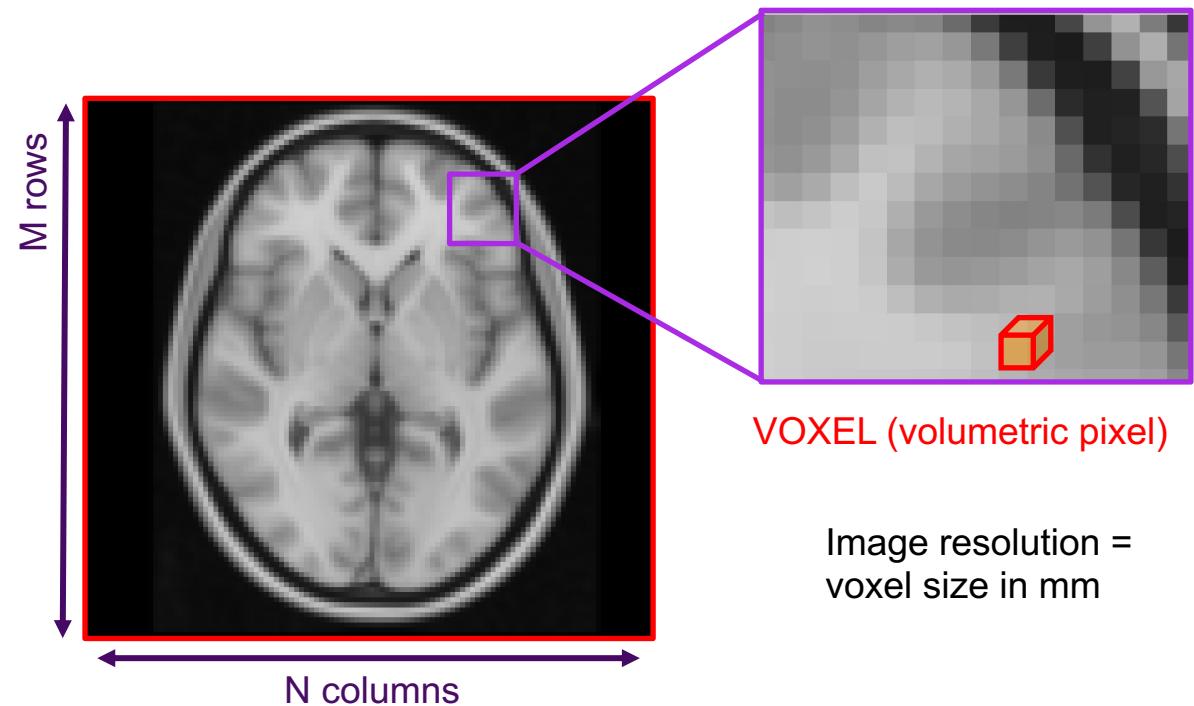
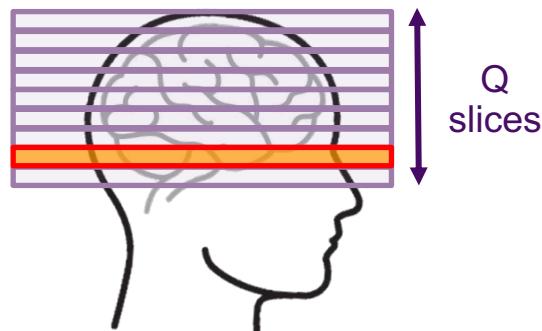
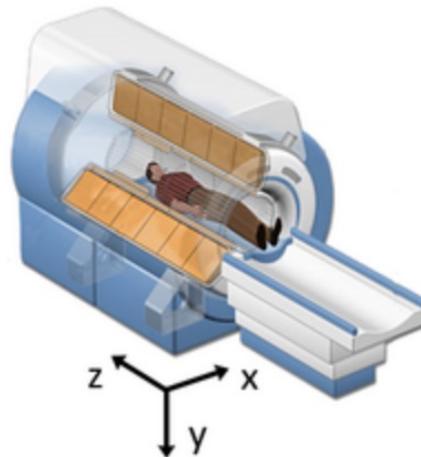


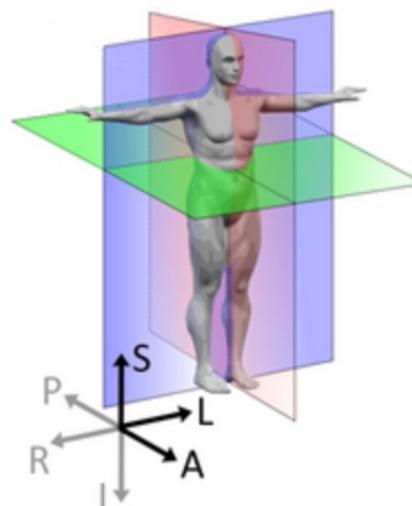
Image resolution =
voxel size in mm

AAIC>23 PROPERTIES OF MEDICAL IMAGING DATA

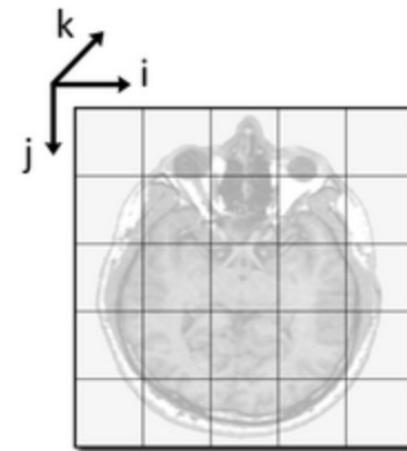
Coordinate systems



Scanner coordinates



Subject
coordinates



Voxel coordinates

IMAGE COORDINATES & IMAGE REGISTRATION

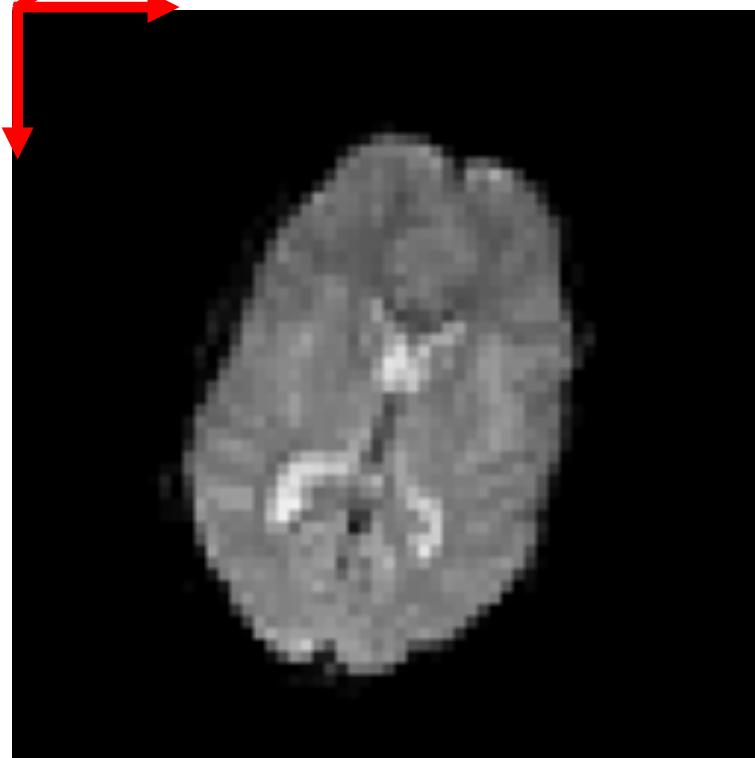


IMAGE COORDINATES & IMAGE REGISTRATION

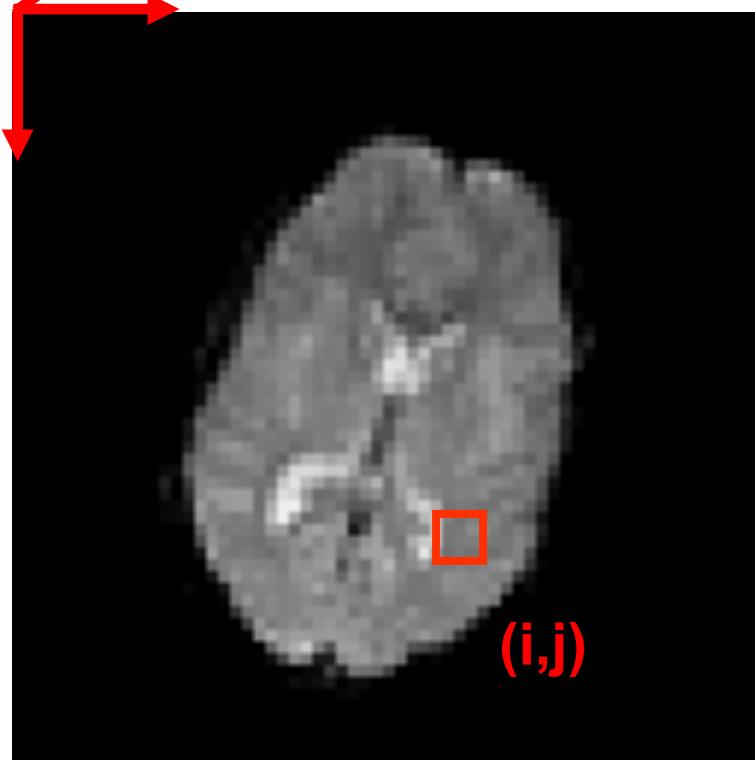
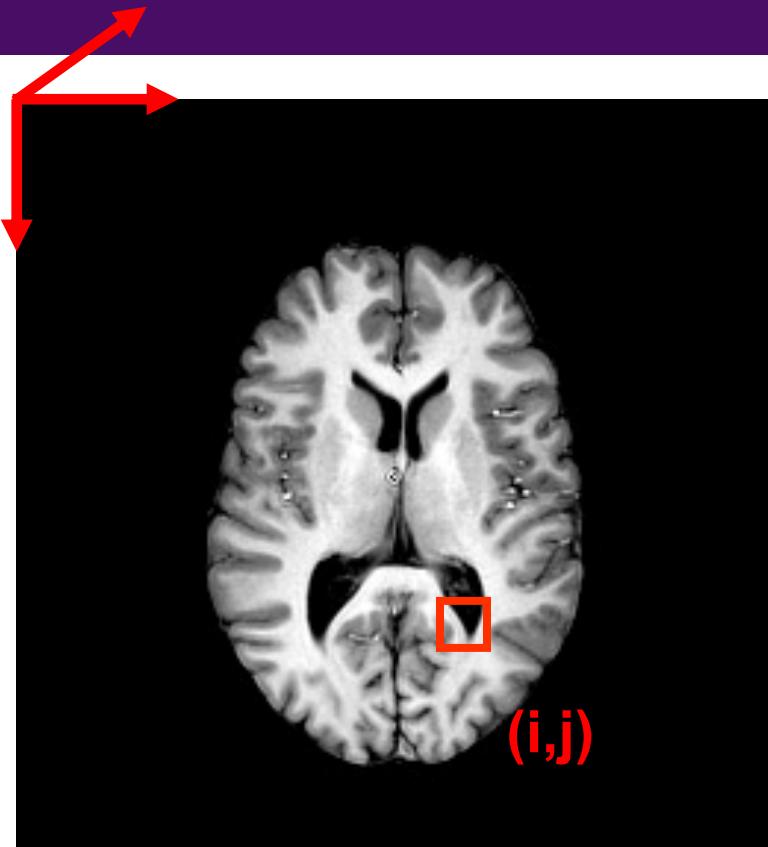
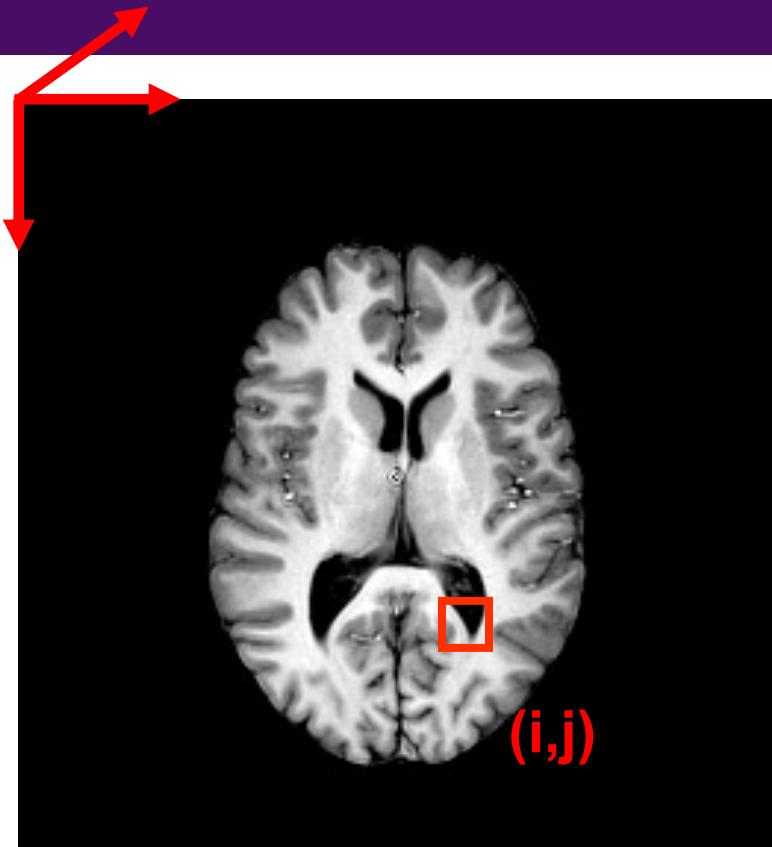
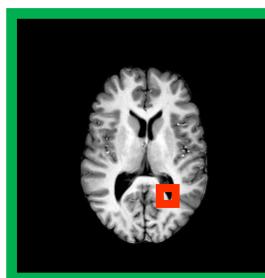
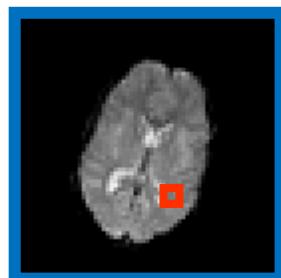
 (i, j)  (i, j)

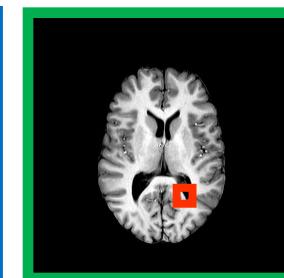
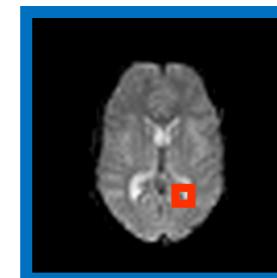
IMAGE COORDINATES & IMAGE REGISTRATION

 (i, j)  (i, j)

- **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)



Transformation,
deformation, warp,
mapping

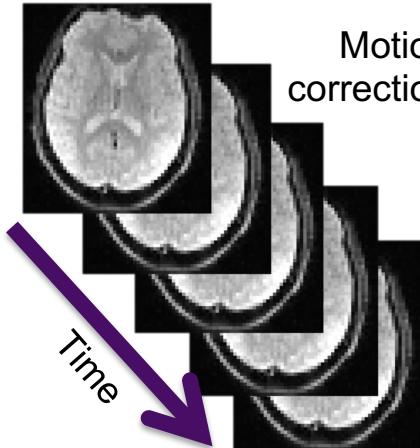


- Input Image
- Moving Image
- Source Image
- Deformed Image

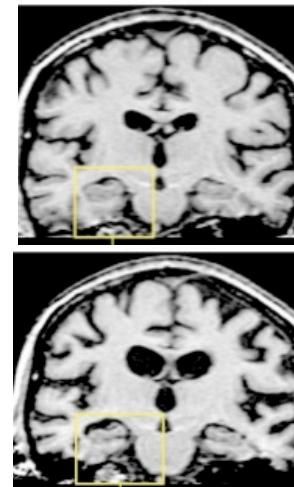
- Reference image
- Stationary Image
- Target Image
- Fixed Image

Within-subject & session

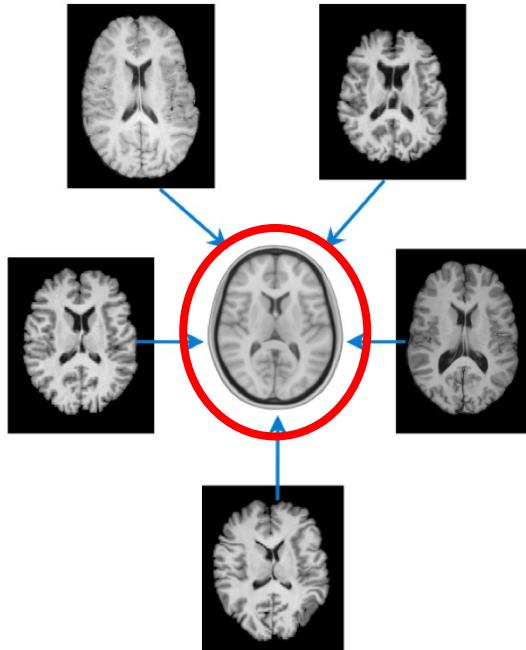
Motion correction



Between-modalities

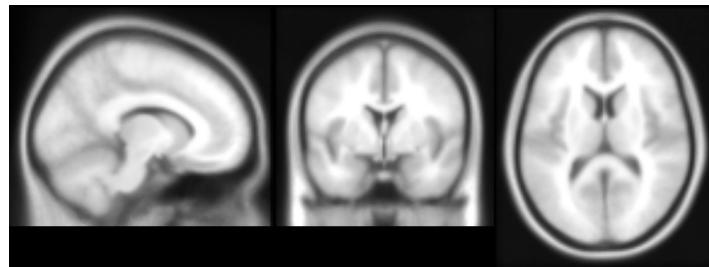
Within-subject,
between sessions

Scheltens et al., 2002

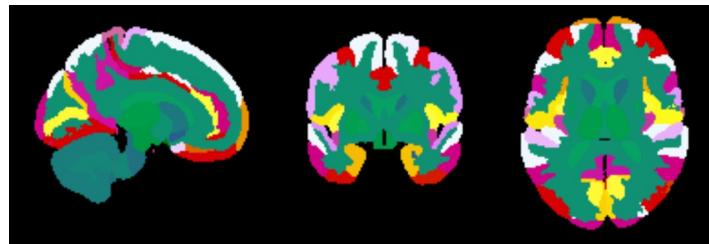
Longitudinal data,
change over timeBetween-subjectsTemplate / Standard space =
"average brain" used as reference

“Average
brain”

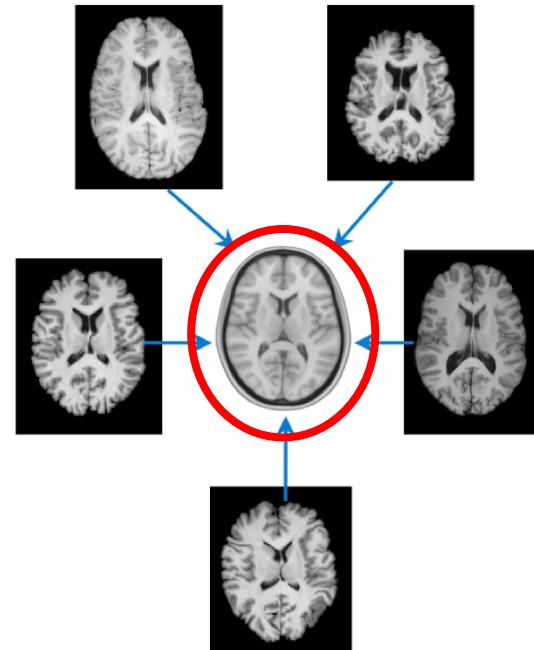
MNI152 Template Space



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

NEUROIMAGING DATA ANALYSIS: DATA STRUCTURE

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.



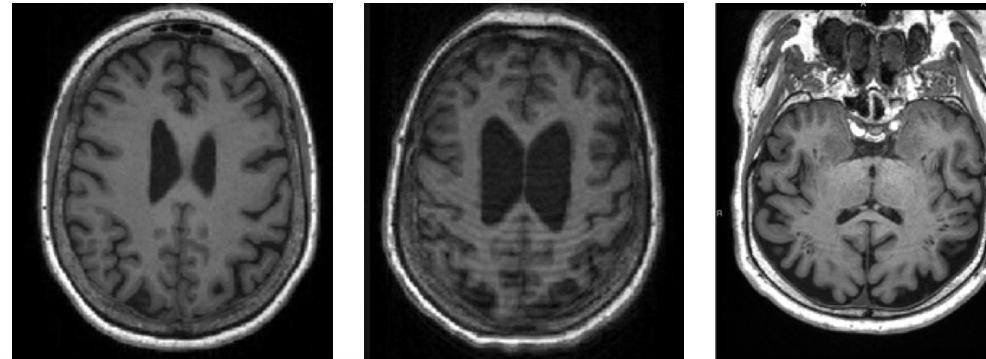
<https://bids.neuroimaging.io>

NEUROIMAGING DATA VISUALIZATION: QUALITY CONTROL

1. Data acquisition



2. Data preprocessing



MRIQC: group anatomical report

Summary

- Date and time: 2017-02-05, 12:27.
- MRIQC version: 0.9.0-rc2.

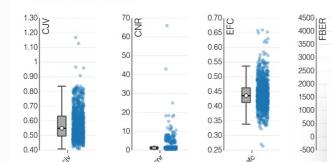


Image quality definition

BWP noise (in percent)

BWP bias (in percent)

resolution RES (mm)

Quality ratings

percentage rating points (rps)

linear rating scale

nominal numbers

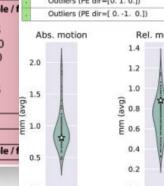
nominal letters

description

	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	4.0
	100	95	90	85	80	75
	1	1.5	2	2.5	3	3.5
	A+	A	A-	B+	B	C+
	excellent	good	satisfactory	sufficient	critical	unacceptable

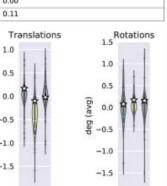
Total outliers (%)	0.11
Avg abs. motion (mm)	0.81
Avg std x translation (mm)	0.88
Avg x translation (mm)	0.17
Avg y translation (mm)	-0.10
Avg z translation (mm)	-0.02
Avg x rotation (deg)	0.07
Avg y rotation (deg)	0.17
Avg z rotation (deg)	0.15

Outliers	
Total outliers (%)	0.11
Outliers ($B=1000 \text{ mm}^2$)	0.22
Outliers ($B=2000 \text{ mm}^2$)	0.00
Outliers (PE dir=[0, 1, 0])	0.00
Outliers (PE dir=[0, -1, 0])	0.11



Total outliers (%)	0.03
Avg std x translation (mm)	0.11
Avg std y translation (mm)	0.04
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 ($B=1000 \text{ mm}^2$)	0.22
Outliers ($B=2000 \text{ mm}^2$)	0.00
Outliers (PE dir=[0, 1, 0])	0.00
Outliers (PE dir=[0, -1, 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>

NEUROIMAGING DATA VISUALIZATION: QUALITY CONTROL

1. Data acquisition



2. Data preprocessing



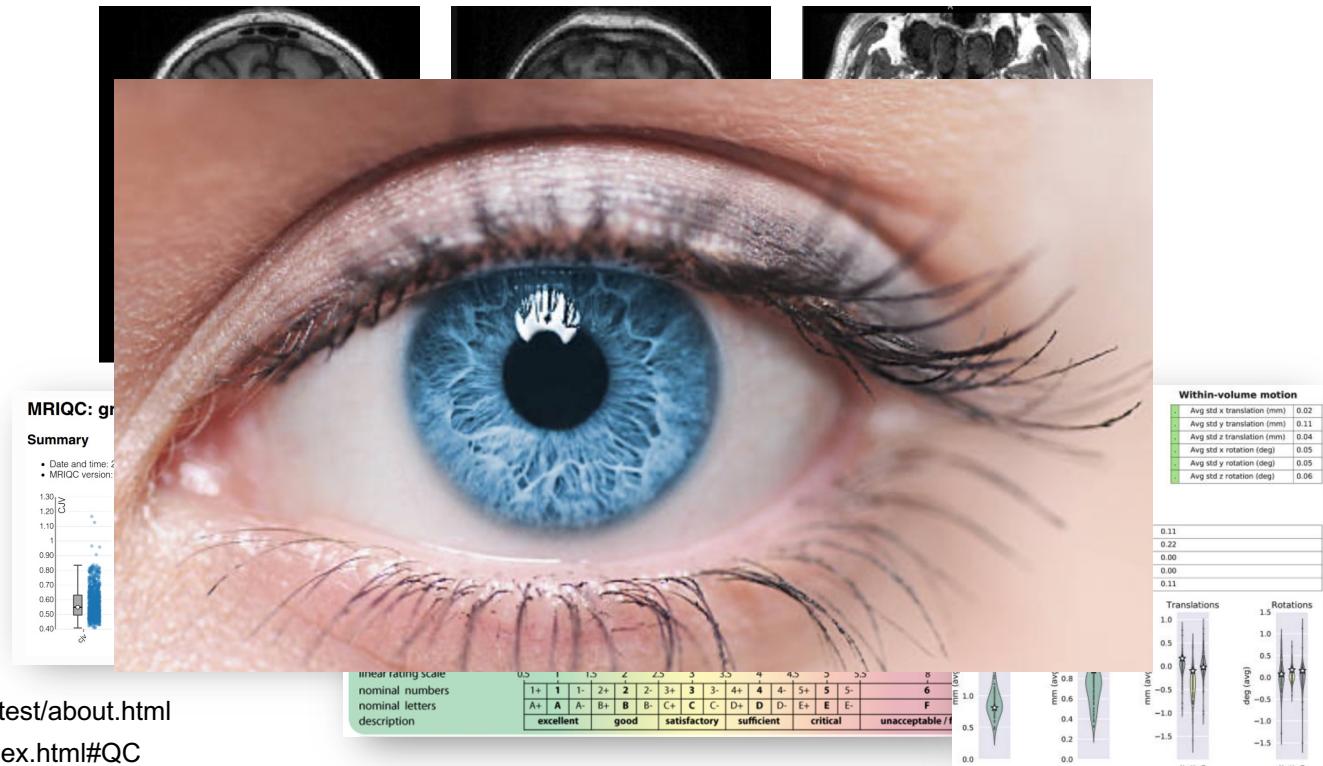
3. Single-subject analysis



4. Group-level analysis



5. Statistical inference

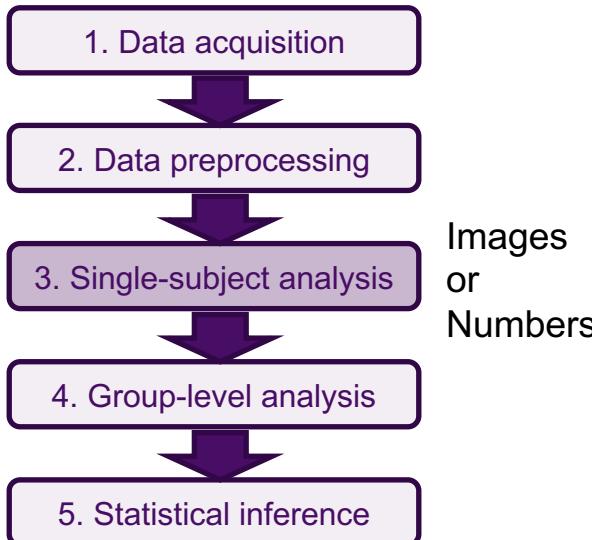


MRIQC - <https://mriqc.readthedocs.io/en/latest/about.html>

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

Images from: Miller et al., *Nat Neurosci.* 2016; FreeSurfer tutorial; Wang et al., *PlosONE* 2013

NEUROIMAGING DATA VISUALIZATION: SINGLE SUBJECT OUTPUT

1. Data acquisition



2. Data preprocessing



3. Single-subject analysis

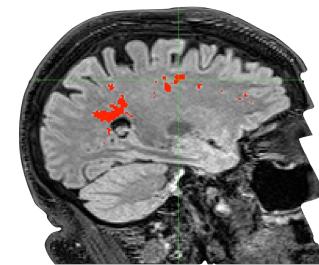


4. Group-level analysis

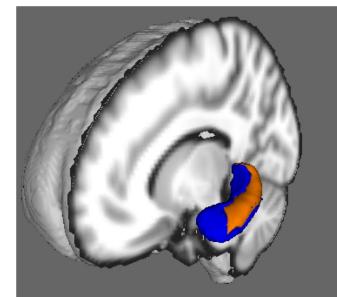


5. Statistical inference

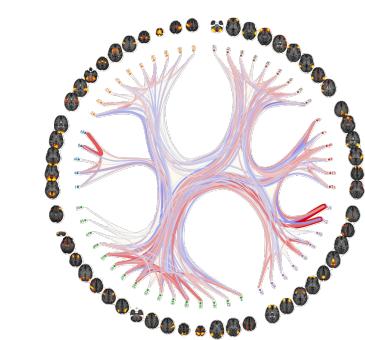
Images
or
Numbers



White matter
lesion load



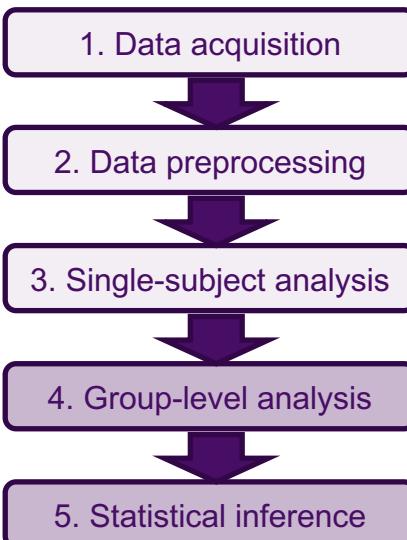
Brain structure
volume



Connection
strength

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

NEUROIMAGING DATA VISUALIZATION: GROUP-LEVEL STATISTICAL OUTPUT



- **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

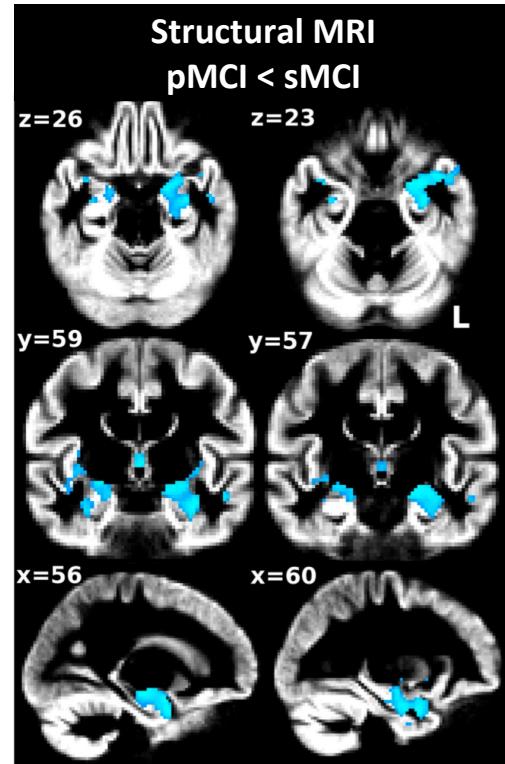
1. Data acquisition

2. Data preprocessing

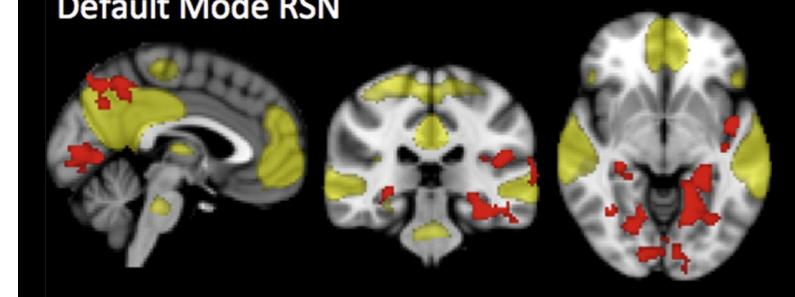
3. Single-subject analysis

4. Group-level analysis

5. Statistical inference

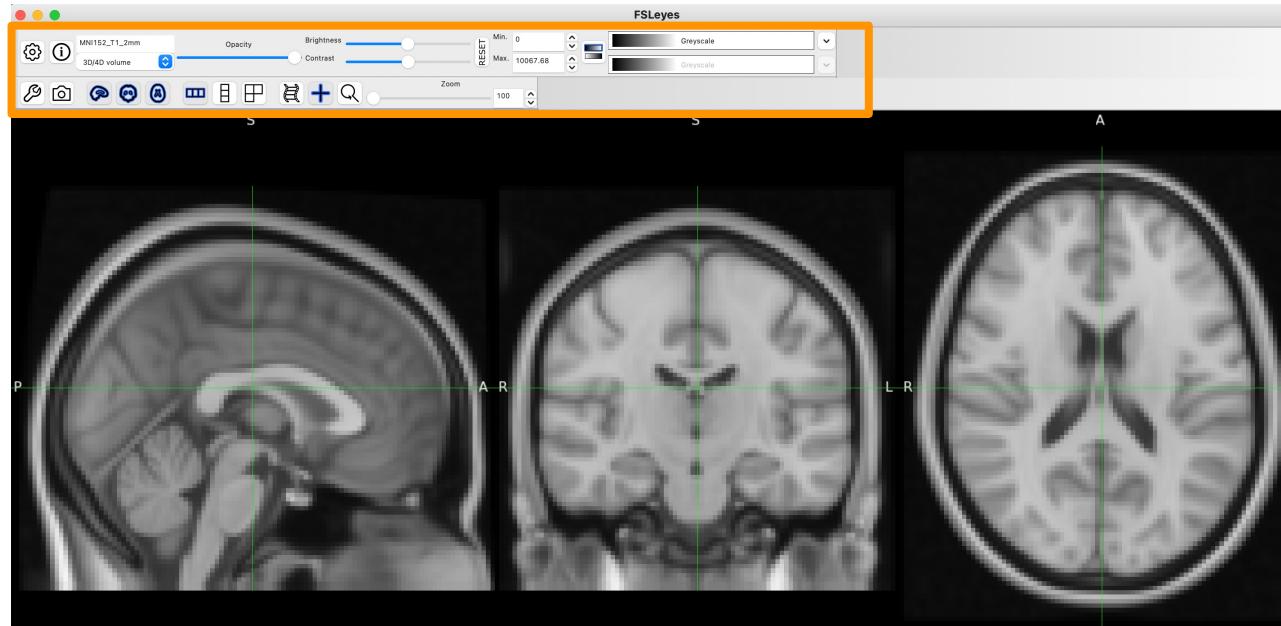


Resting-state fMRI - Controls > AD
Default Mode RSN

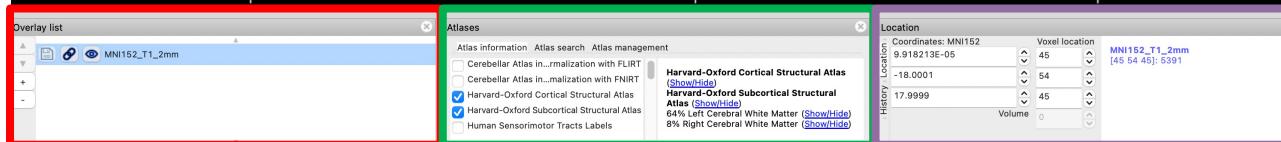


NEUROIMAGING DATA VISUALIZATION: FSLEYES

1. Getting started



2. Viewing multiple images



4. Viewing Atlases

3. Image information

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The Basics of Neuroimaging Series

THANK YOU!



@ludogriffanti



ludovica.griffanti@psych.ox.ac.uk

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 Betthauser
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