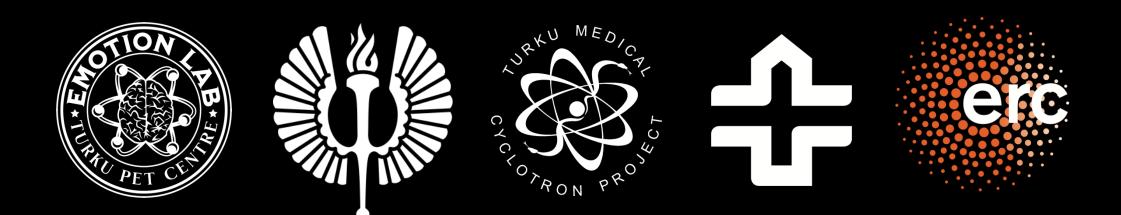
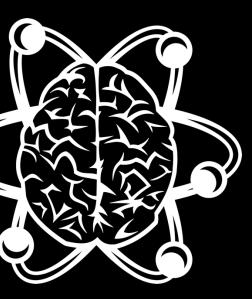
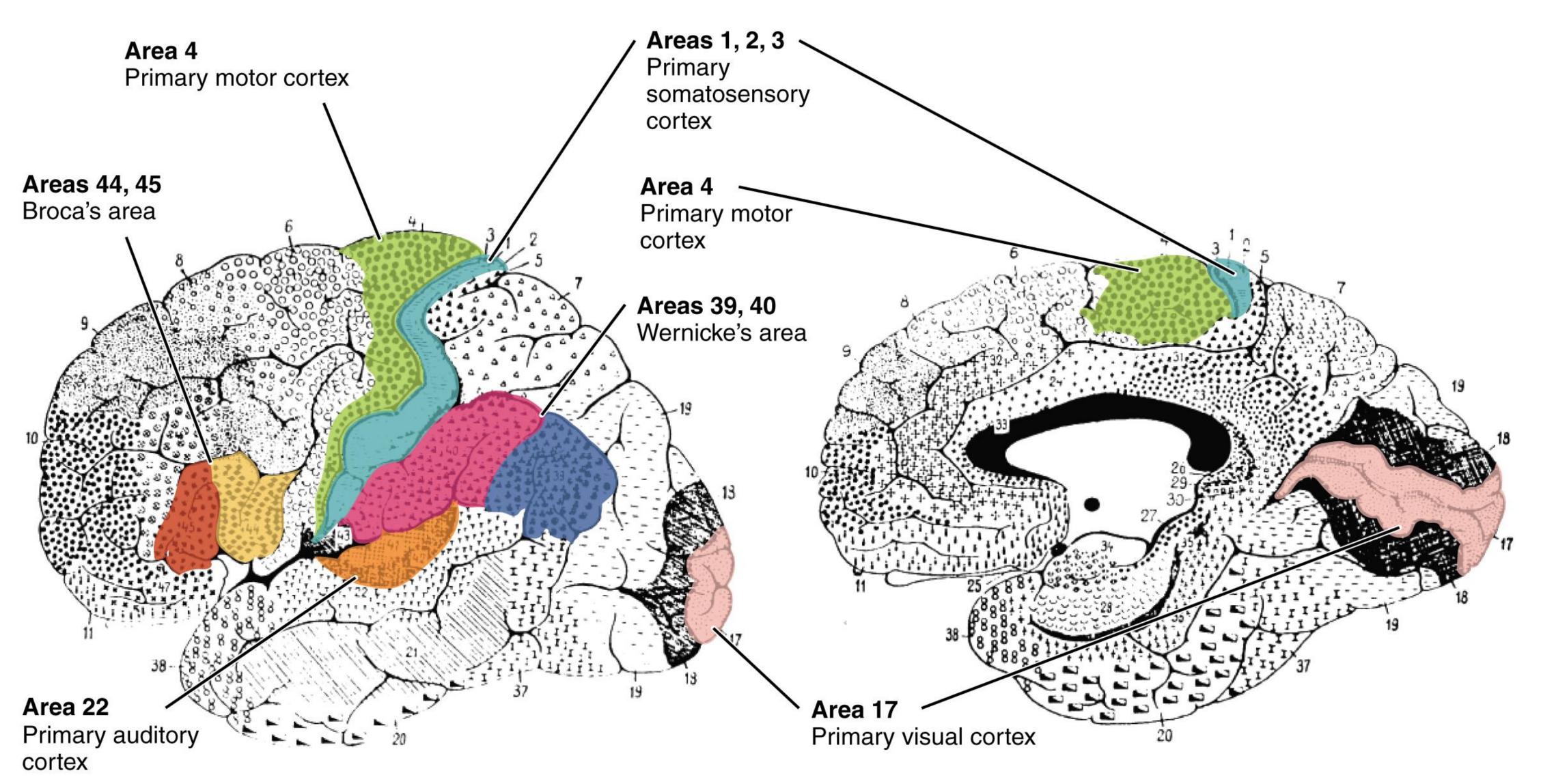
REGION-OF INTEREST ANALYSIS

Lauri Nummenmaa, Turku PET Centre





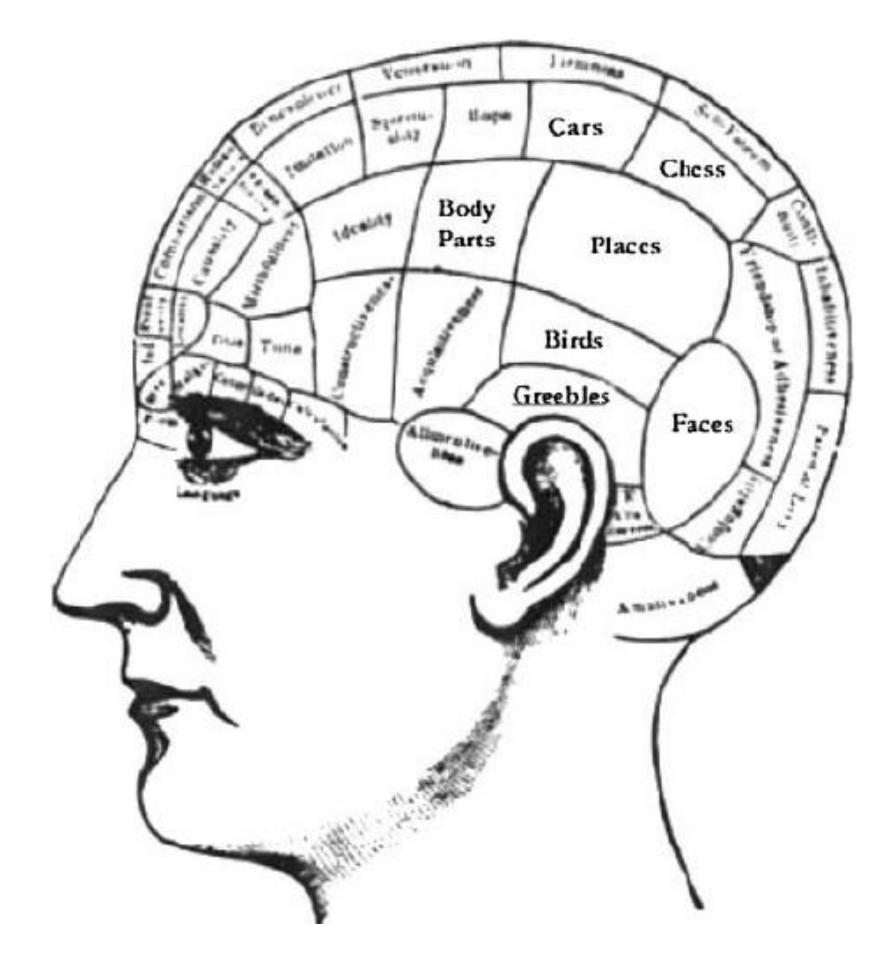
Turku PET Centre Brain Imaging Course 2024



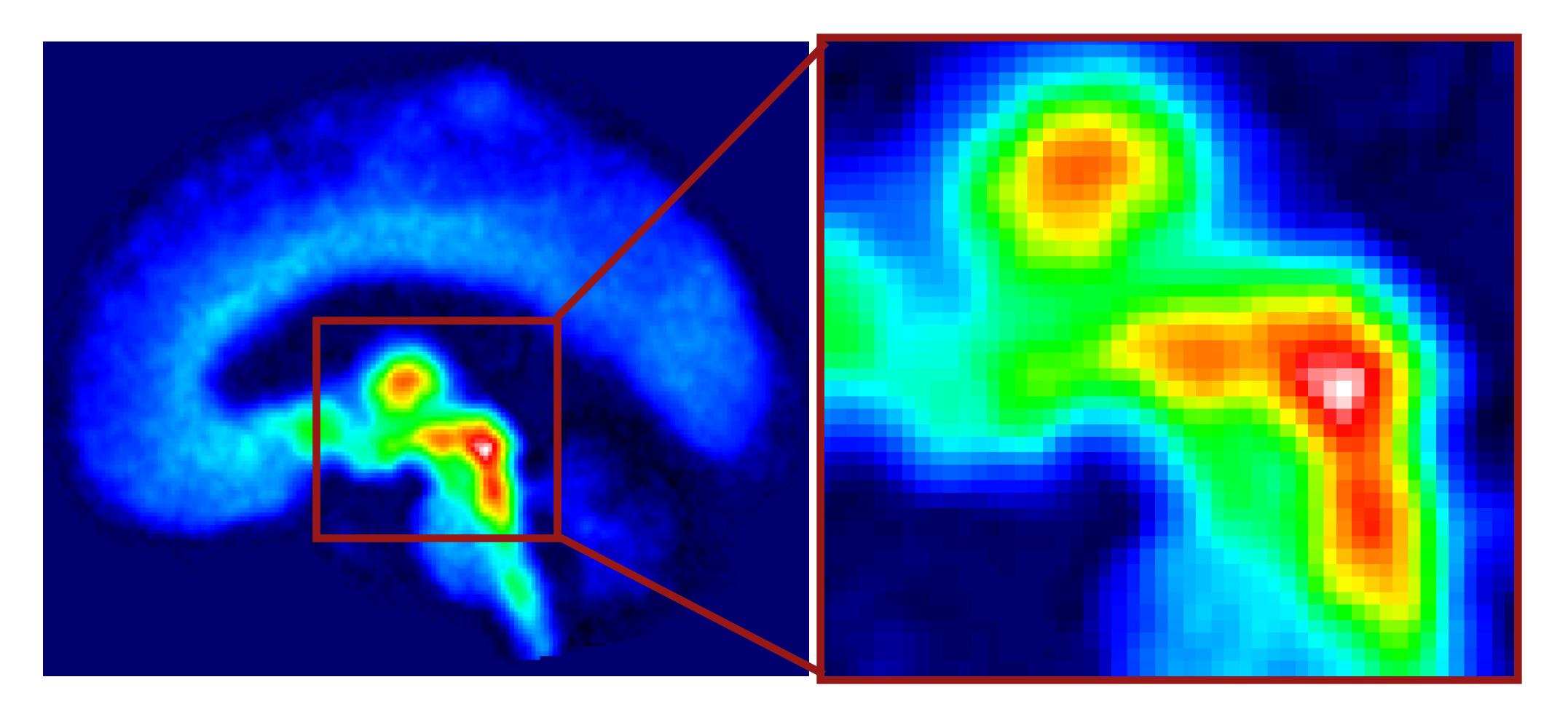
Brodmann's cytotechtonic map (1909): Lateral surface Brodmann's cytotechtonic map (1909): Medial surface

Brodmann areas

- Korbinian Brodmann (1909)
- Based on cytoarchitectural (cellular) organization of neurons observed in Nissl staining
- Initial idea of dividing brain into different regions based on their similarity — but similarity in what?



Why ROI analysis



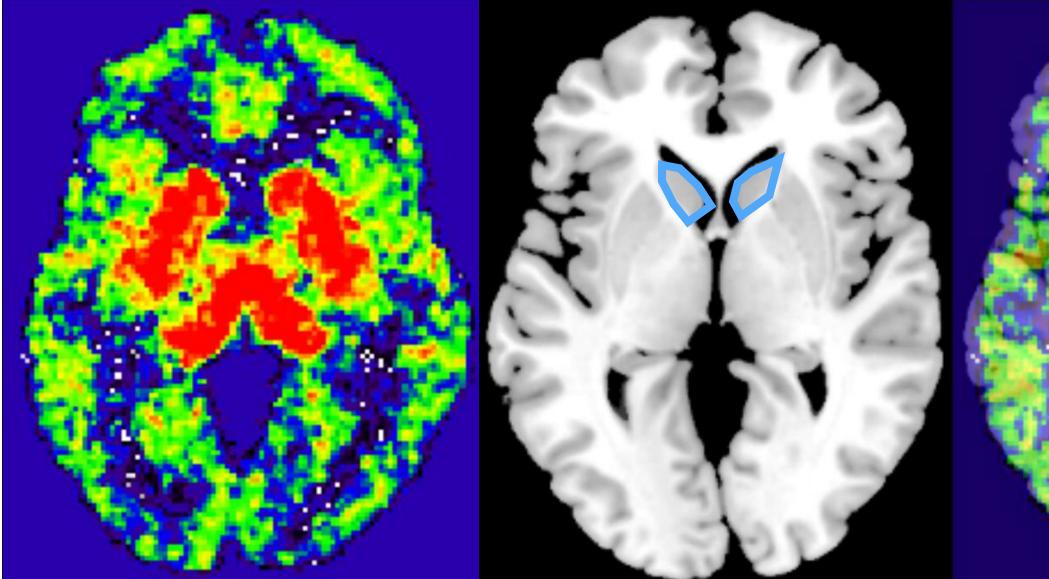
We still have good reasons! ROI data extraction = dimension reduction

- (e.g. multivariate analysis, Bayesian analyses...)
- Comparison across modalities
- Visualisation
- Need for high anatomical accuracy

Complex statistical models that cannot be accommodated to voxel space

Difficulties in interpretation of factorial statistical models at voxel level

Dimension reduction



- statistical tests

Univariate data regularly shaped can use univariate stats

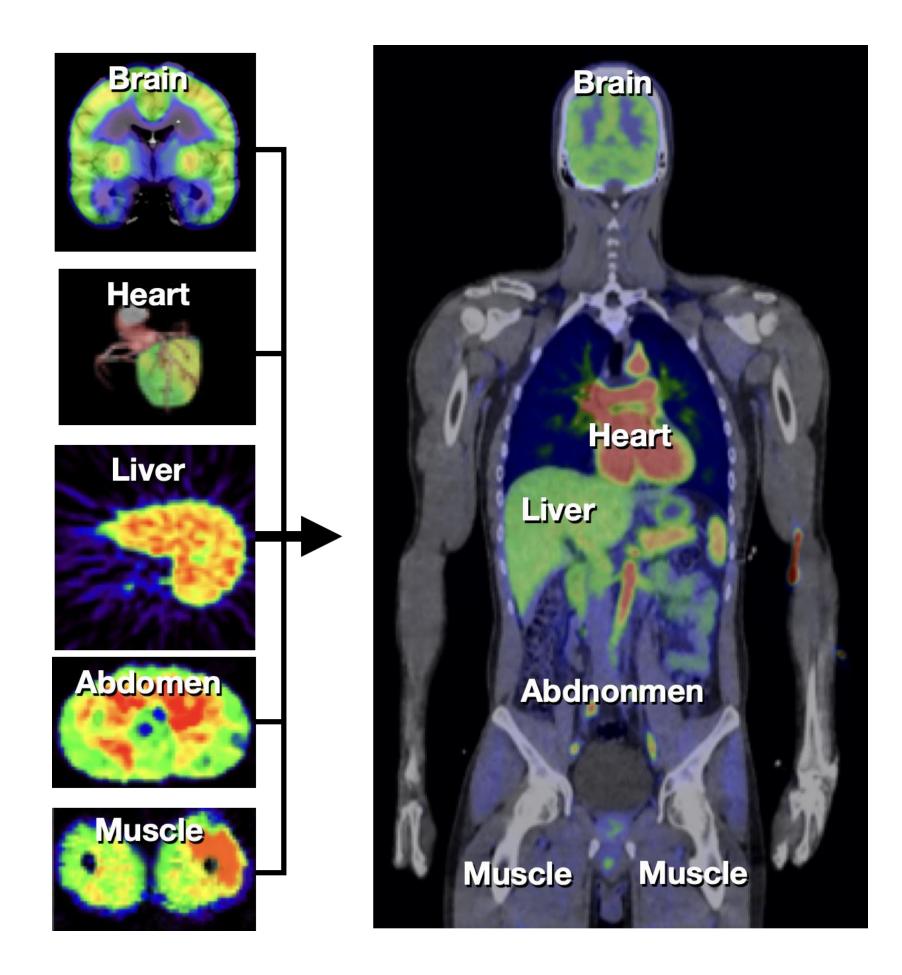
	Controls	Patien
	3	5
E ytroot	4	4
Extract outcome	5	6
measure	6	7
in ROI	3	6
	2	5
	3	2
	5	6
	2	8

Pros: Anatomically accurate if ROIs well definied, data can be analyzed with simple univariate

Cons: Laborious, using many ROIs not feasible, averaging within ROI not always appropriate



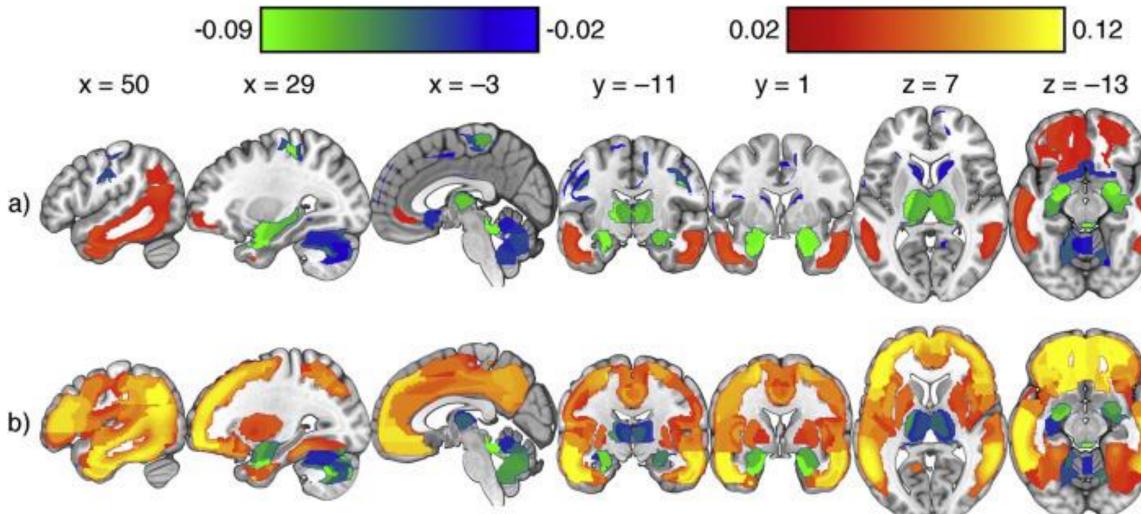
Major issue in total-body analysis

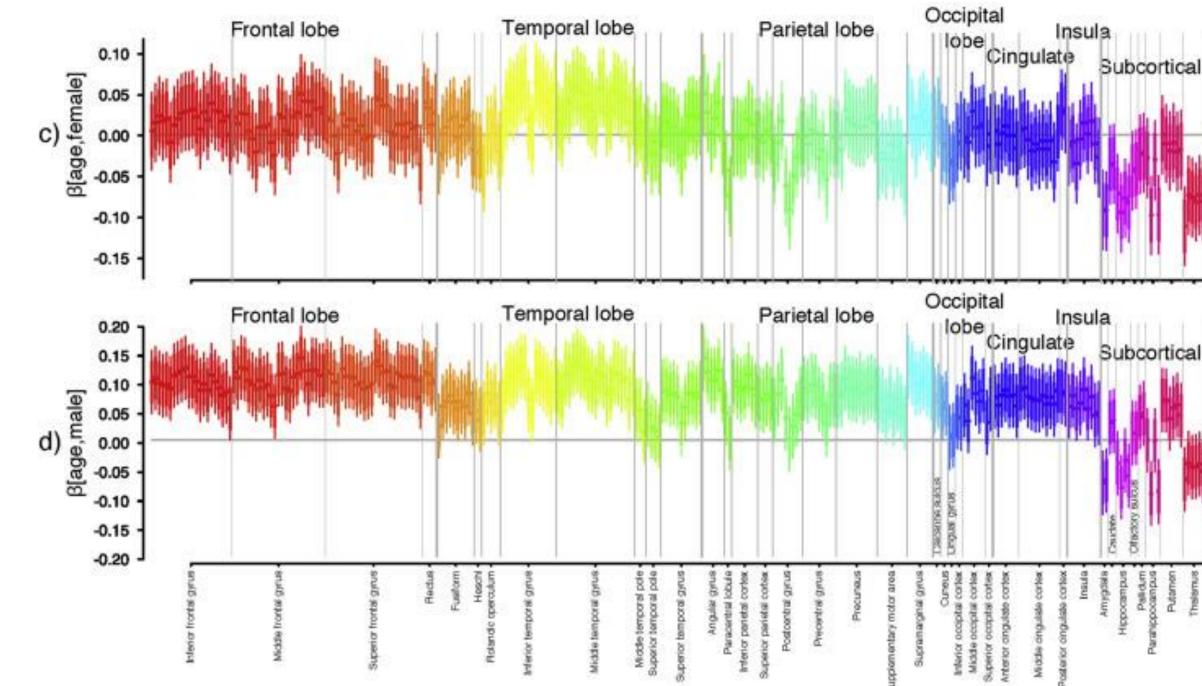


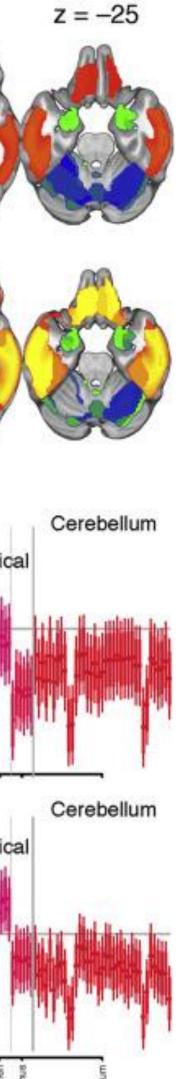


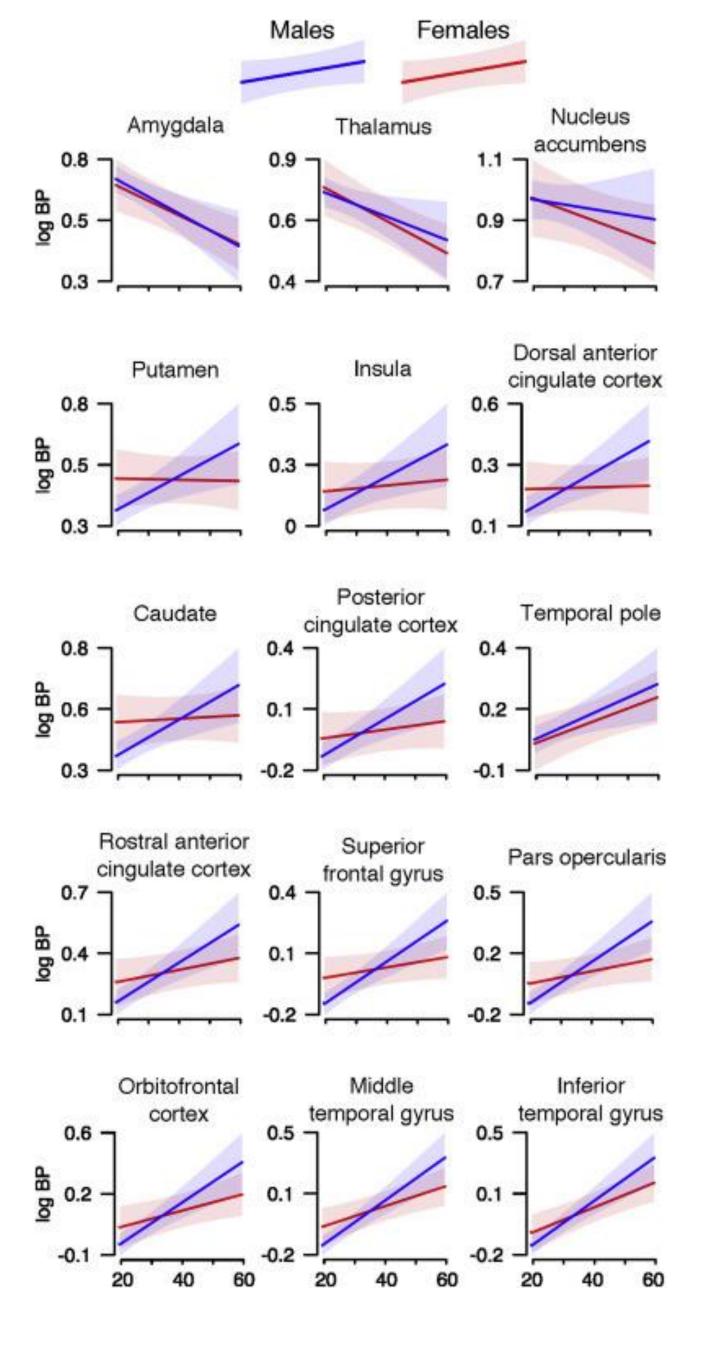
CT segments





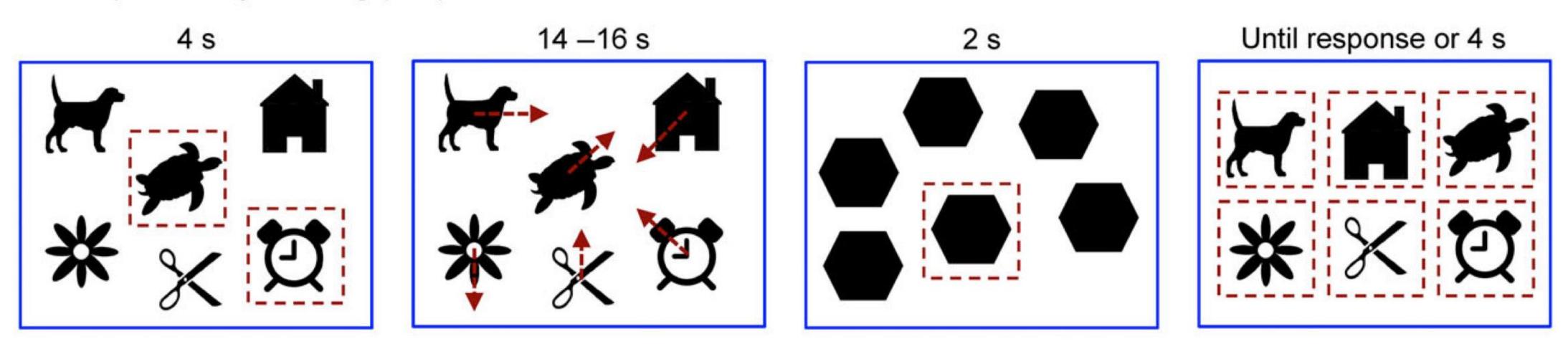




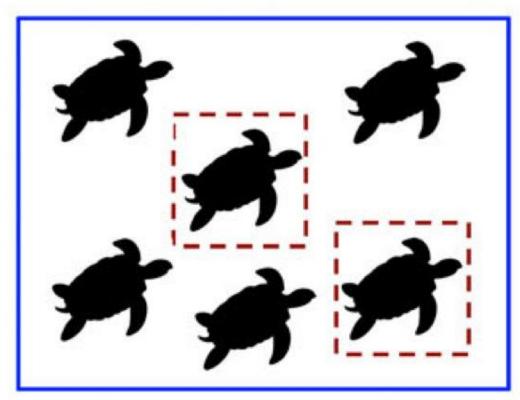


Kantonen et al (2020 Neurolmage)

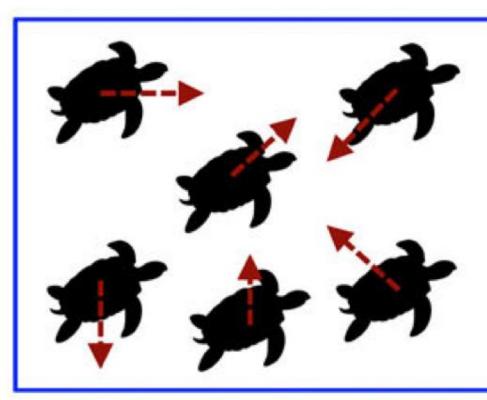
A Multiple Identity Tracking (MIT)



B Multiple Object Tracking (MOT)

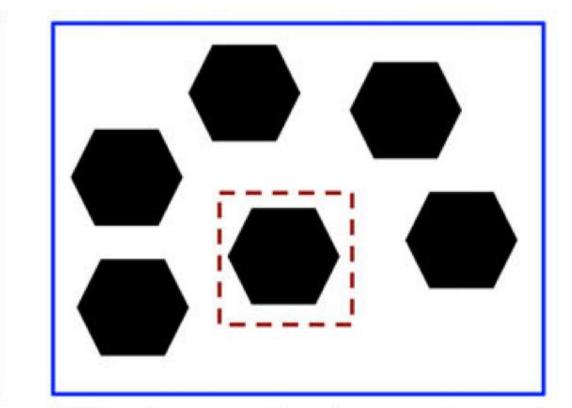


Target designation

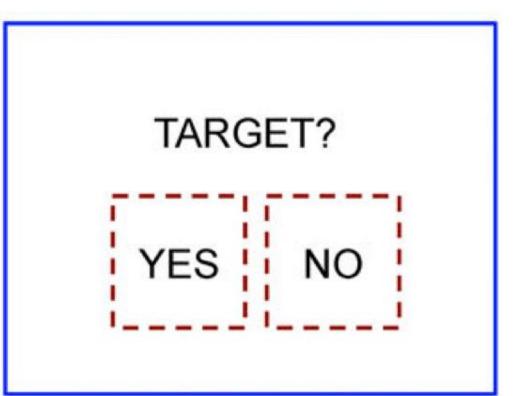


Target movement





Objects masked, one target marked



Probe phase Response with trackball

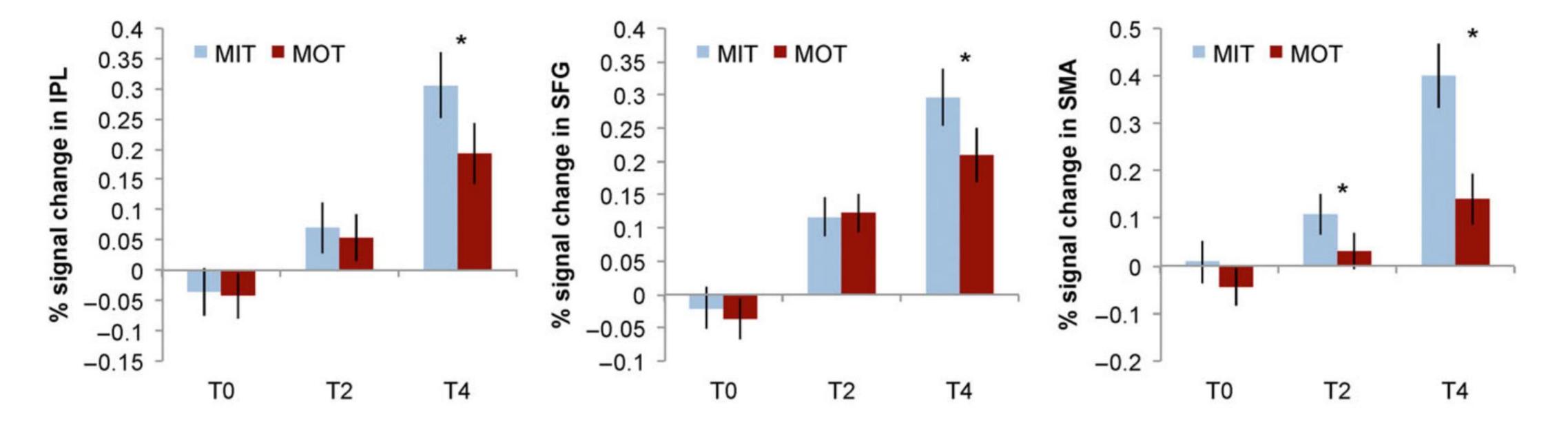
Nummenmaa et al (2017)

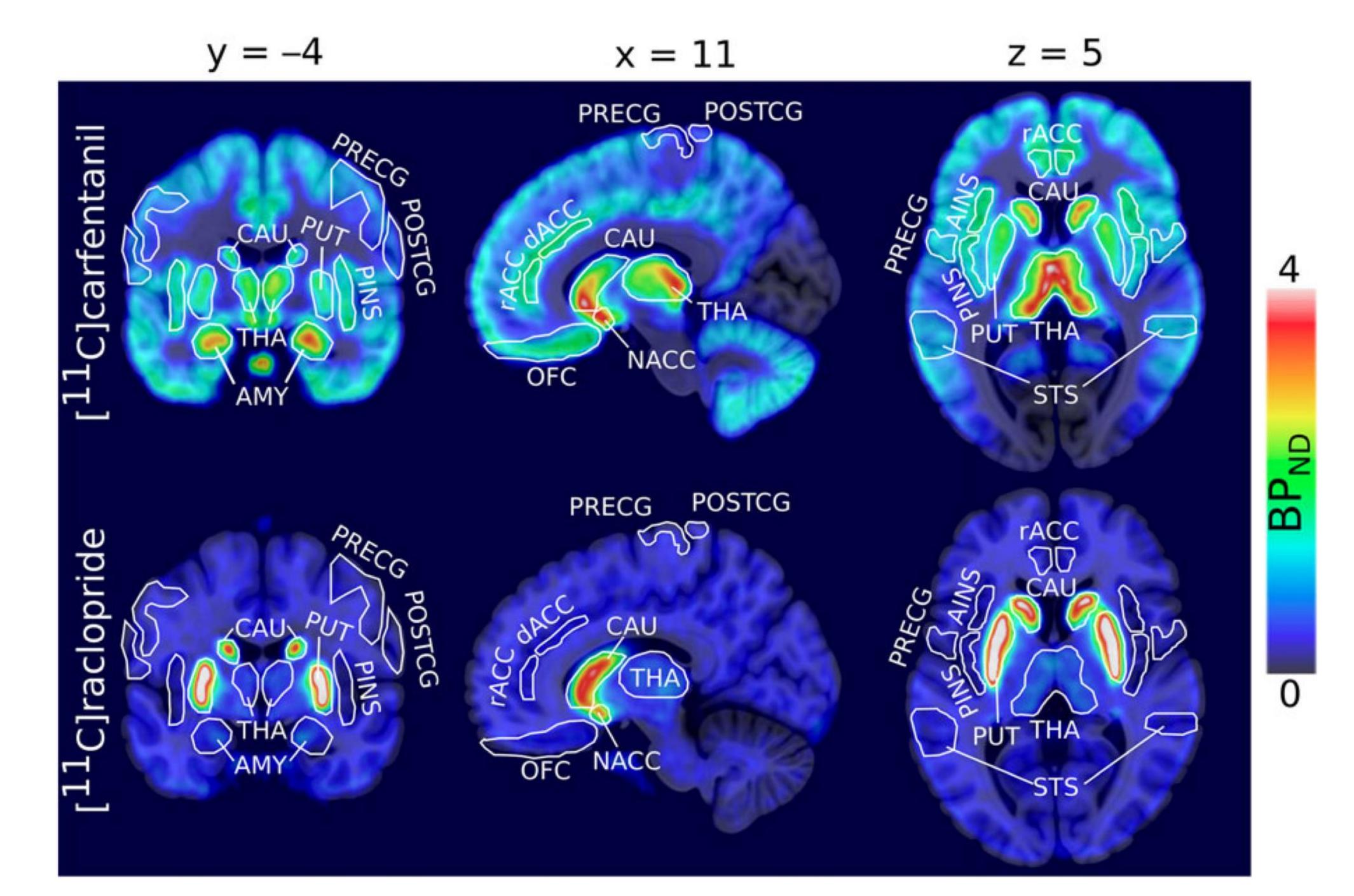


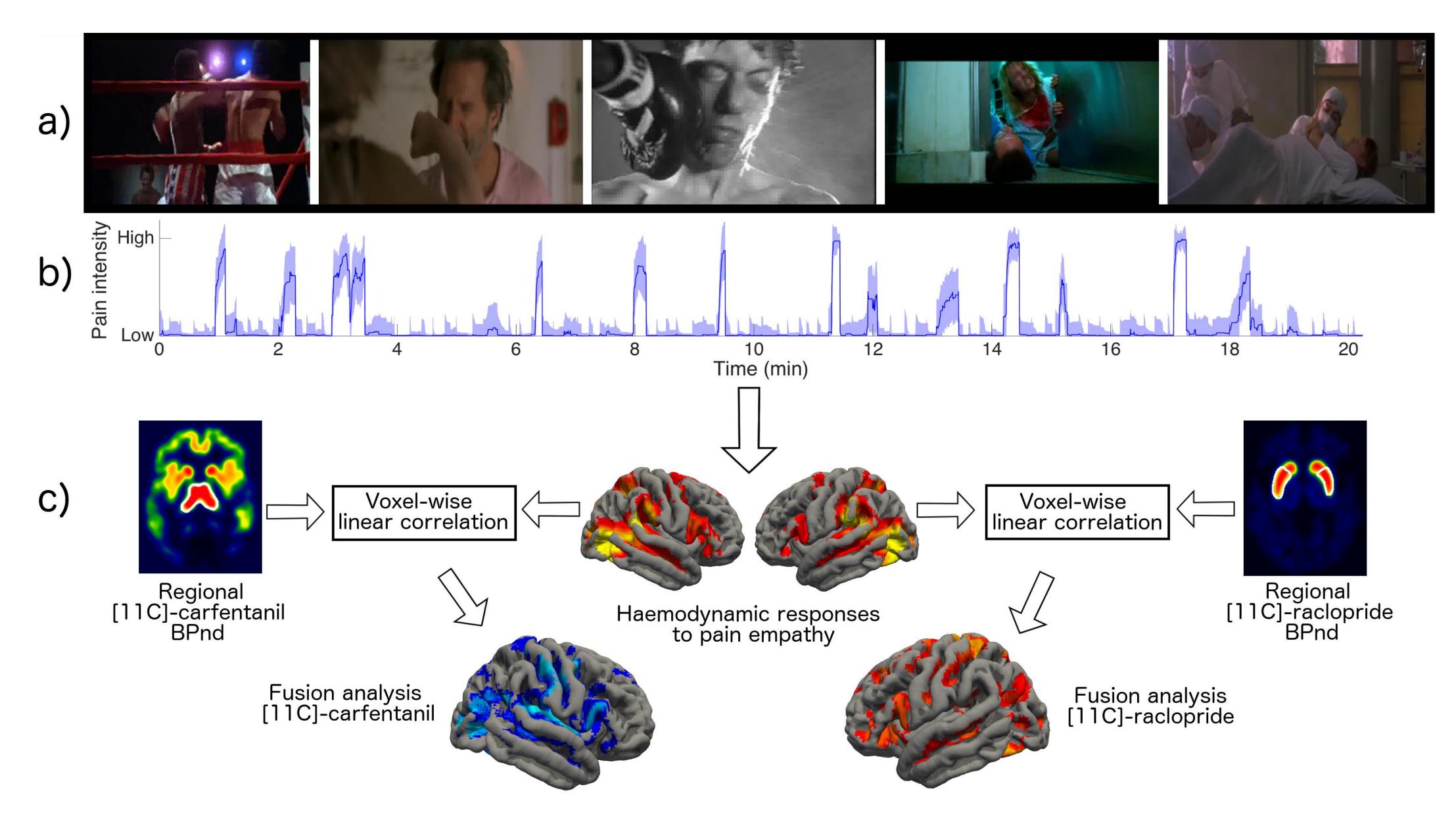
bitis going on tharvir gaue



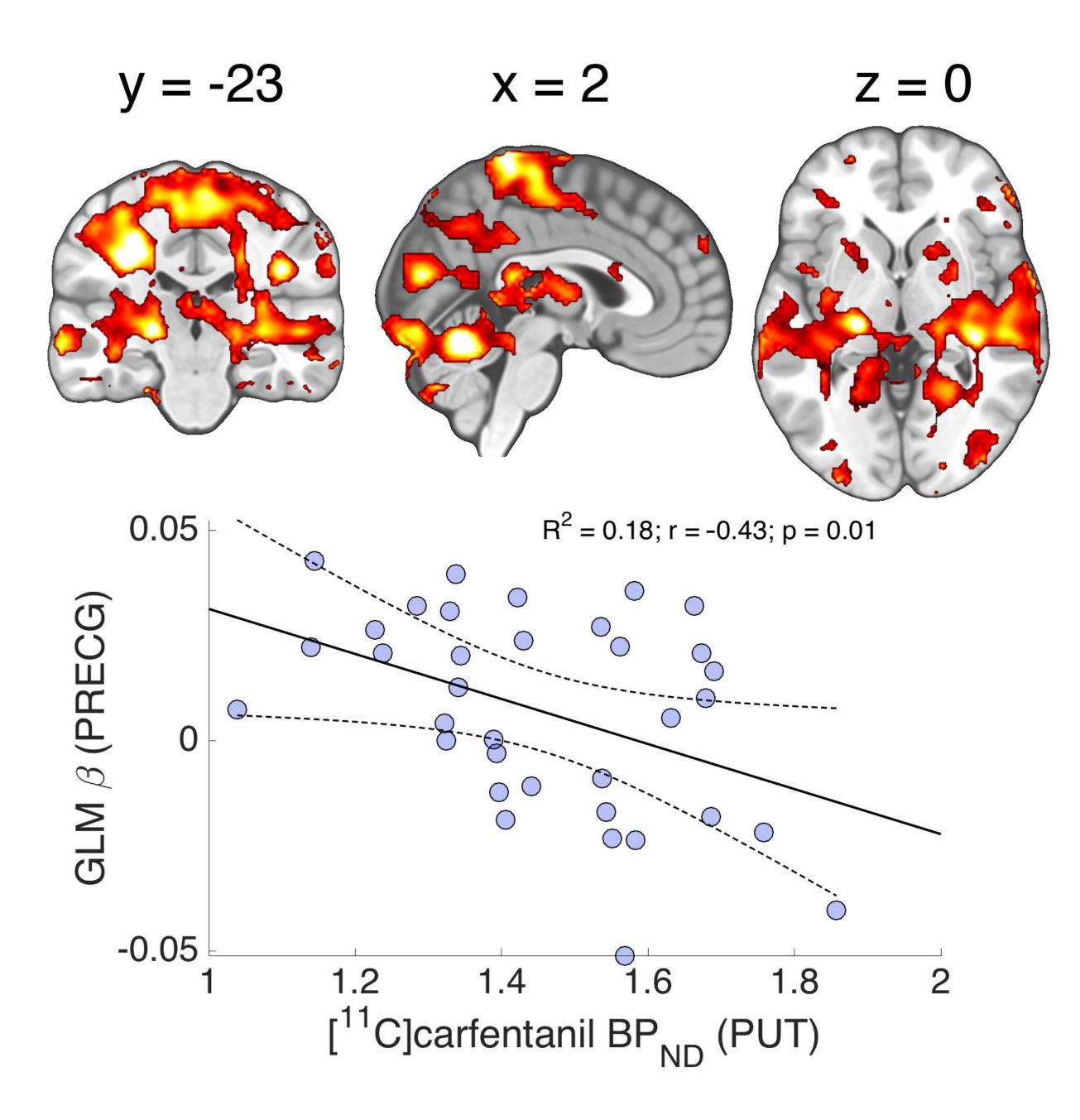
ROI analysis helps in breaking down the interaction







Karjalainen et al (2017 Cereb Cortex)

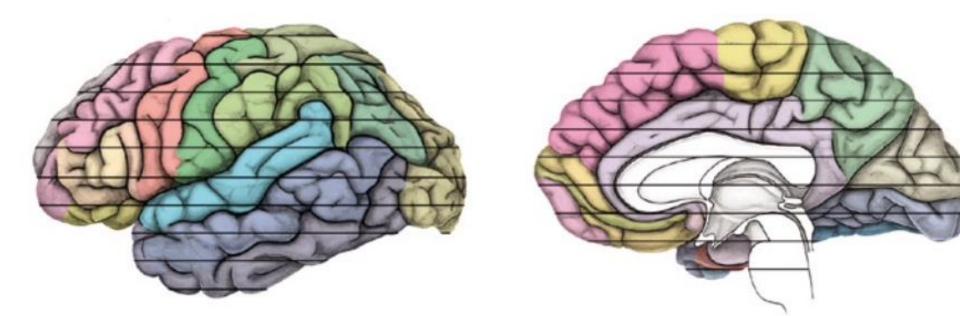


Karjalainen et al (2017; 2019 Cereb Cortex)

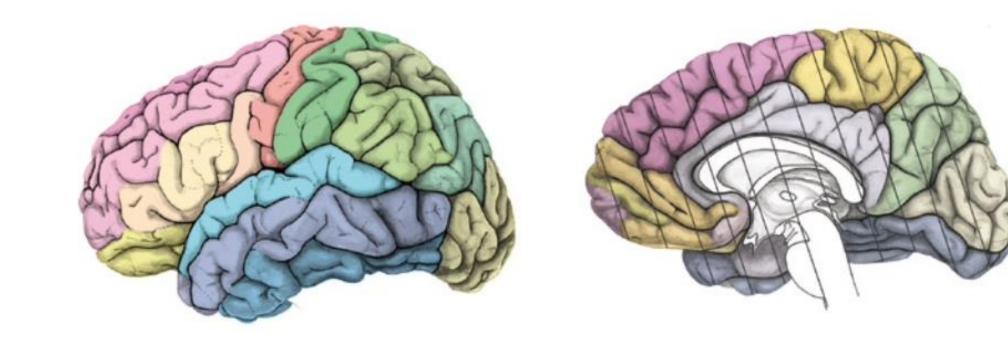


How to define ROIs?

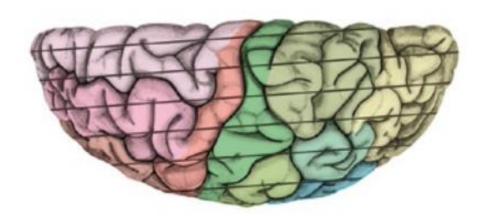
Horizontal Atlas:

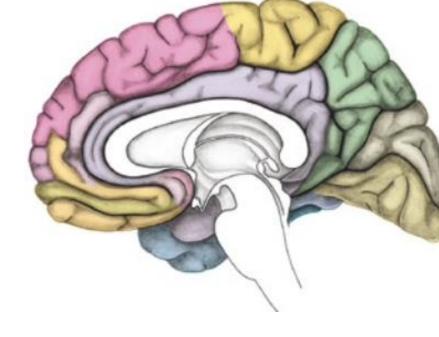


Coronal Atlas:



Sagittal Atlas:



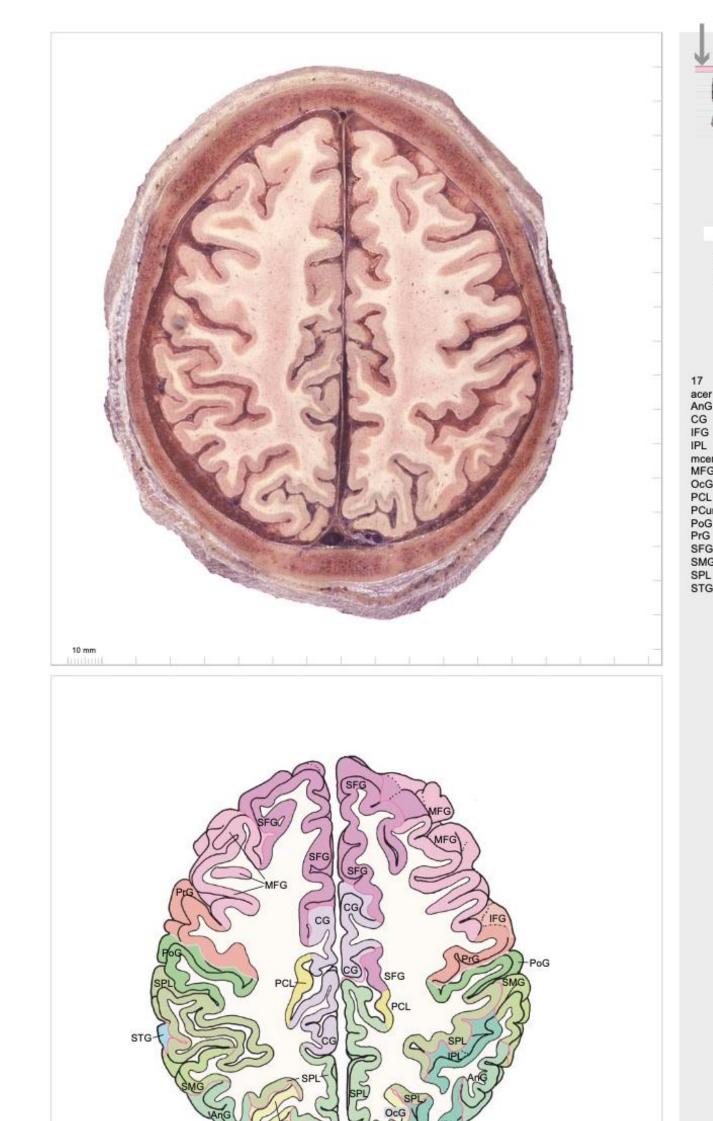


Mai et al (2008)









Geve

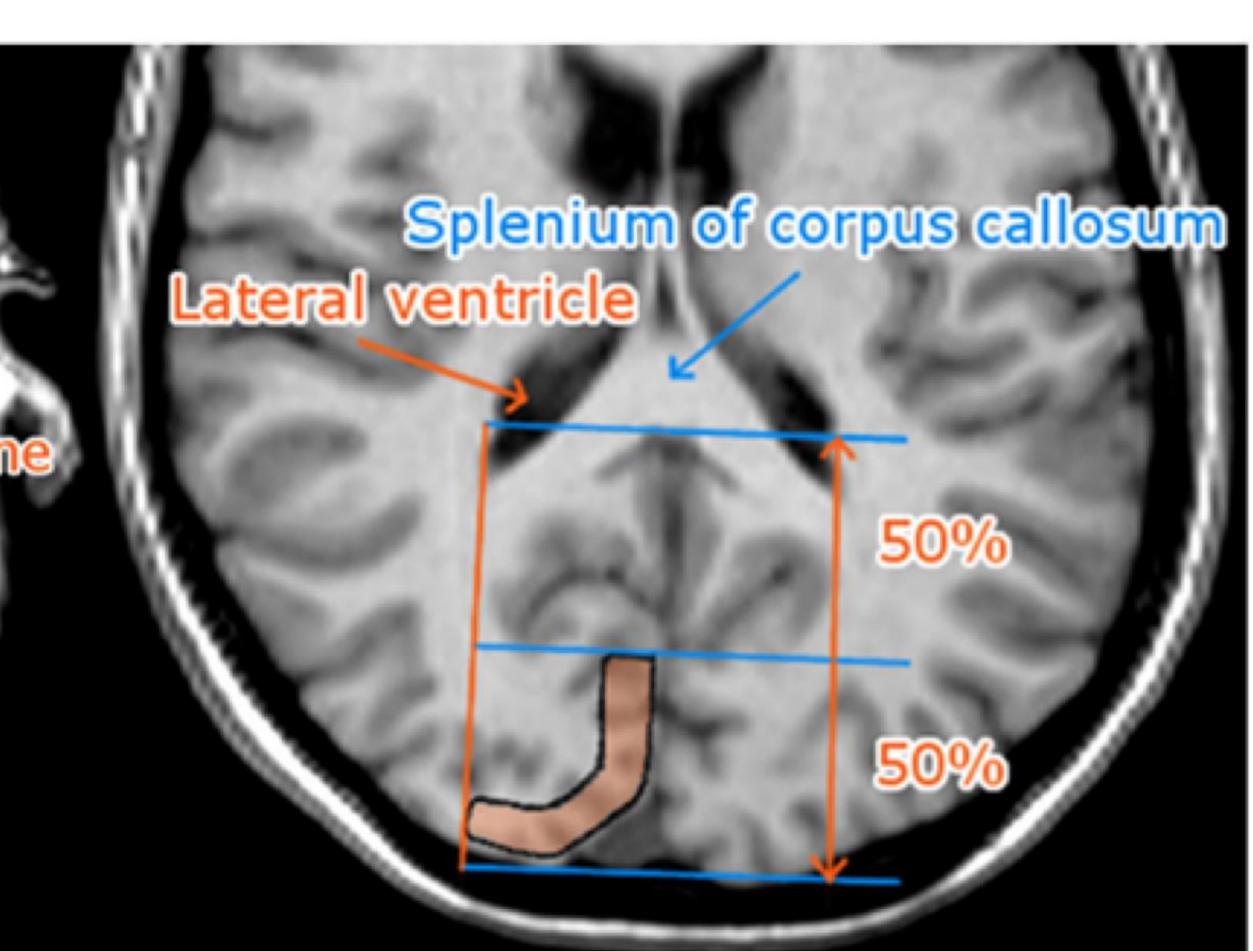
0	(dorsal view
striate area	

cer	anterior cerebral artery	
nG	angular gyrus	
G	cingulate gyrus	
G	inferior frontal gyrus	
۹L	inferior parietal lobule	
cer	middle cerebral artery	
IFG	middle frontal gyrus	
cG	occipital gyri	
CL	paracentral lobule	
Cun	precuneus	
oG	postcentral gyrus	
rG	precentral gyrus	
FG	superior frontal gyrus	
MG	supramarginal gyrus	
PL	superior parietal lobule	
TG	superior temporal gyrus	

Temporal lobe

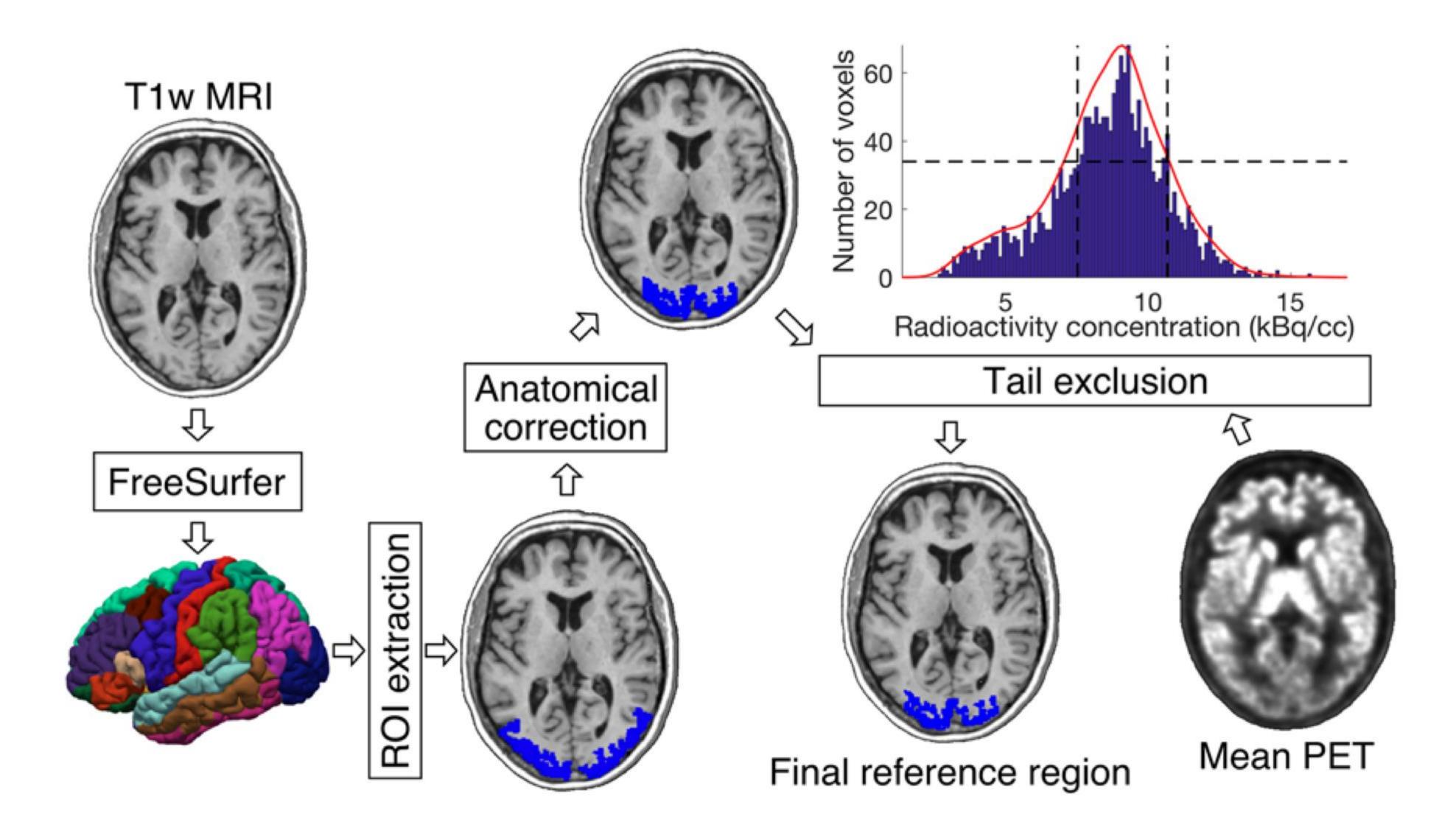
Petrosal part of temporal bone

Venous sinus



Karjalainen et al (2017)

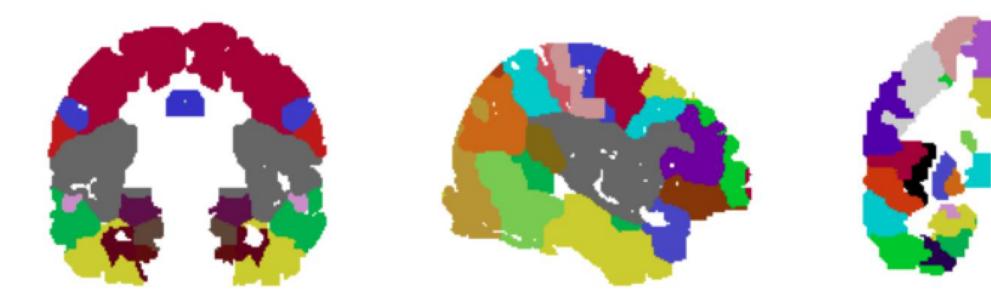




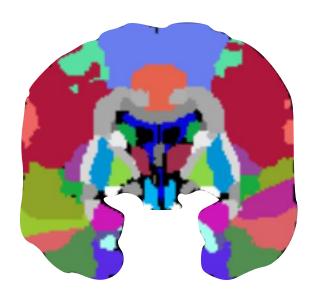
Karjalainen et al (2017)

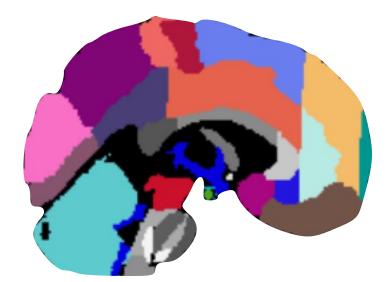
Anatomical and functional parcellations

Brodmann atlas



JHU atlas

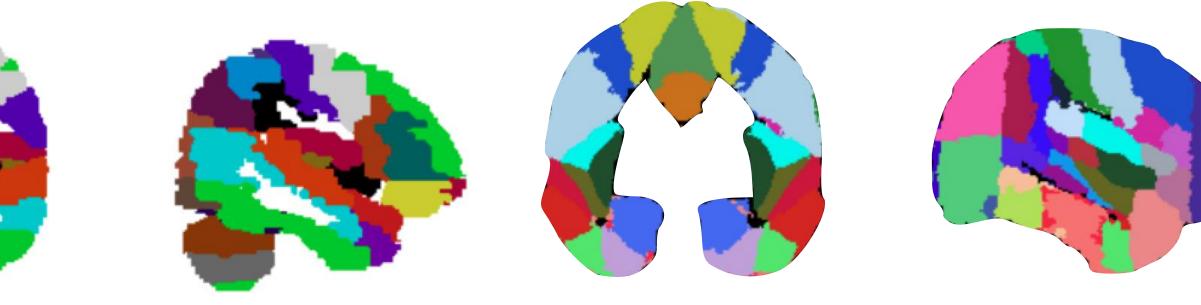






AAL atlas

Harvard-Oxford



AICHA atlas



And many, many more!



Turku PET Centre molecular atlases

Type 2 dopamine receptors



[11C]raclopride Malen et al (2022)

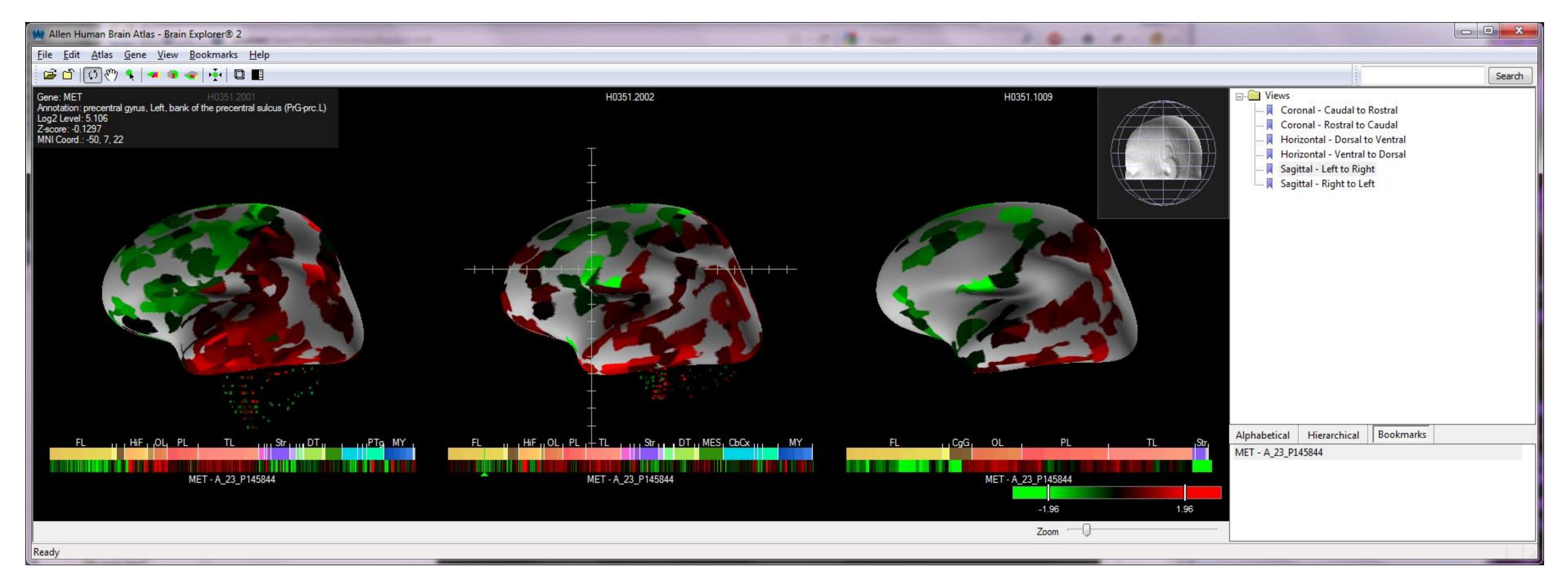
µ-opioid receptors

Serotonin transporters

[11C]carfentanil Kantonen et al (2020)

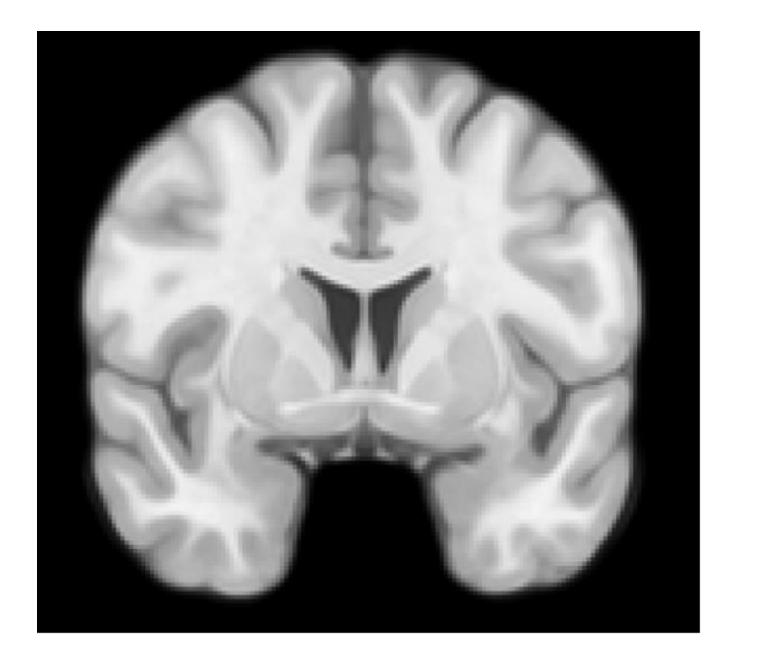
[11C]MADAM Manninen et al (2021)

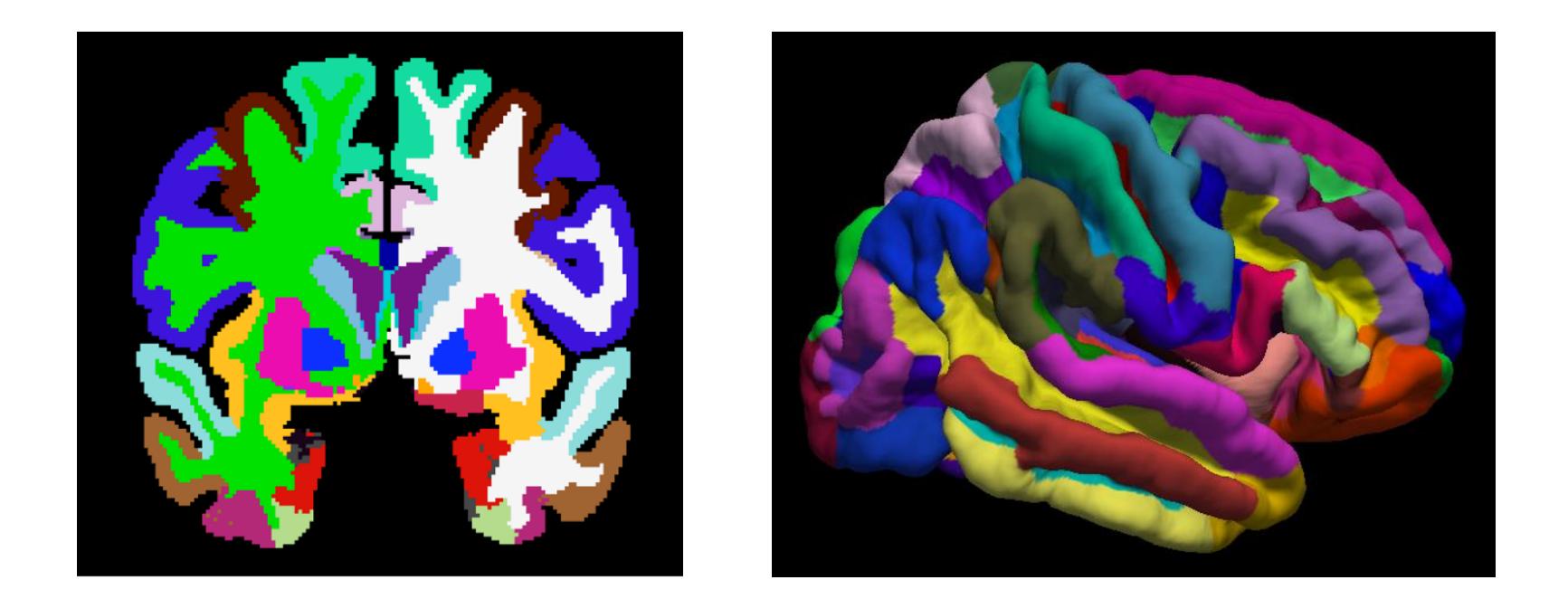
Transcriptomic atlases



http://human.brain-map.org

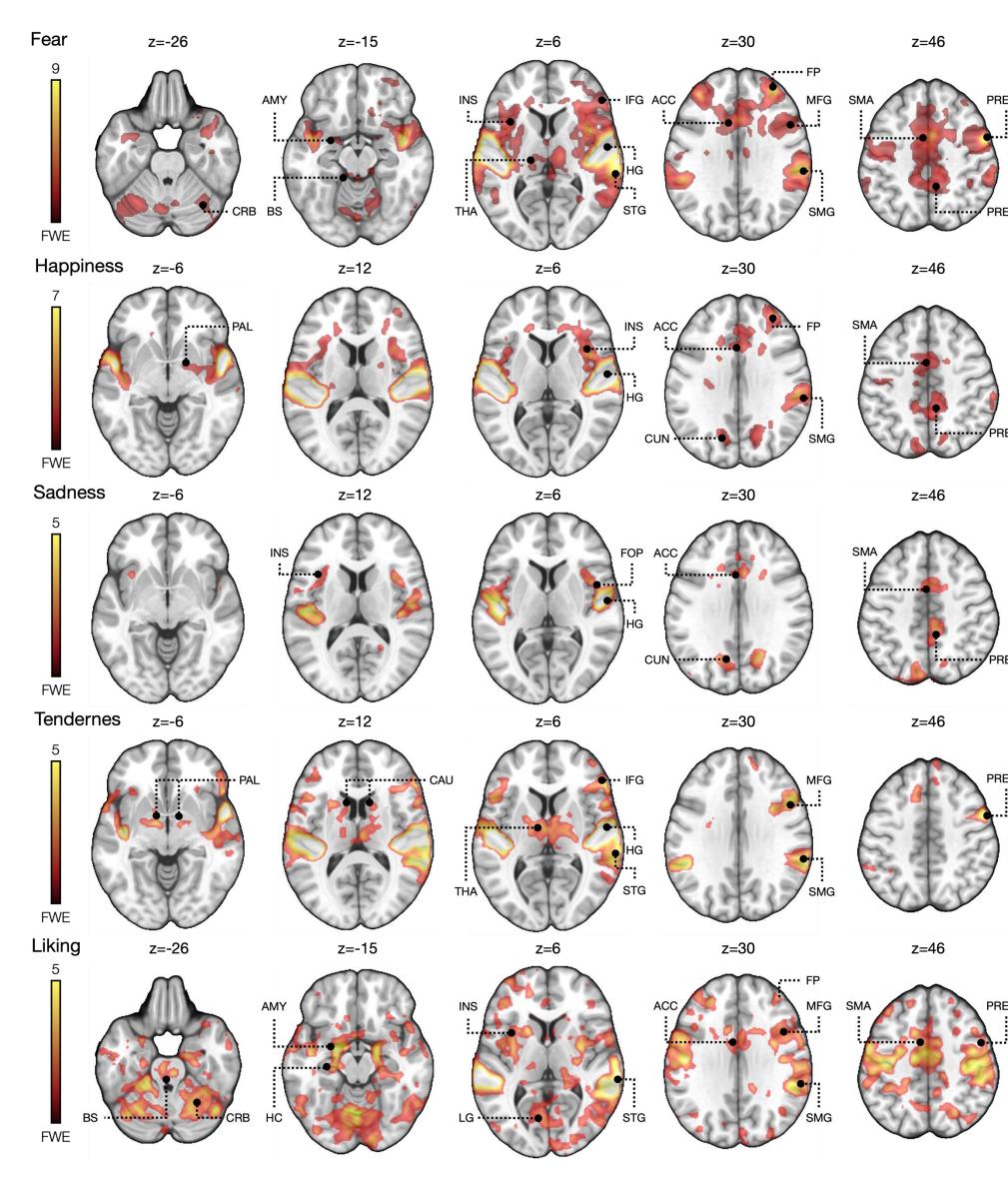
Surfin' U.S.A.



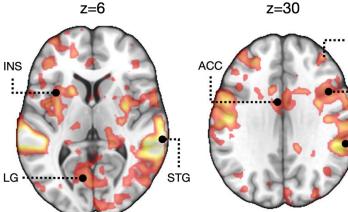


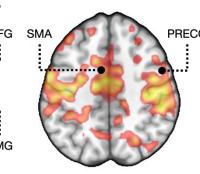
Fressurfer parcellations allow definition of subject-specific and anatomically specific ROIs - however response homogeneity across large ROIs remains an issue

What about functional ROIs?



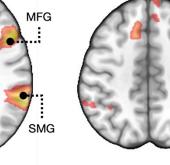
Putkinen et al (2020)



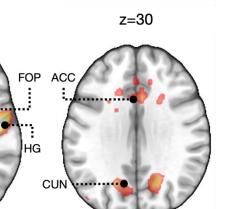


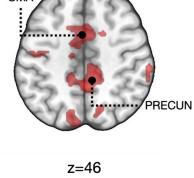
z=46

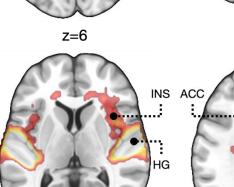
z=30







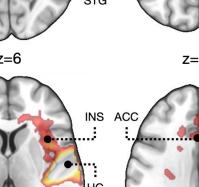


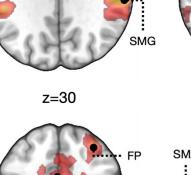


z=6

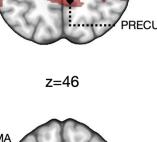
z=6

z=6





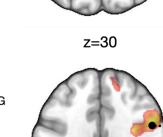
z=30



z=46

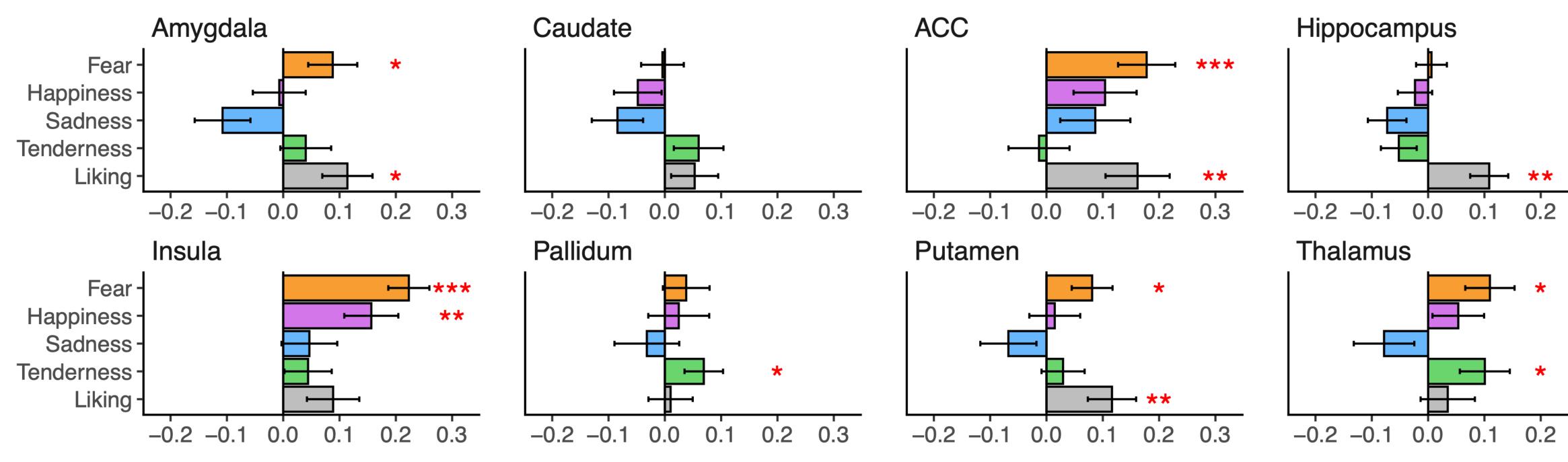
PRECUN

RECUN



z=46

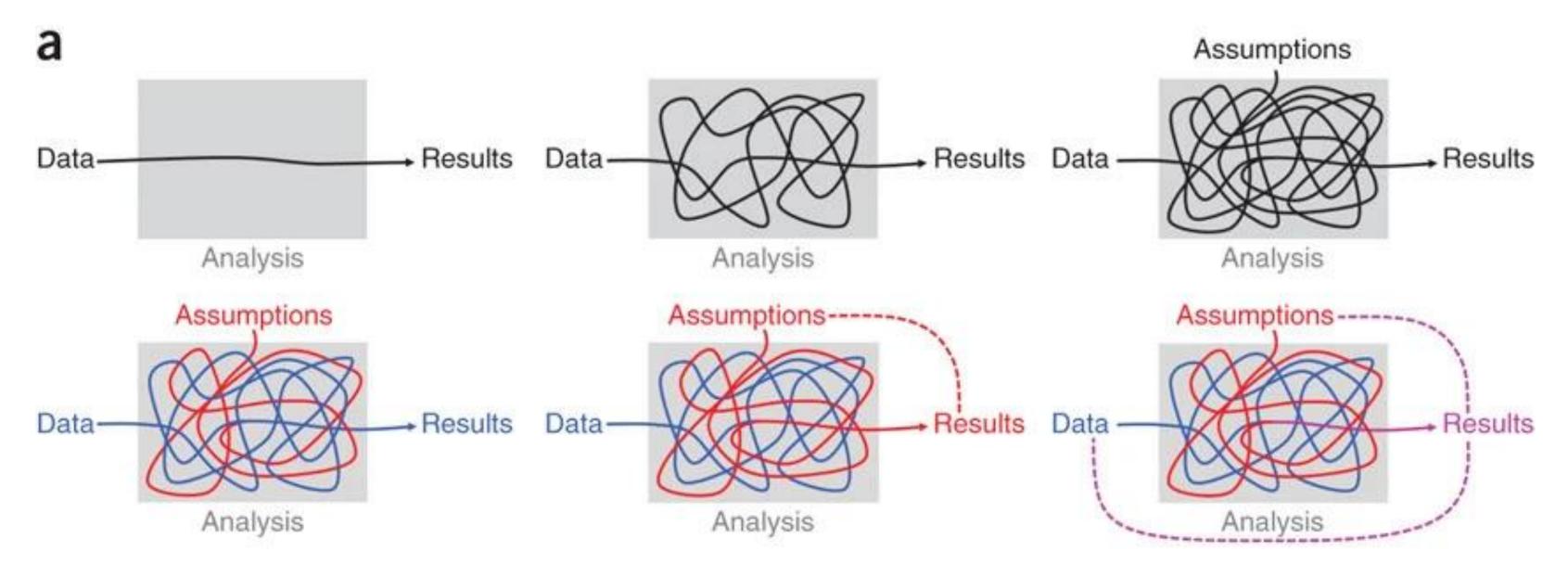
Why can't we analyze regional data for clusters?



Putkinen et al (2020)



Dangers of double dipping





Kriegeskorte et al (2009)

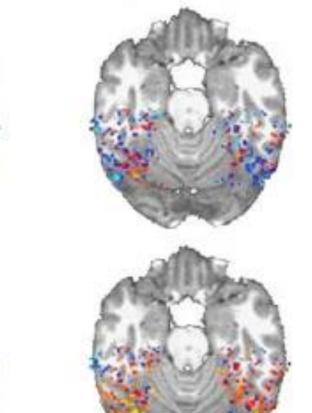
Double dipping in action

а

Task (property judgment)

Animate?

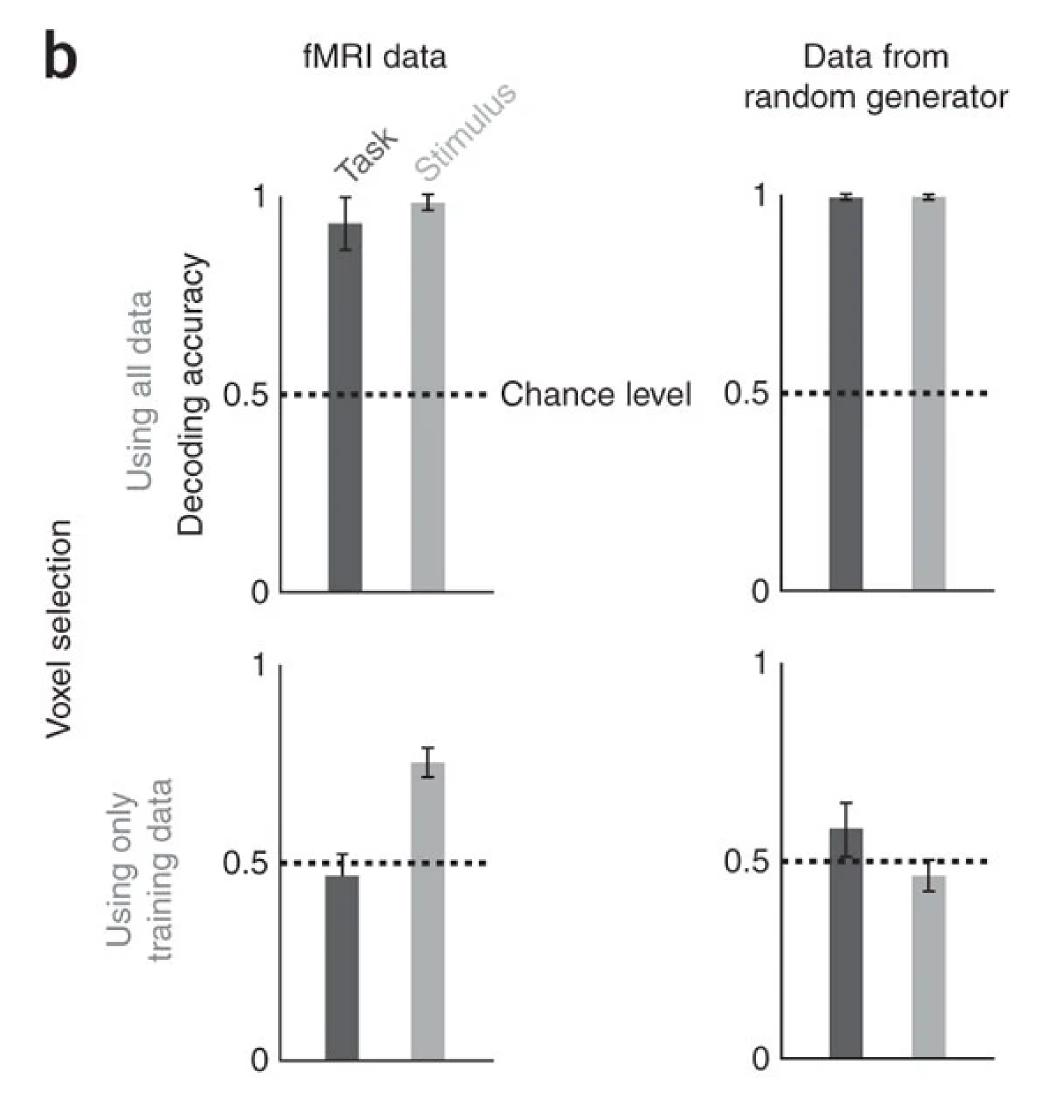
Pleasant?



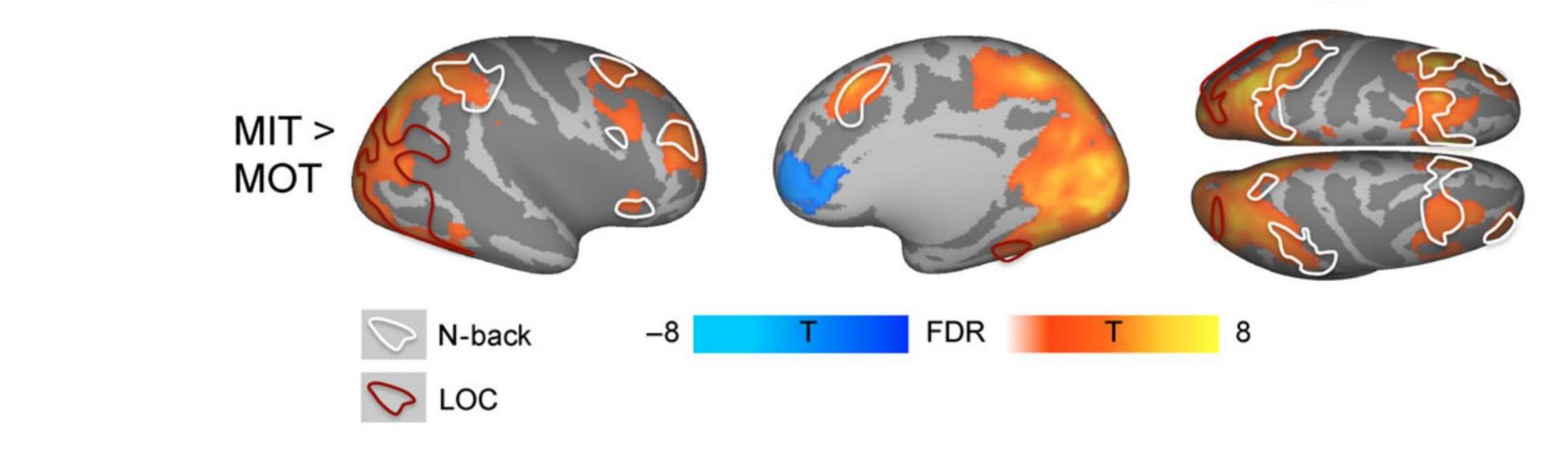
Stimulus (object category)

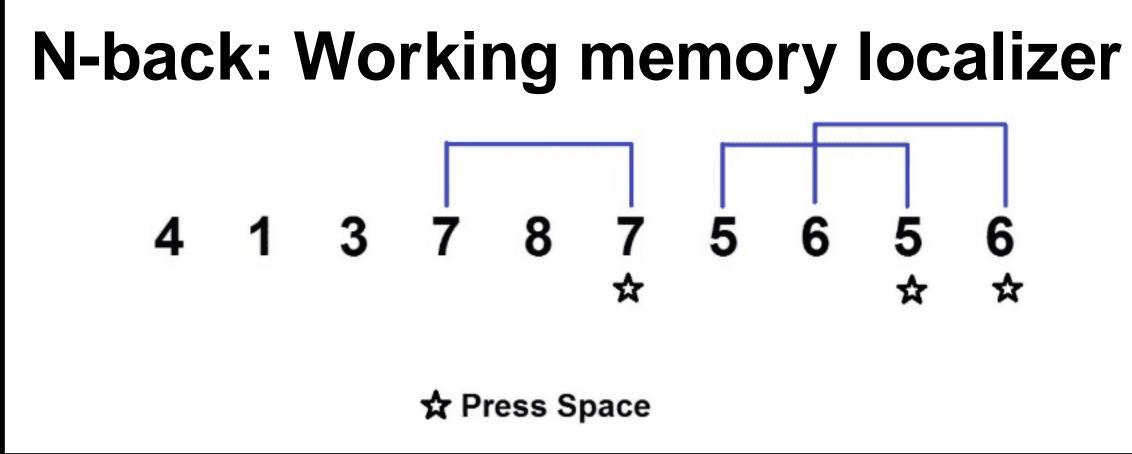


Kriegeskorte et al (2009)



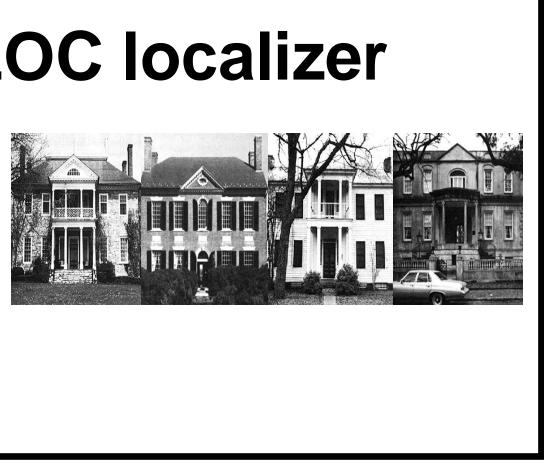
Solution Anatomical ROIs and functional localizers





Picture task: LOC localizer





Nummenmaa et al (2017)

How to extract?

- MarsBaR toolbox allows extracting regional data from already estimated 1st and 2nd level SPM models
- ROI definition also supported
- Useful for extracting %signal changes or regional PET outcome measures (BPND, SUVR...)
- A model has to exist to extract data, you may need to set up e.g. dummy 2nd level model testing voxel values against zero
- Note in the end really easy to use, although confusing to start with

🚺 MarsBaR - Marseille R 💶 🗙
ROI definition 🔻
■ Design
■ Data
Results 🕶
Options 🕶
Quit

How to extract?

- NIFTI is a well-standardised image format
- Numerous packages allow accessing and manipulating NIFTI structures
- Often the easiest way for ROI data extraction is simply to define your mask in the same space as the target image and just work on the data matrices

Edit View Zoom on Interp off 🛷 Image Information × Dimension: [91 109 91] Voxel Size: [2.000 2.000 2.000] Data Type: [32-bit single float] Max Value: [100] == X => <= Y == Min Value: [-99.9999] Brightness: Reset rast: Reset File Name: [T1.nii] Axes Unit Colormap OK Negative Bipolar Color On Crosshair: 46 64 37 [XYZ] at origin: 62 48 46 [XYZ] at crosshair: 21.0744 Value at crosshair: [XYZ] at cursor:

Value at cursor:

