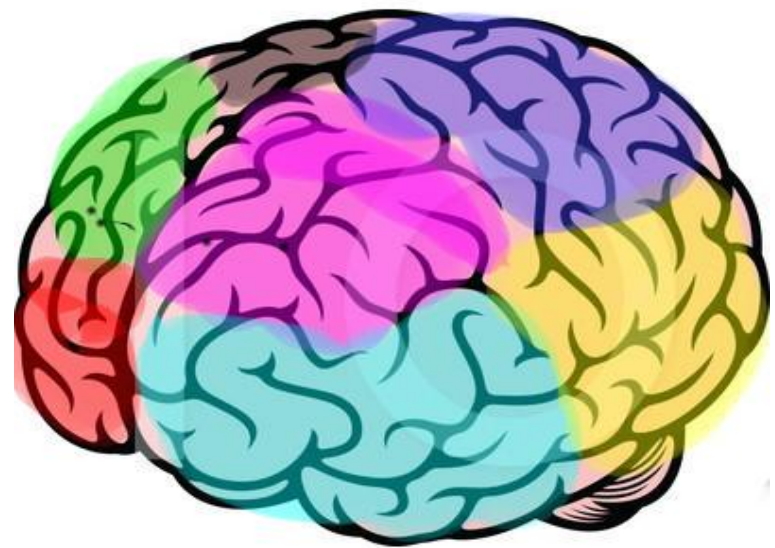
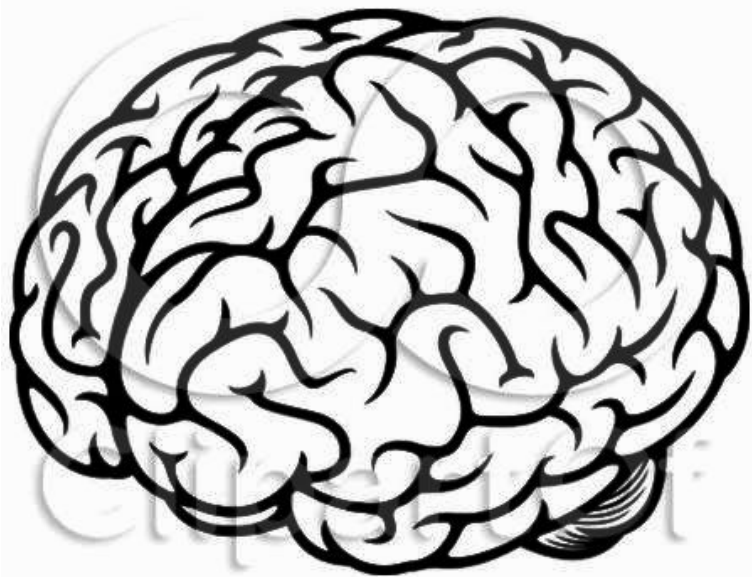
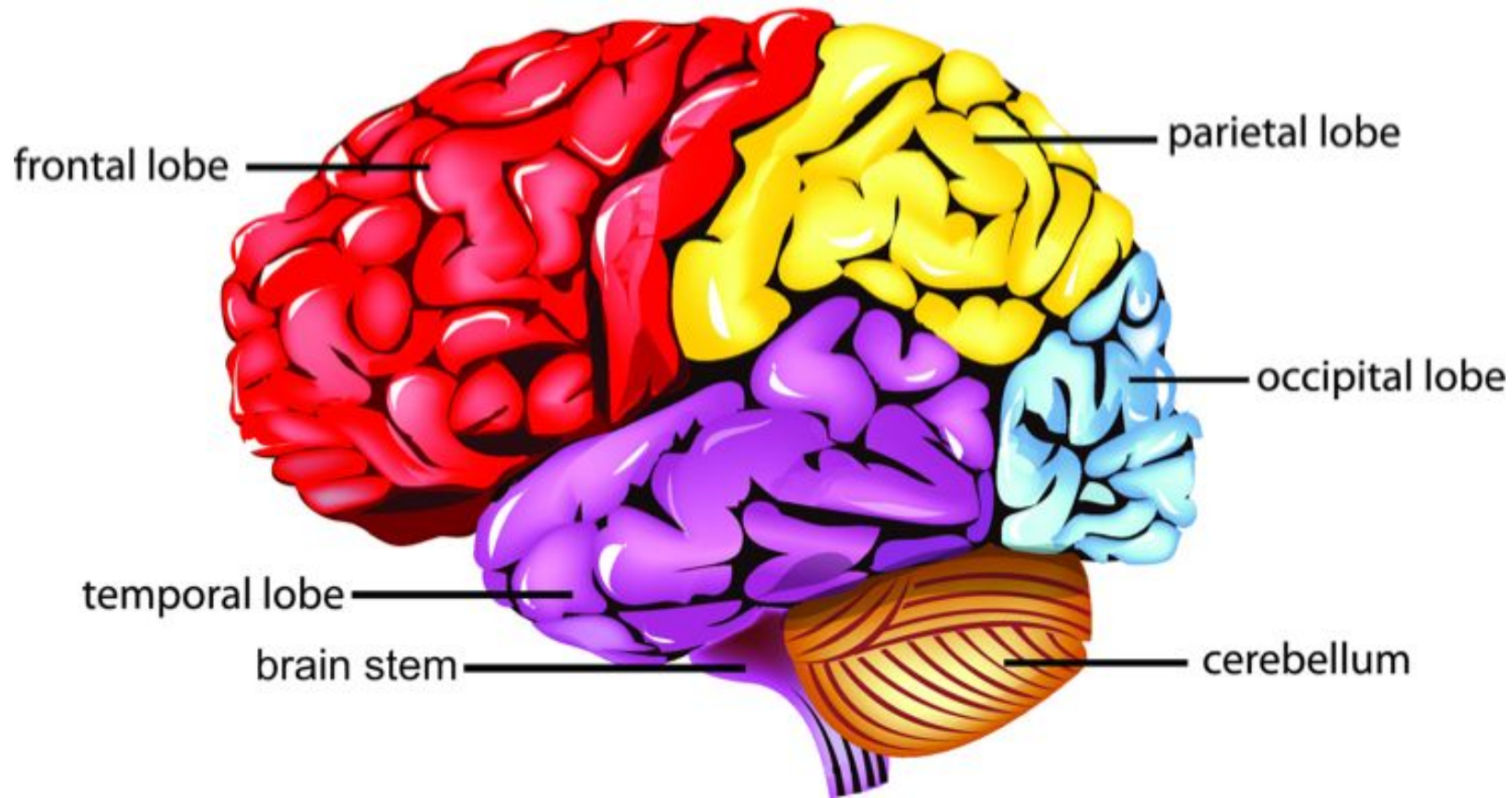


MRI based brain parcellation

Kurmukov Anvar, 2019

1. What is brain
parcellation?





frontal lobe

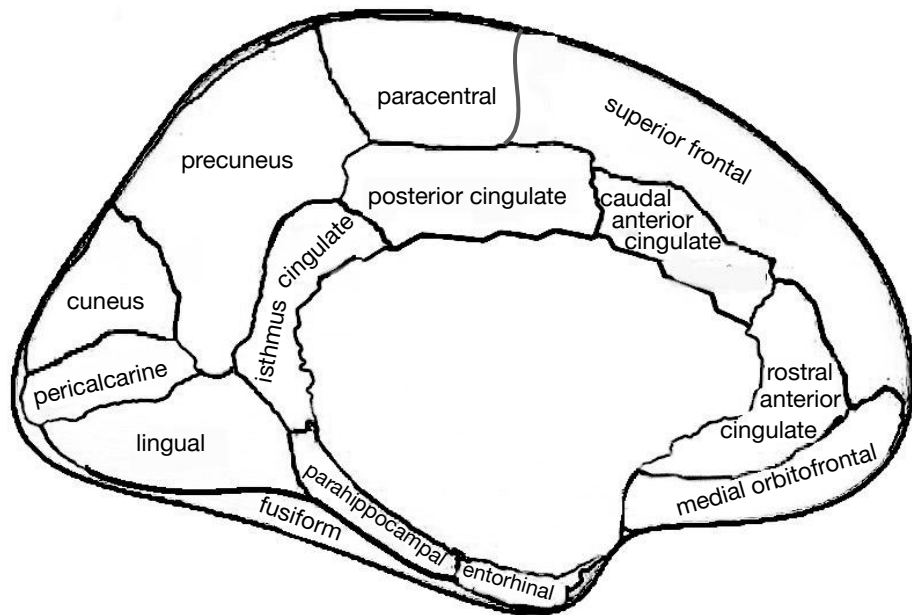
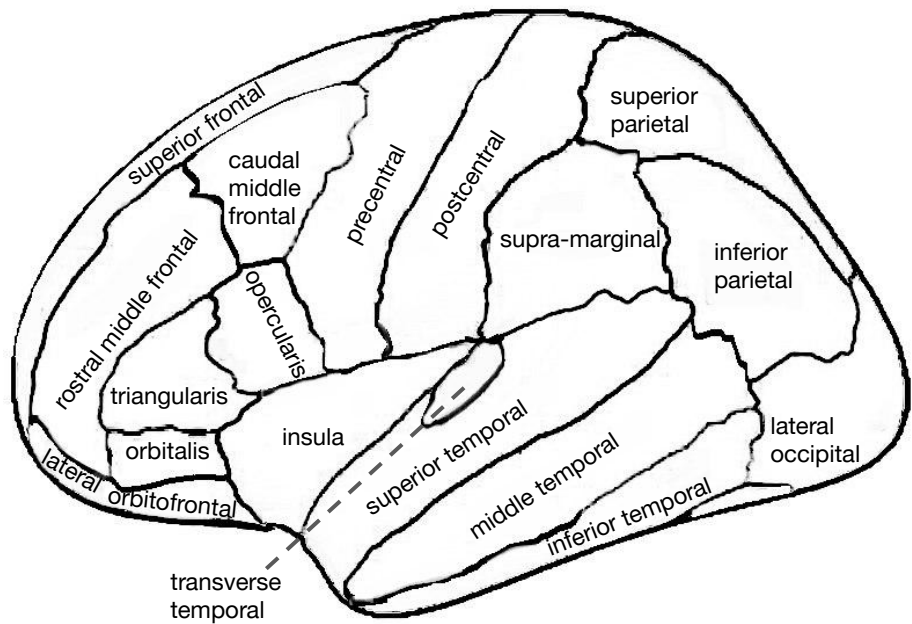
parietal lobe

occipital lobe

temporal lobe

brain stem

cerebellum

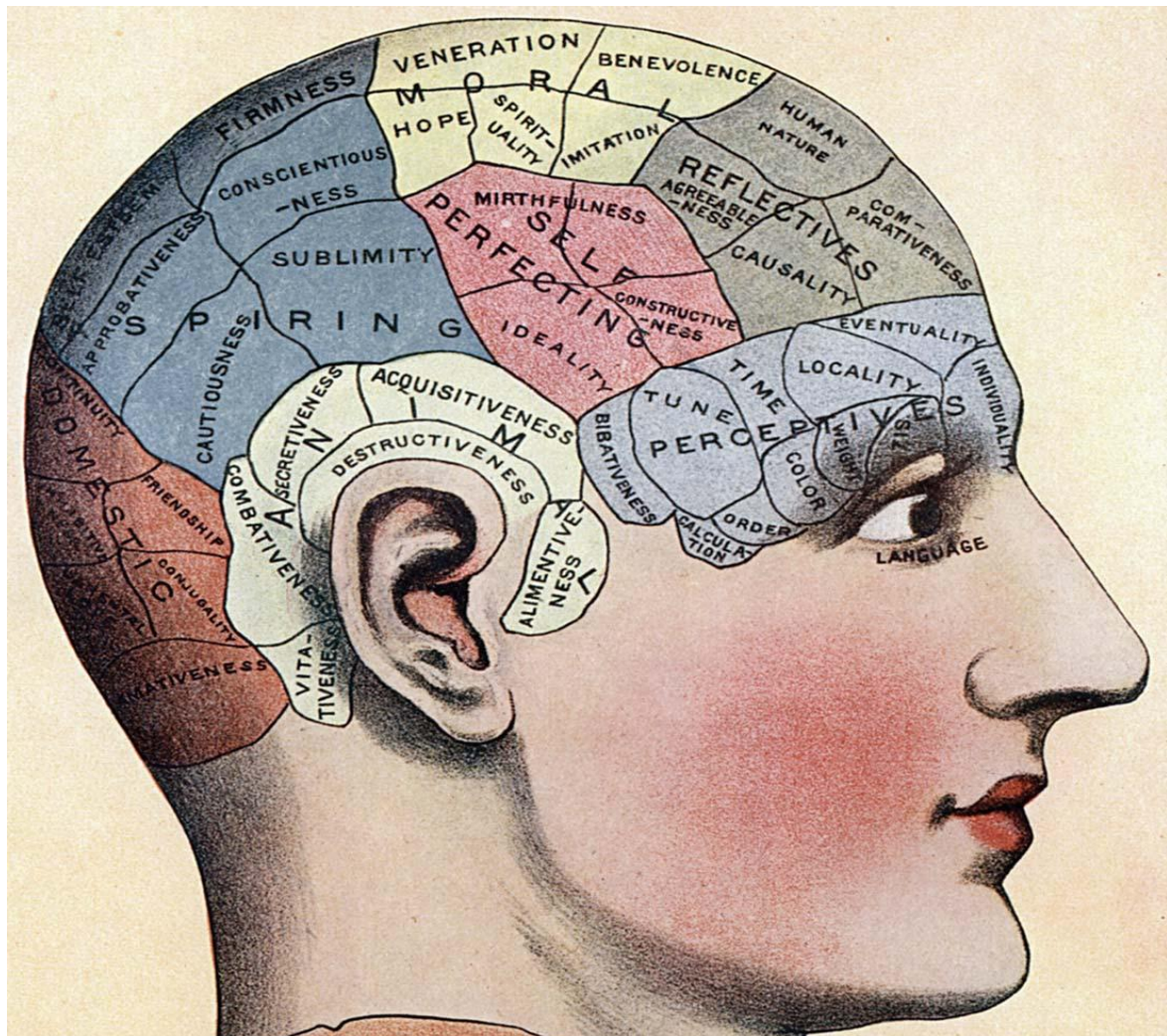


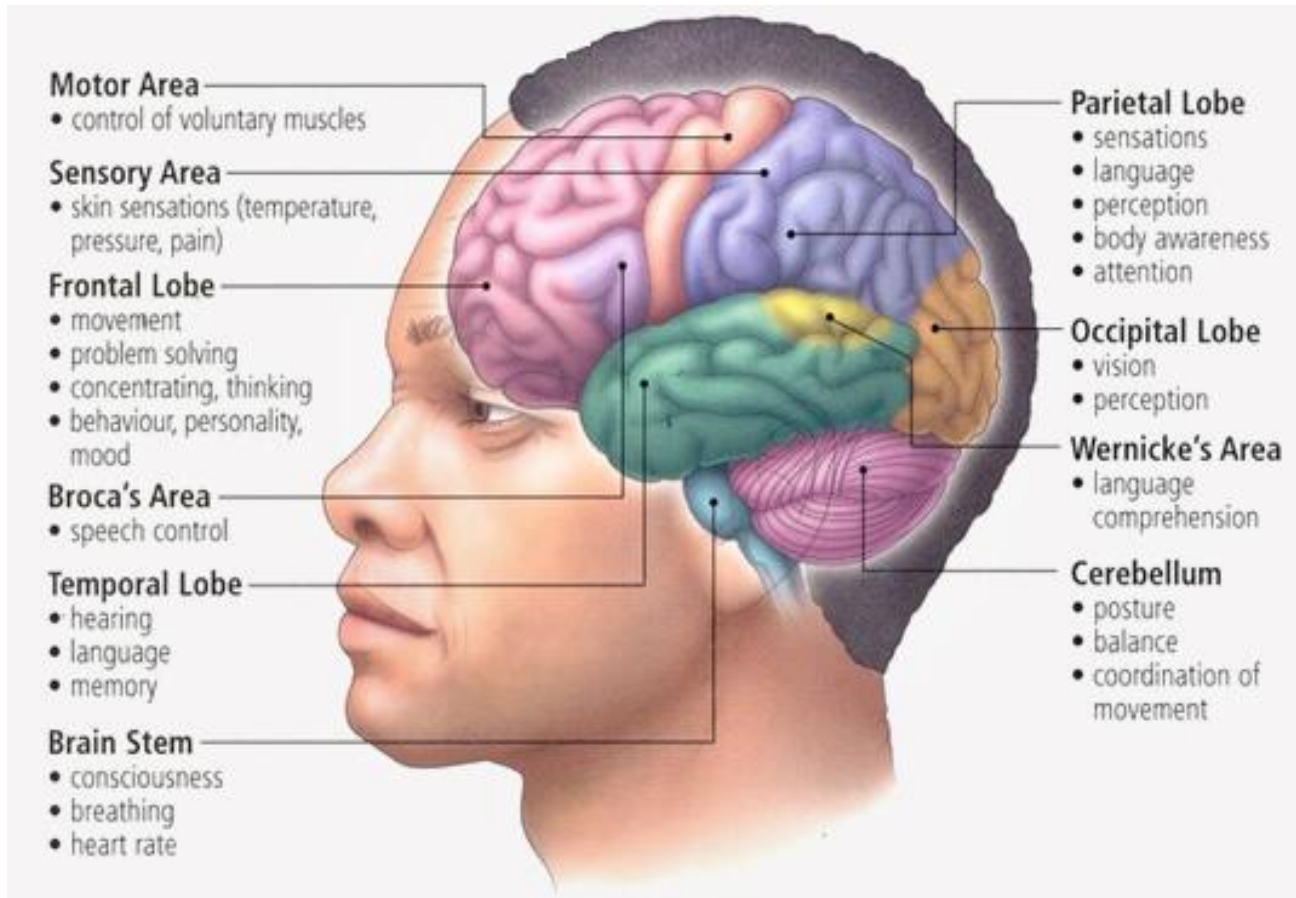
In computational neuroimaging, brain parcellation methods subdivide the brain into individual regions <...> to study its **structure and function.**

[Brain parcellation based on information theory](#)

2. Why do we need
parcellations?

1. Because brain structure is somehow related to its function.
2. Because typical MRI consists of $\sim 10^5$ up to $\sim 10^7$ voxels and typical study has $\sim 10^1$ up to $\sim 10^3$ observations (dimensionality reduction).
3. Because in multimodal studies we need to have 1 to 1 correspondence between different modalities.
4. Because in population studies we need to have 1 to 1 correspondence between subjects.
5. To build connectomes;)





Motor Area

- control of voluntary muscles

Sensory Area

- skin sensations (temperature, pressure, pain)

Frontal Lobe

- movement
- problem solving
- concentrating, thinking
- behaviour, personality, mood

Broca's Area

- speech control

Temporal Lobe

- hearing
- language
- memory

Brain Stem

- consciousness
- breathing
- heart rate

Parietal Lobe

- sensations
- language
- perception
- body awareness
- attention

Occipital Lobe

- vision
- perception

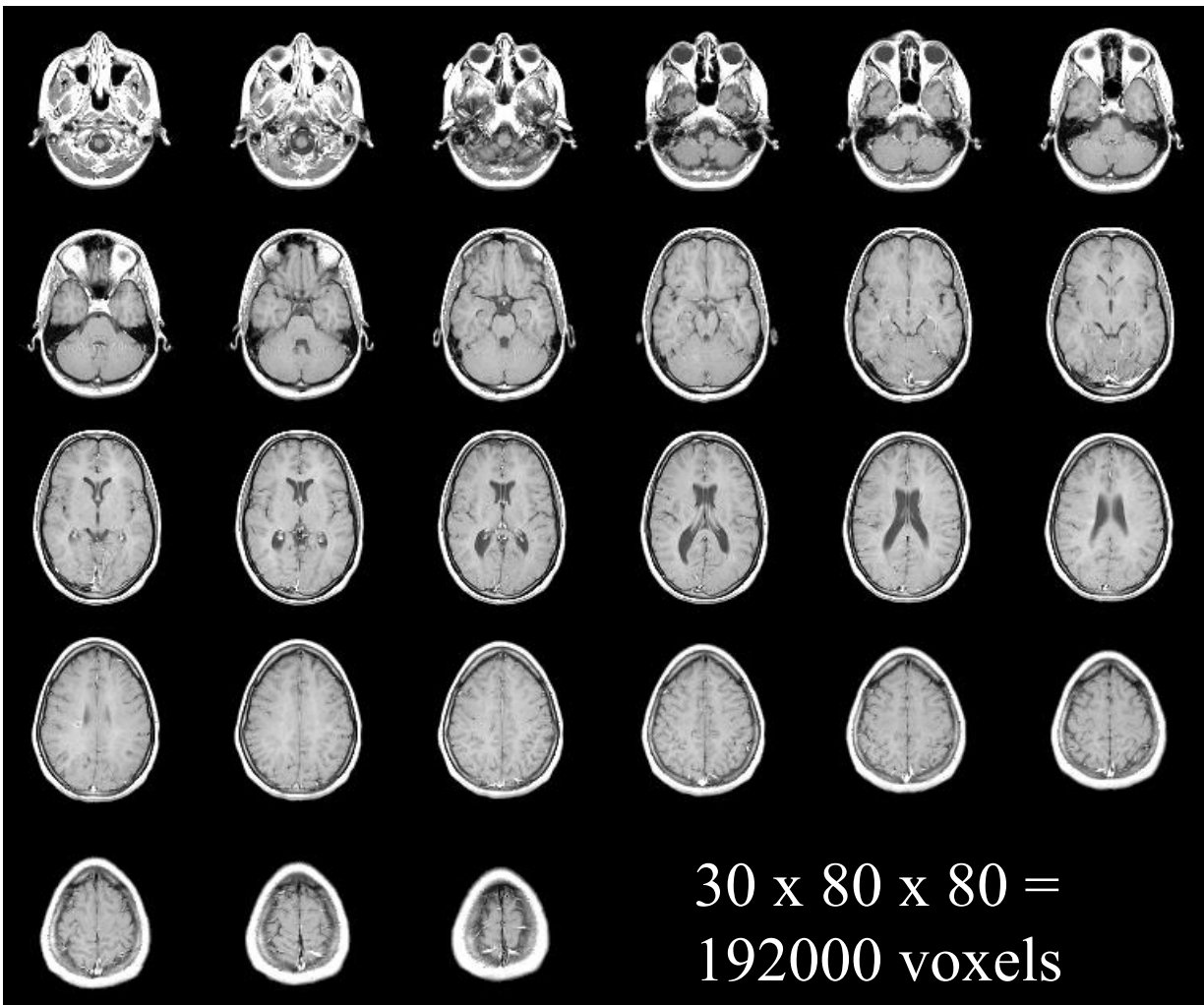
Wernicke's Area

- language comprehension

Cerebellum

- posture
- balance
- coordination of movement

1. Because brain structure is somehow related to its function.
2. Because typical MRI consists of $\sim 10^5$ up to $\sim 10^7$ voxels and typical study has $\sim 10^1$ up to $\sim 10^3$ observations (dimensionality reduction).
3. Because in multimodal studies we need to have 1 to 1 correspondence between different modalities.
4. Because in population studies we need to have 1 to 1 correspondence between subjects.
5. To build connectomes;)



$30 \times 80 \times 80 =$
192000 voxels



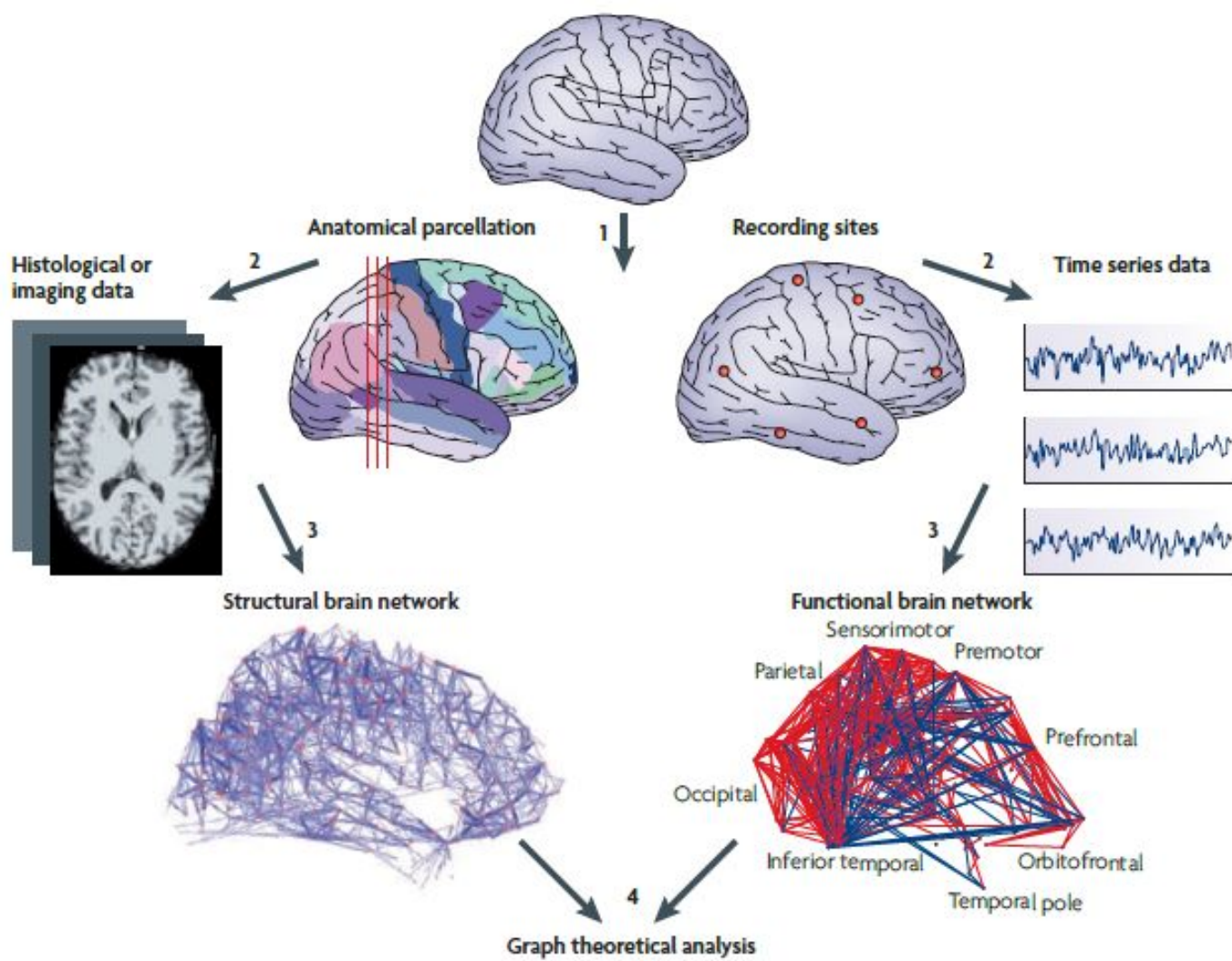
Matrix size of slice:
64 x 64

Voxel size:
3mm x 3mm x 3mm

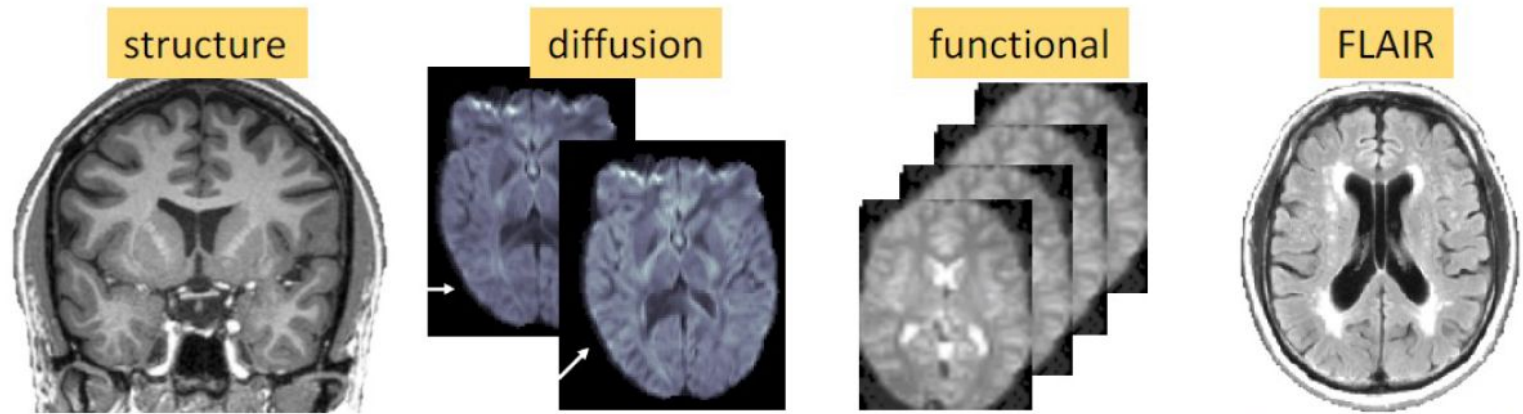


**Disclaimer: voxels are
not necessarily squared!**

1. Because brain structure is somehow related to its function.
2. Because typical MRI consists of $\sim 10^5$ up to $\sim 10^7$ voxels and typical study has $\sim 10^1$ up to $\sim 10^3$ observations (dimensionality reduction).
- 3. Because in multimodal studies we need to have 1 to 1 correspondence between different modalities.**
4. Because in population studies we need to have 1 to 1 correspondence between subjects.
5. To build connectomes;)

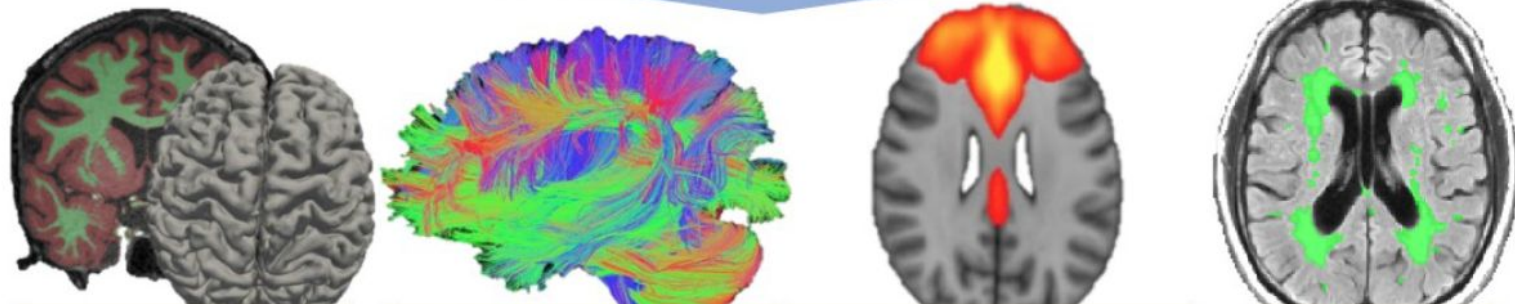


Source is lost ;(



Fully automated processing pipelines
on HPC clusters

unit: subject



ANALYSES

unit: features

1. Because brain structure is somehow related to its function.
2. Because typical MRI consists of $\sim 10^5$ up to $\sim 10^7$ voxels and typical study has $\sim 10^1$ up to $\sim 10^3$ observations (dimensionality reduction).
3. Because in multimodal studies we need to have 1 to 1 correspondence between different modalities.
4. Because in population studies we need to have 1 to 1 correspondence between subjects.
5. To build connectomes;)

Narrow-sense heritability: $\Omega = 2 \cdot \Phi \cdot \sigma_g^2 + I \cdot \sigma_e^2$

$\text{Var}(y)\sigma_p^2 = \Omega$ – pedigree covariance, Φ – kinship matrix
 σ_g^2 – genetic variance, σ_e^2 – environmental variance

*Typically, done in family/twin studies

$$\sigma_p^2 = \sigma_g^2 + \sigma_e^2$$

$$h^2 = \sigma_g^2 / \sigma_p^2$$



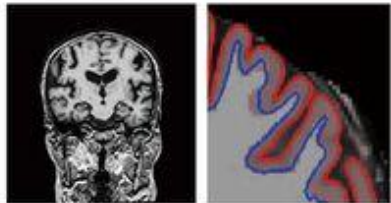
Interpretation:

Relative importance of **genetics vs. environment** for a given trait

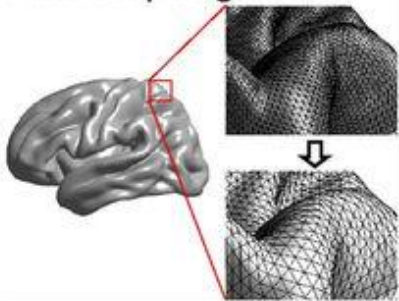
[Heritability of the shape of subcortical brain structures in the general population](#)

Image preprocessing

Freesurfer

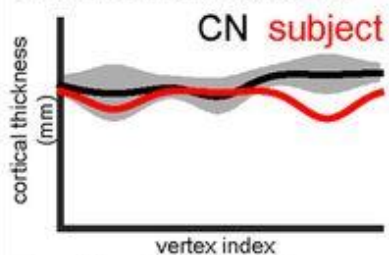


Resampling

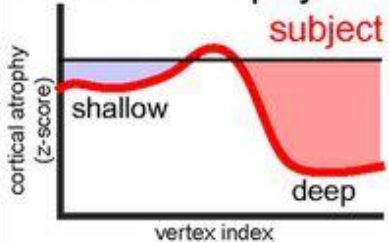


Computing cortical atrophy map

Cortical thickness

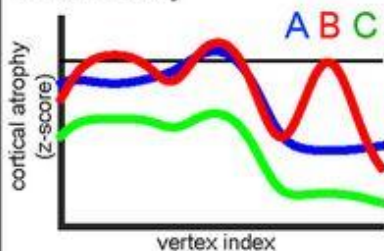


Cortical Atrophy



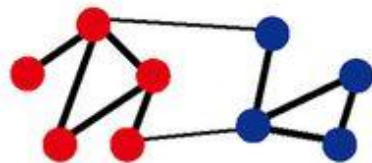
Subtyping based on similarity

Similarity



$$Corr_{AB} < Corr_{AC}$$

Modularity

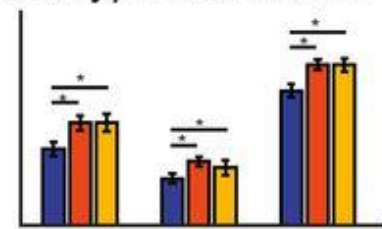


Statistical analysis

Atrophy pattern



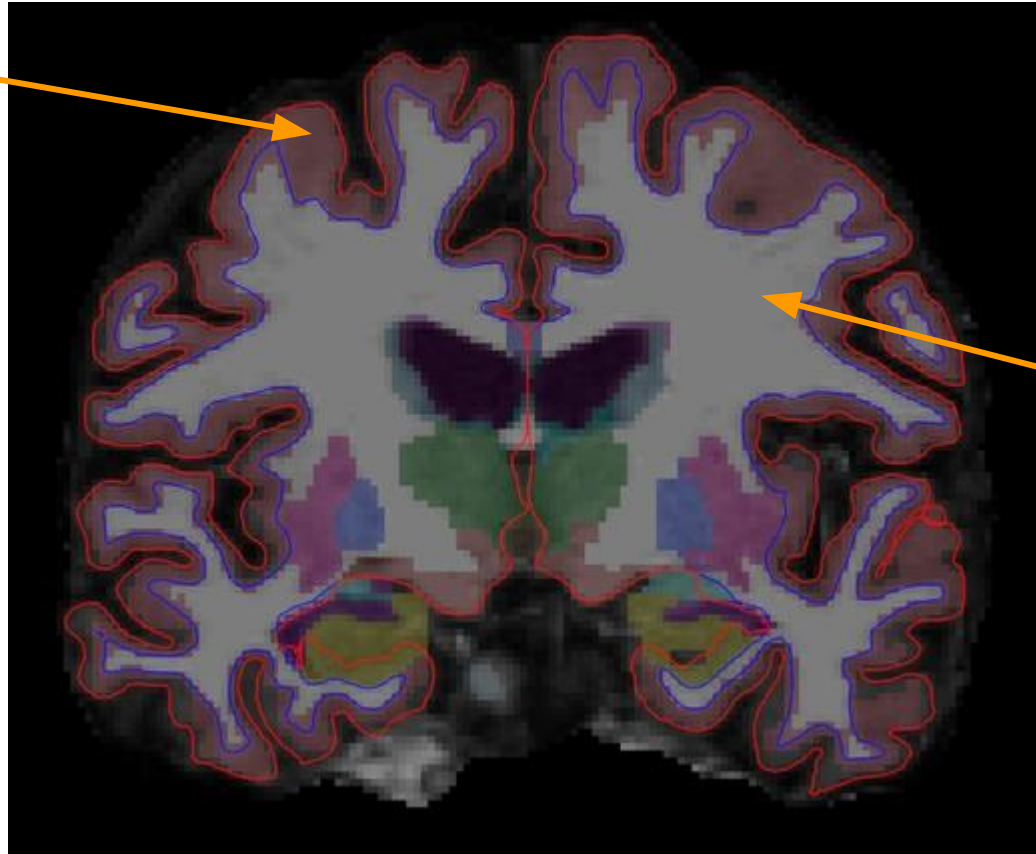
Subtype Hallmarks



[Robust Identification of Alzheimer's Disease subtypes based on cortical atrophy patterns](#)

Gray
matter

Freesurfer
segmentation

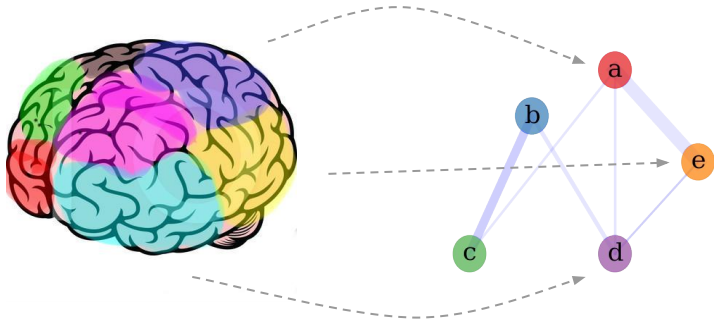


White
matter

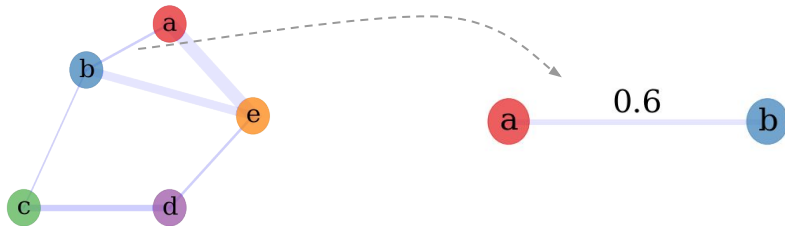
1. Because brain structure is somehow related to its function.
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5. To build connectomes;)

Brain network = Connectome

Brain regions become nodes



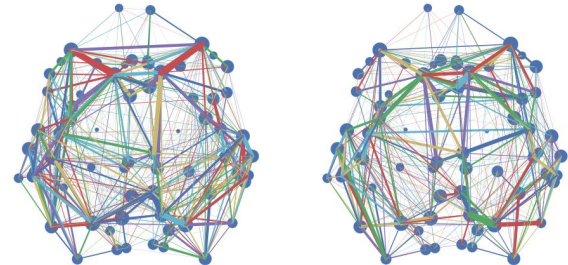
Neural connections between regions become edges



Graph $G = (V, E, l, w)$, where

- V is the set of nodes
- E is the set of edges
- l is node's labeling mapping
- w is edge's weighting mapping

is called a brain network or a **connectome**



[Classification of Normal and Pathological Brain Networks Based on Similarity in Graph Partitions](#)

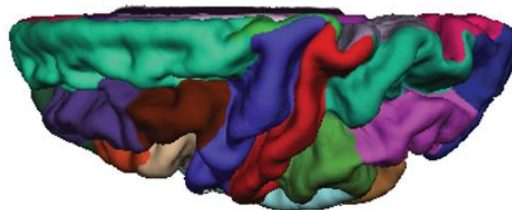
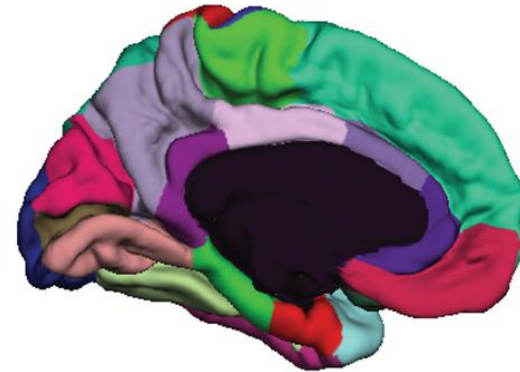
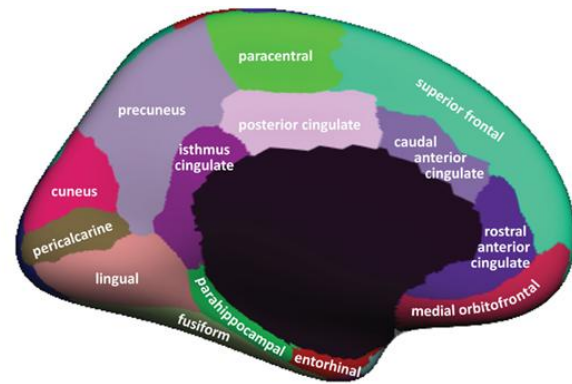
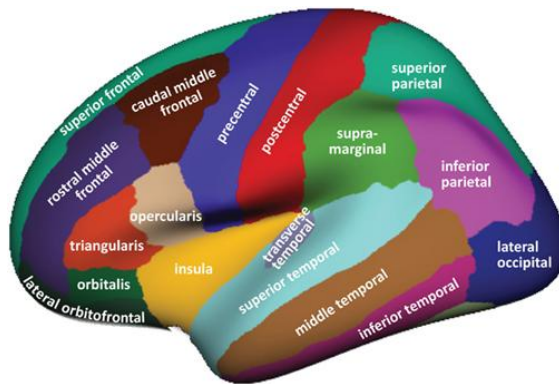
3. Anatomical parcellations

Cortical parcellations = parcellation of the brain surface, popular examples are:

1. [Destrieux Atlas](#)
2. [Desikan-Killiany Atlas](#)
3. [DKT Atlas](#)
4. [Lausanne atlas](#)
5. [Harvard-Oxford atlas](#)
6. [Automated Anatomical Labeling](#)
7. [more](#)

101 labeled brain images
and a consistent human
cortical labeling protocol

They used DKT protocol.
Manual segmentation of
anatomical areas



Lausanne atlas



83 ROI



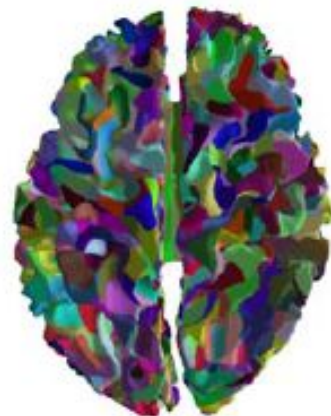
129 ROI



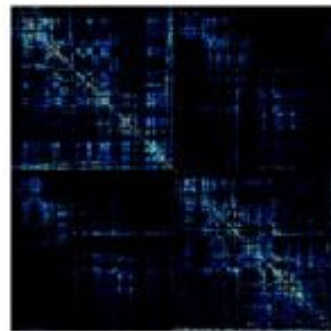
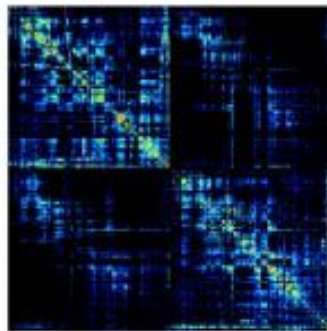
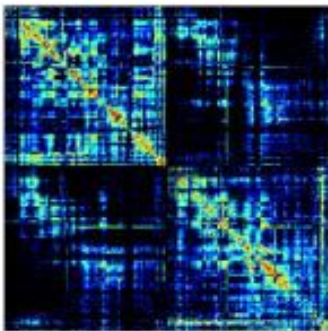
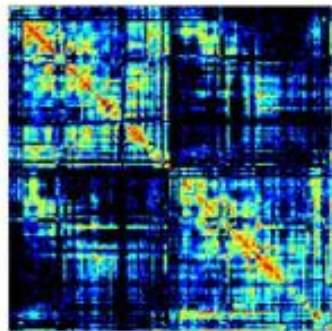
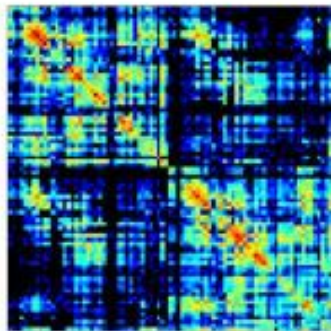
234 ROI



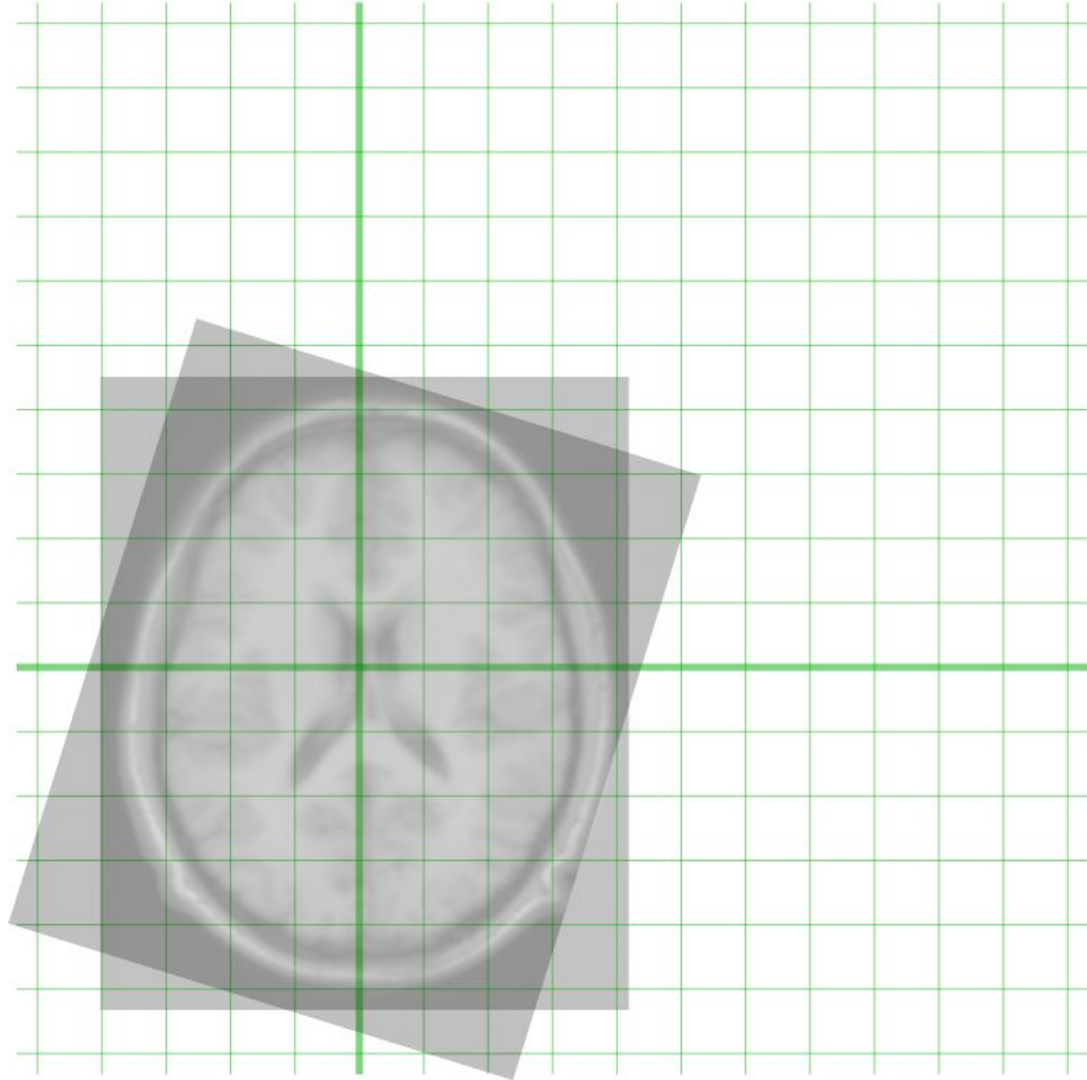
463 ROI



1015 ROI



4. Couple of words on Image Registration



Voxel-wise registration

Register on template



Original Brain

+

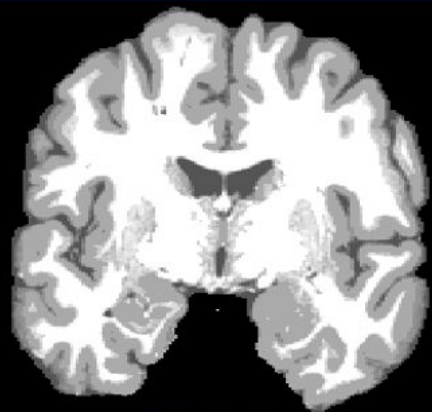


MNI152 Template

=



Registered Brain



Atlas Brain



Subject Brain

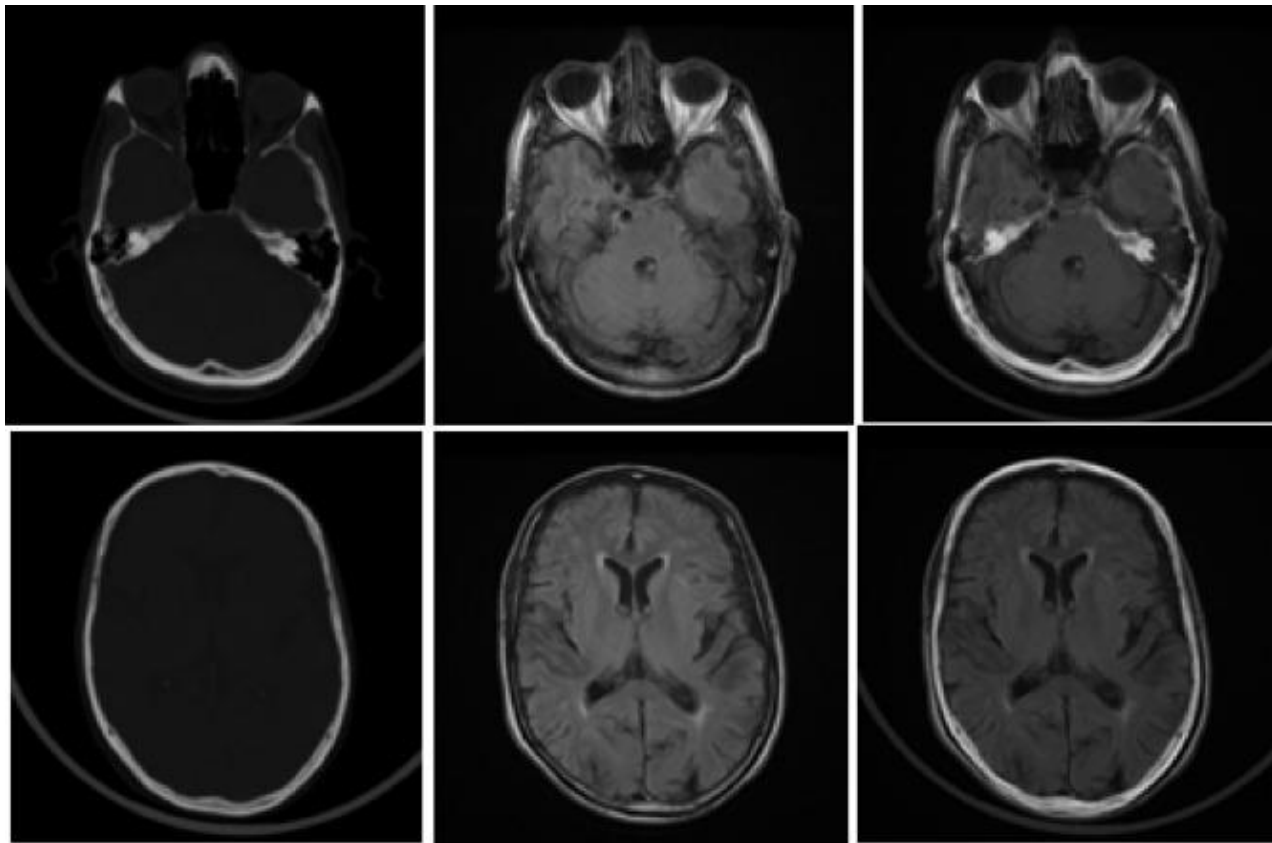


Warped atlas by surface-
constrained mapping

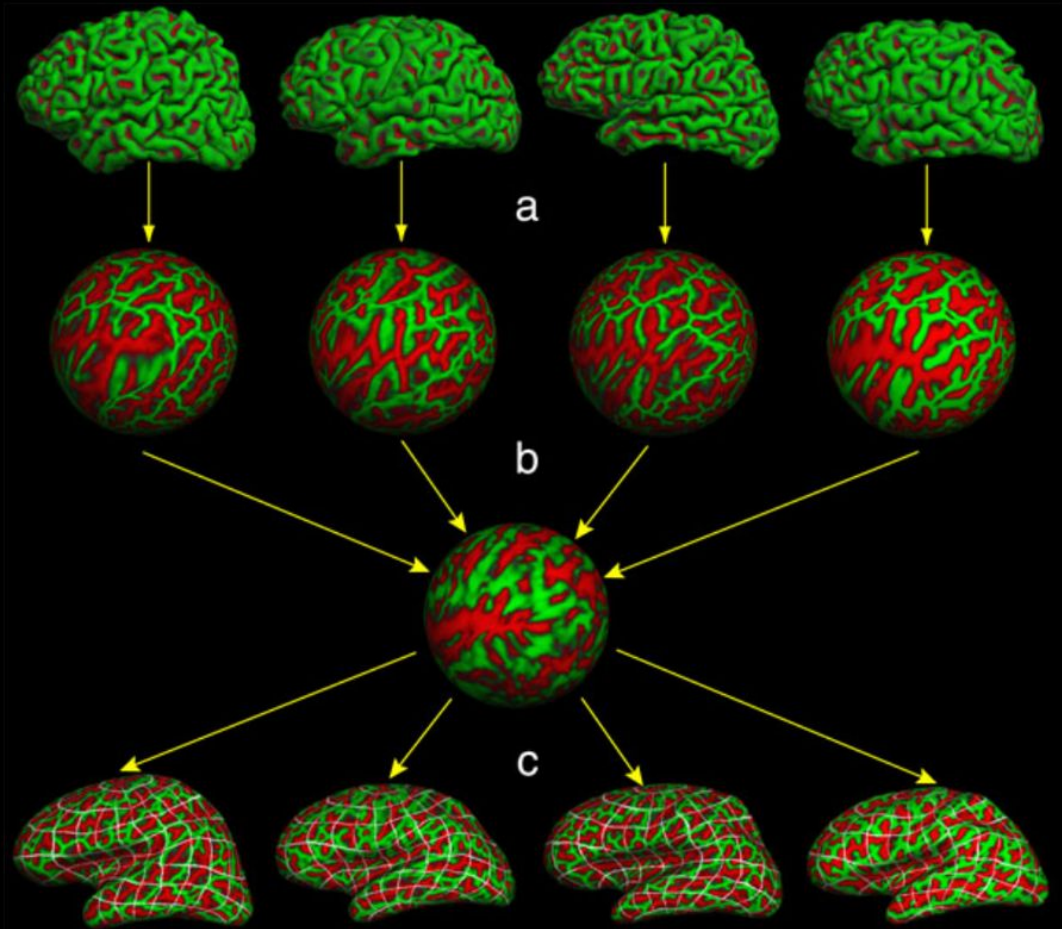


Warped atlas after
intensity registration

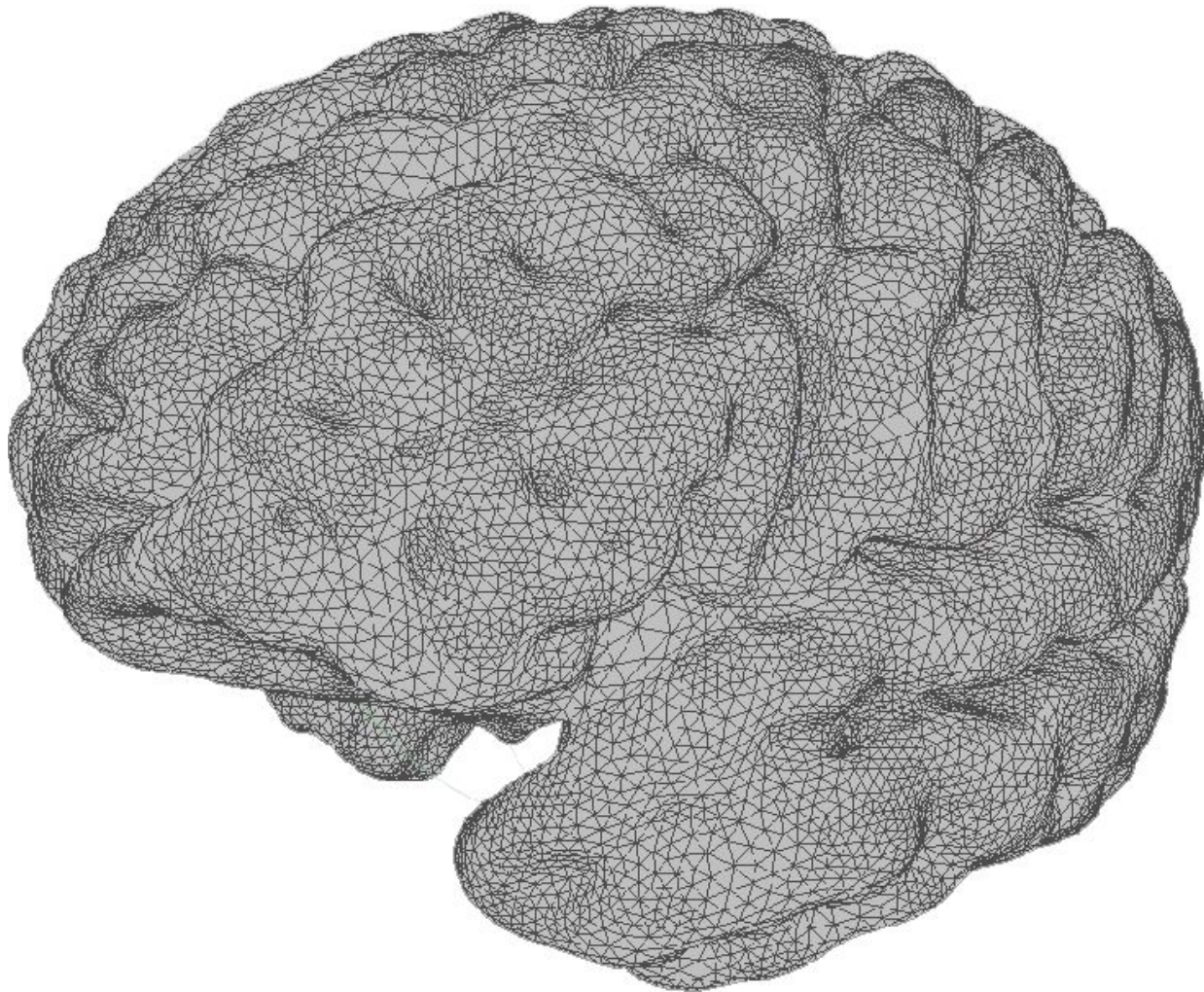
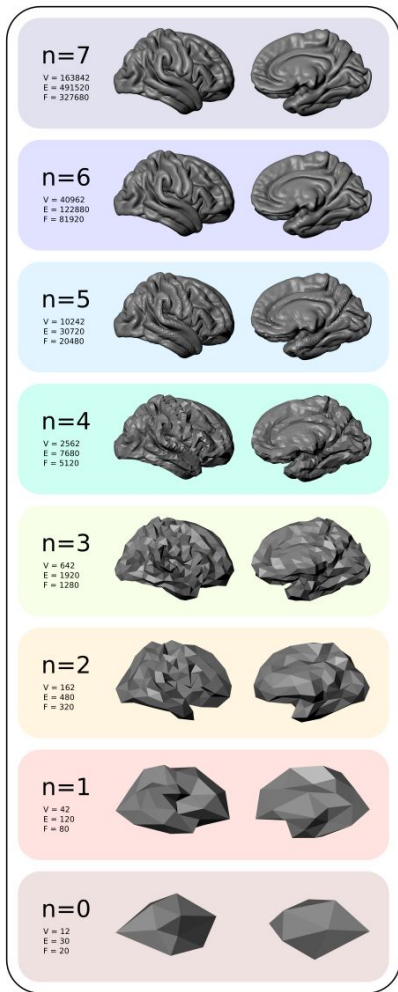
Register on modality



Surface-based registration

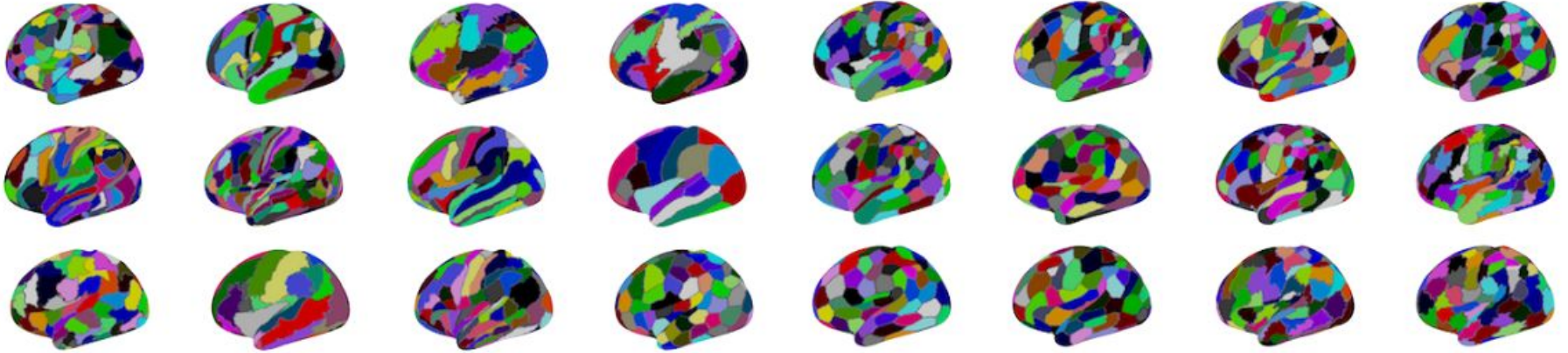


[Freesurfer](#)



4. Data driven brain parcellations

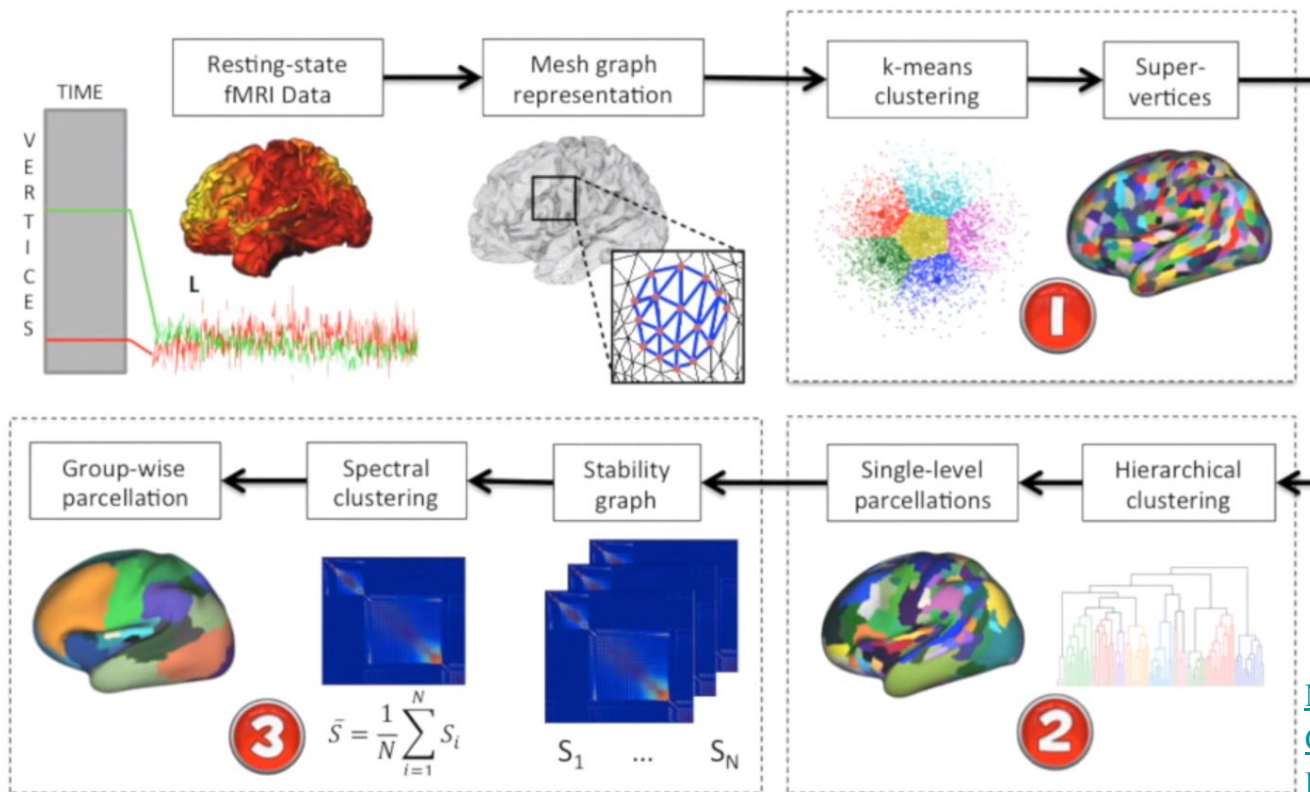
Dozens (or even hundreds) of them



[Human Brain Mapping: A Systematic Comparison of Parcellation Methods for the Human Cerebral Cortex](#)

1. **2015** Multi-Level Parcellation of the Cerebral Cortex Using Resting-State fMRI, Salim Arslan, Daniel Rueckert https://www.doc.ic.ac.uk/~sa1013/pub/2015_S_Arslan_MICCAI.pdf
 - a. 100 HCP subjects. Initial parcellation using k-means, distance is a combination of geodesic distance and time series correlation (from **fMRI**) -> hierarchical (agglomerative) clustering of these supervertices into larger ones -> Construct meta-graph edges - #times vertices co-occur in the same community -> Cluster this graph using n-cut
2. **2015** A Continuous Flow-Maximisation Approach to Connectivity-driven Cortical Parcellation, Sarah Parisot, Martin Rajchl, Jonathan Passerat-Palmbach, Daniel Rueckert, 2015
 - a. Start with random parcellation (spatially constrained) -> Update parcel centers seeking for a point with maximum correlation with all other nodes in a parcel (correlation from **fMRI**) -> Attach each node to a parcel with the highest correlation (s.t. Spatially smoothness constraints) -> Repeat until convergence.
 - b. 25 HCP Subjects
3. **2008** Normalized Cut Group Clustering of Resting-State fMRI Data, Martijn van den Heuvel, Rene Mandl, Hilleke Hulshoff Pol, 2008, <https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0002001#s2>
 - a. 2-level procedure. 1 construct subject level parcellation from **fMRI** (correlation, 0.4 cutoff) - 20 clusters (authors call them resting-state networks RSNs). Individual graph consist of 8500-9500 nodes. Use ncut clustering, no spatial constraints. Construct group graph edge +1 between 2 nodes if they were in the same cluster (for a subject). Finally group graph was clustered using ncut. 26 fMRI subjects.
4. **2012** A whole brain fMRI atlas generated via spatially constrained spectral clustering, R. Cameron Craddock, G. Andrew James, Paul E. Holtzheimer, Xiaoping P. Hu, and Helen S. Mayberg, 2012
 - a. Build a network from **fMRI**, such that every node is a voxel and an edge between two nodes exist only if they are in 3D neighborhood (for every voxel there are 26 neighborhood voxels), the weight on an edge is a correlation. Cluster it using ncut, group atlas generated either by averaging subject networks and cluster an averaged one, or by using the same technique as in [3] (Martijn van den Heuvel, 2008)
5. **2014** OPTIMIZING BRAIN CONNECTIVITY NETWORKS FOR DISEASE CLASSIFICATION USING EPIC Gautam Prasad, Shantanu H. Joshi, and Paul M. Thompson <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4232940/>
 - a. Start with Desikan atlas proposed algorithm combine separate regions into bigger ones, recompute connectivity matrix and run a classification algorithm on it (using PCA as prep step). In such a way authors find “optimal” in terms of classification (AD vs NC) brain parcellation. The search of best combination of regions is done in a probabilistic manner (simulated annealing/random search).

Methodology

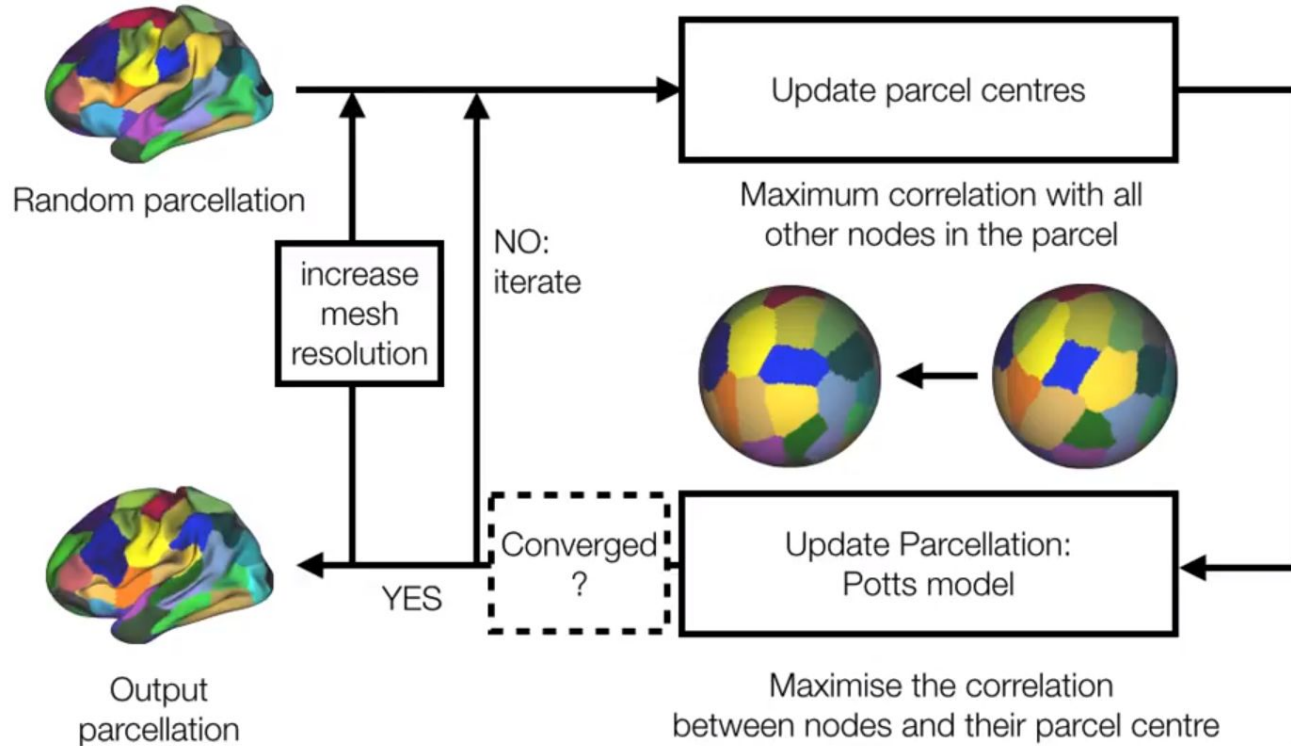


[Short presentation video](#)

[Multi-Level Parcellation of the Cerebral Cortex Using Resting-State fMRI](#)



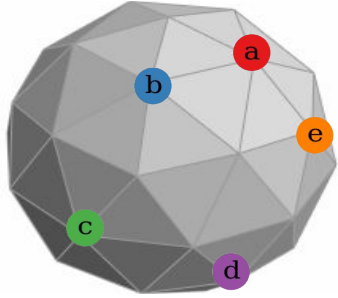
Method Overview



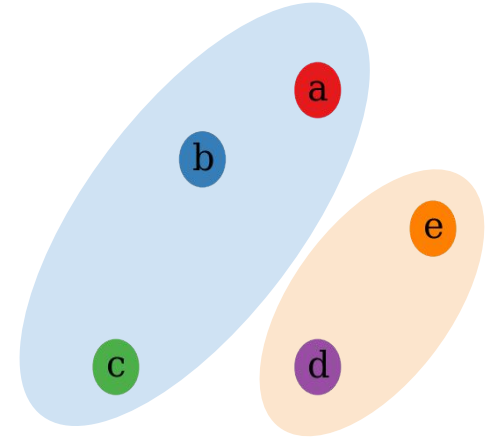
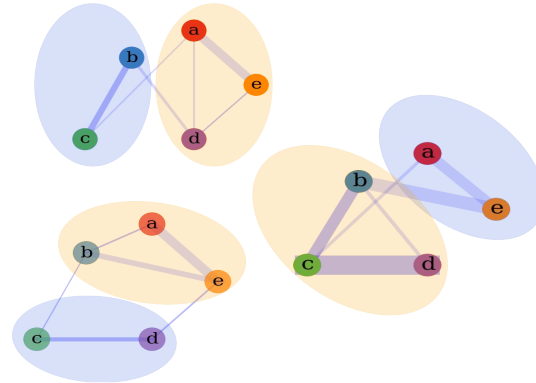
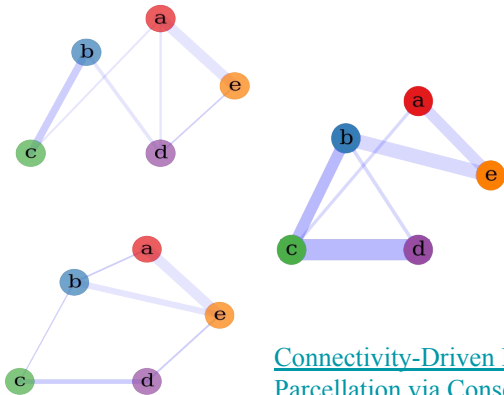
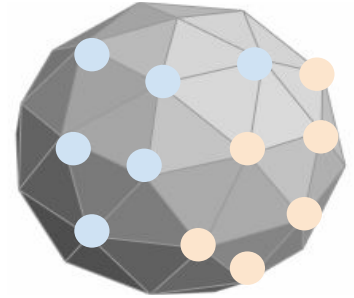
[Short presentation video](#)

[A Continuous Flow-Maximisation Approach to Connectivity-driven Cortical Parcellation](#)

Connectivity-Driven Brain Parcellation via Consensus Clustering

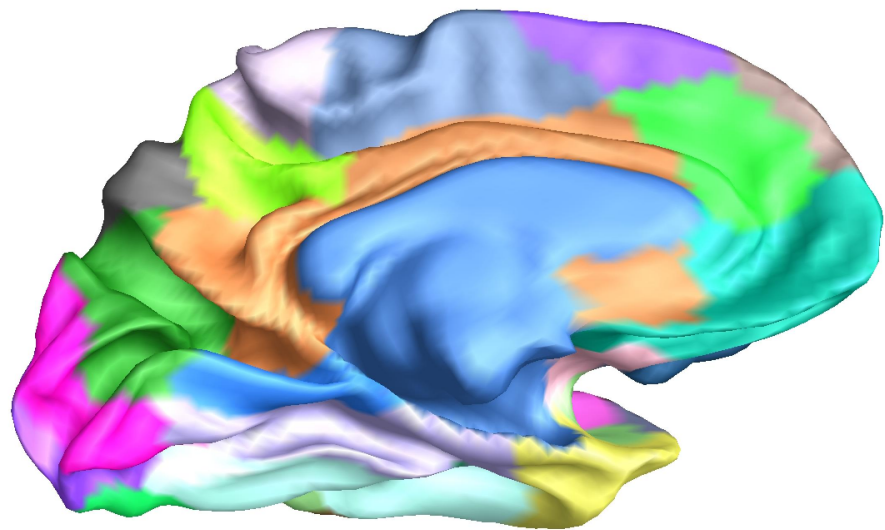
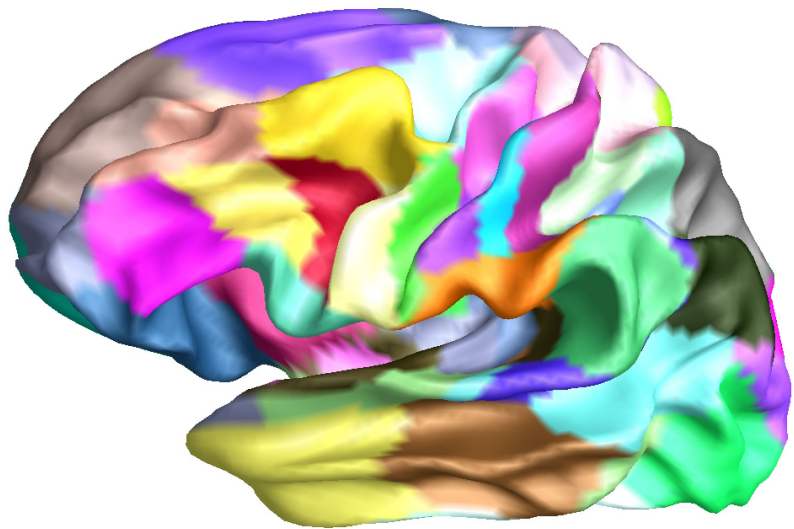


1. Surface mesh
2. Individual continuous connectomes
3. Partitions of the individual connectomes
4. Consensus vertex clustering
5. Mapping clusters on brain surface



Features

- Obtained parcellation is highly symmetrical (left vs right hemisphere)
- Has substantial intersection with classical gyral based parcellations
- Spatially continuous without specific spatial constraints
- Arbitrar subject to clustering approach and averaging approach
- Could be used for subject or group analysis



Random picture sources

1. http://www.clipartpanda.com/clipart_images/black-and-white-human-brain-3-29489506
2. <https://braintumor.org/brain-tumor-information/signs-and-symptoms/brain-illustration/>
3. <https://www.britannica.com/topic/phrenology>
4. <https://www.humanbrainfacts.org/basic-structure-and-function-of-human-brain.php>
5. <http://cs.wellesley.edu/~cs112/courseMaterials/assignments/assign5/assign5.html>
6. <https://miykael.github.io/nipype-beginner-s-guide/neuroimaging.html>
7. http://www.clinica.run/doc/Pipelines/T1_FreeSurfer/
8. <https://fcp-indi.github.io/docs/user/anat.html>
9. <http://brainsuite.org/processing/svreg/details/>
10. https://en.wikipedia.org/wiki/Image_registration
11. <https://www.spiedigitallibrary.org/conference-proceedings-of-spie/10133/1/Multi-atlas-based-CT-synthesis-from-conventional-MRI-with-patch/10.1117/12.2254571.short?SSO=1>
12. <https://www.sciencedirect.com/science/article/pii/S2319417017300653>
13. <https://www.semanticscholar.org/paper/Shape-analysis-of-the-human-brain.-Nitzken-Joseph/c47a238ce4a943a30da38f1047c01580a47fab7d/figure/63>
14. <https://www.semanticscholar.org/paper/Shape-analysis-of-the-human-brain.-Nitzken-Joseph/c47a238ce4a943a30da38f1047c01580a47fab7d>
15. <https://brainder.org/2016/05/31/downsampling-decimating-a-brain-surface/>

Conclusion

- Problems:
 - Huge amount of different parcellation approaches
 - No obvious way to choose amongst them
- Solution:
 - Use common sense
 - Use anatomical parcellations, they are good enough in most cases.
 - Do not use parcellation at all.
 - In case of structural connectomes use Connectivity-driven parcellation!

Szemerédi regularity lemma

From Wikipedia, the free encyclopedia

In [mathematics](#), the **Szemerédi regularity lemma** states that every large enough [graph](#) can be divided into subsets of about the same size so that the edges between different subsets behave almost randomly. [Szemerédi \(1975\)](#) introduced a weaker version of this lemma, restricted to bipartite graphs, in order to prove [Szemerédi's theorem](#),^[1] and in [\(Szemerédi 1978\)](#) he proved the full lemma.^[2] Extensions of the regularity method to [hypergraphs](#) were obtained by [Rödl](#) and his collaborators^{[3][4][5]} and [Gowers](#).^{[6][7]}

