



Brain Dynamics on the Connectome Summer School 2021

Analysis of Brain Networks

Brain Dynamics on the Connectome Summer School 2021

12/10/2021

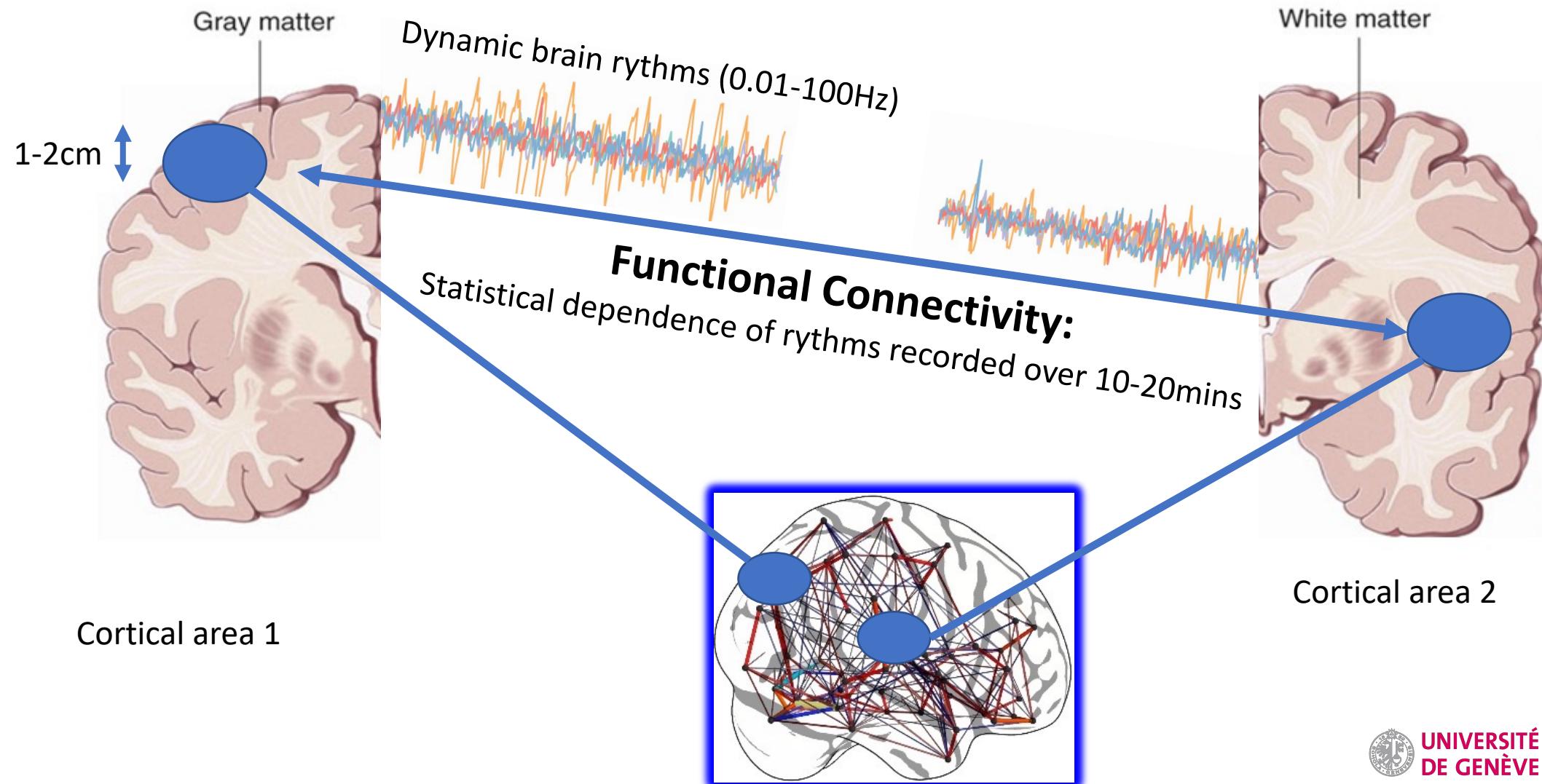
Jonathan Wirsich

University of Geneva, University Hospitals of Geneva and the Faculty of
Medicine

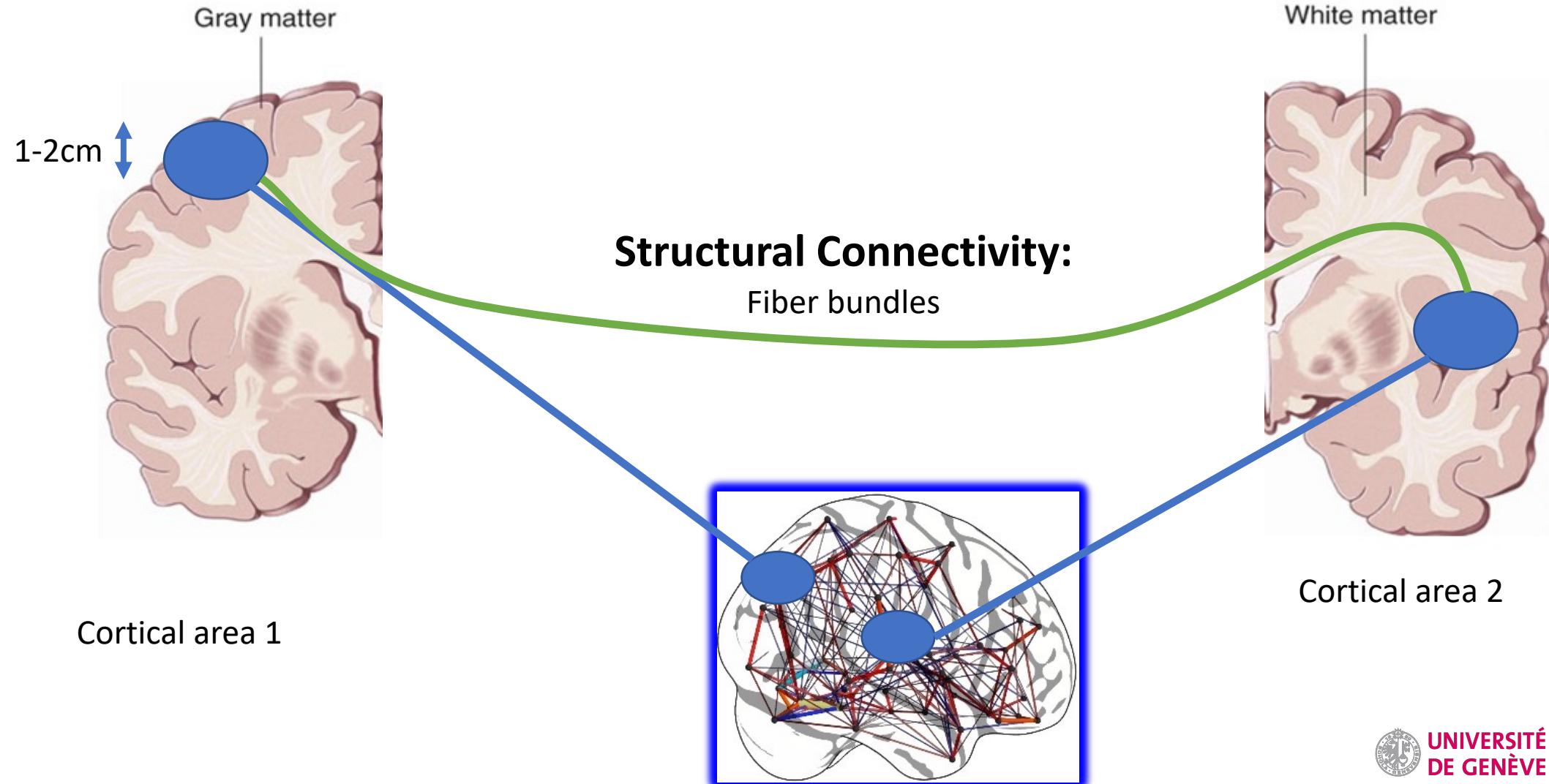
Outline

- EEG and fMRI connectivity
- Graph analysis
- Combination of EEG and fMRI

Accessing how brain regions-communicate in a network at multiple scales



Accessing how brain regions-communicate in a network at multiple scales



Non-invasive means to measure the human connectome

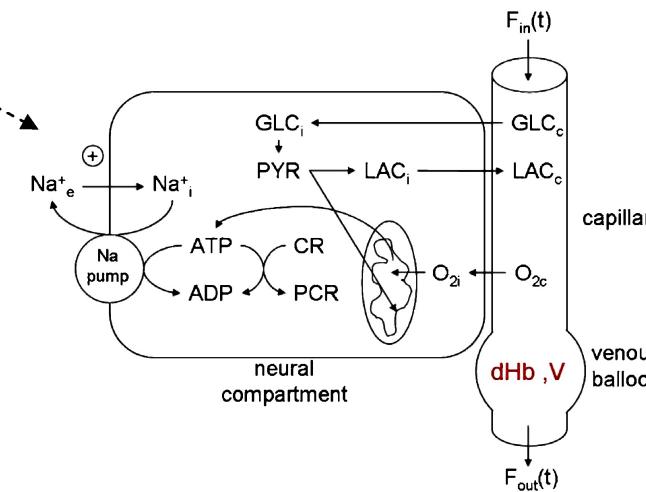
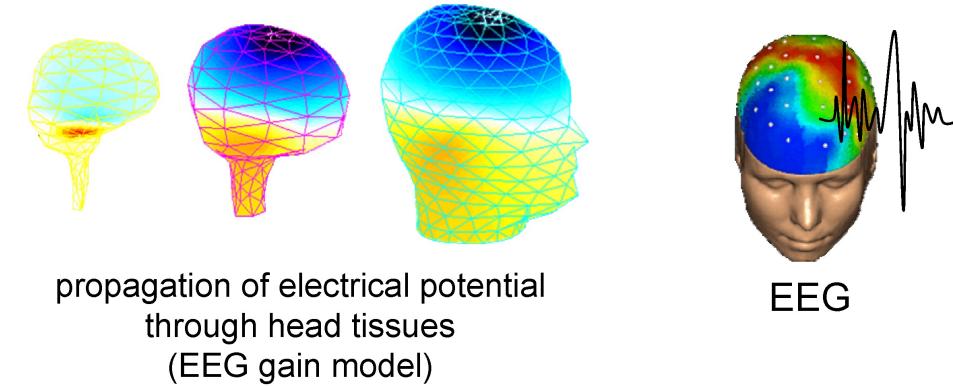
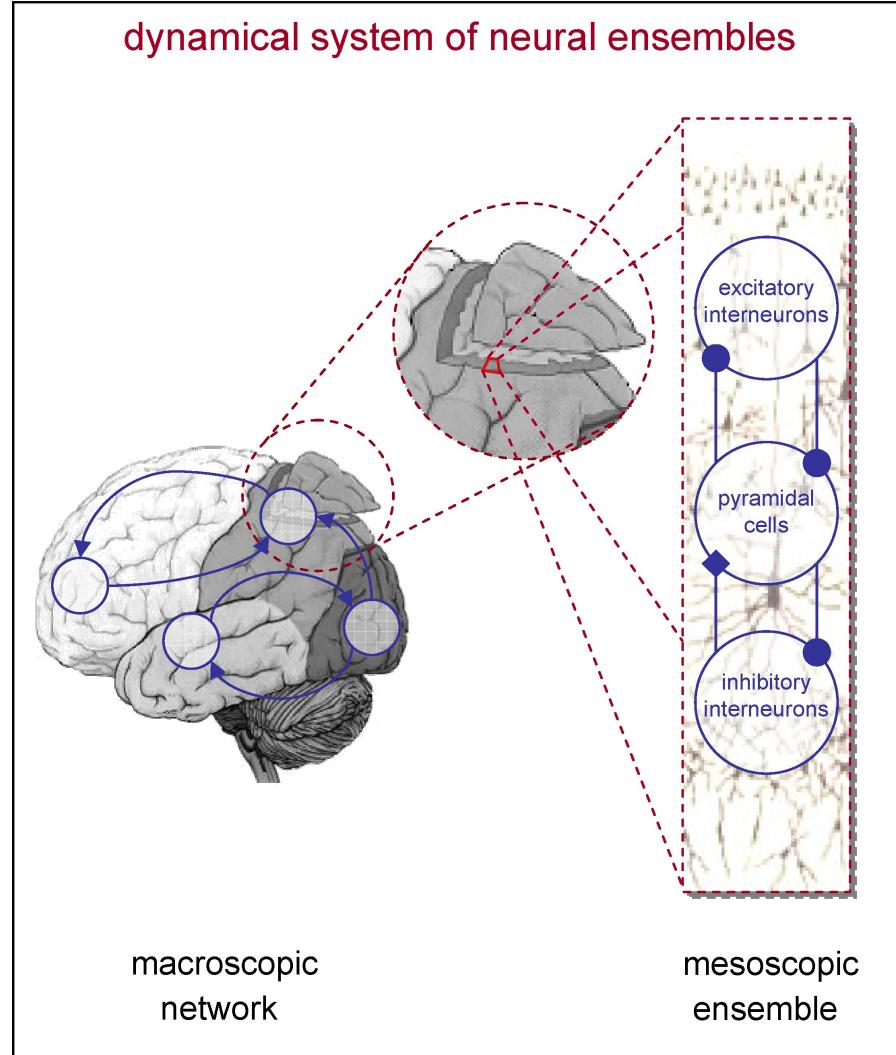
EEG



MRI



EEG and fMRI signal



metabolic and hemodynamic cascade
(extended fMRI balloon model)

[Rosa 2010, J Integr Neurosci]

Brain Connectivity

- Brain = City with a lot of connected streets
- Brain activity: Number of cars in one region



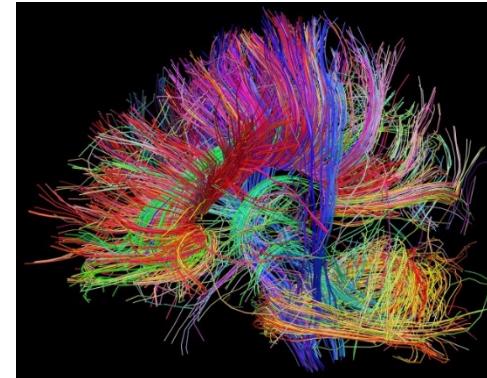
Building Network Graphs



Measures of connectivity



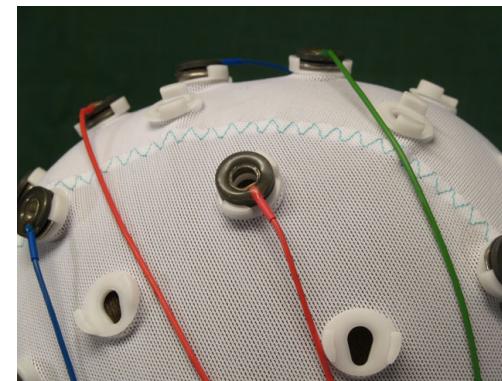
Number of lanes =
dMRI based number
of fiber tracts



Consumption of gas
measured at a gas station =
functional MRI based
BOLD signal



Smog caused by cars =
EEG signal



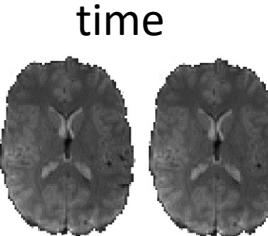


Constructing the multimodal connectome

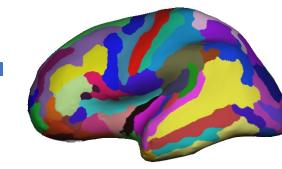
EEG
1ms
>1cm



Functional
MRI
1-2s
3mm

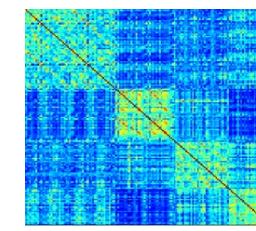
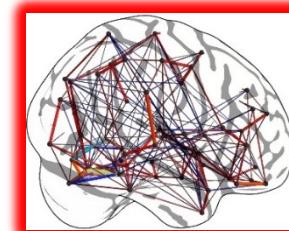
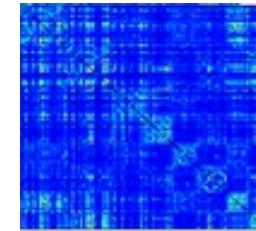
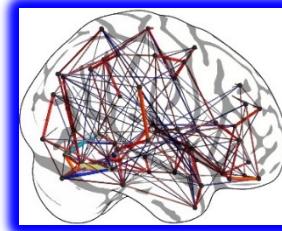


Tractography
diffusion
MRI
3mm

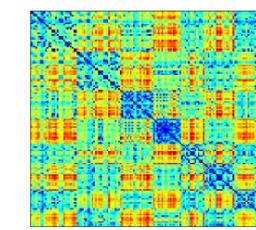
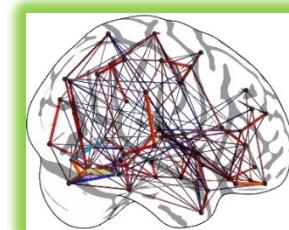


Atlas

Connectome Connectivity matrix



regions



Connectome

Connectivity matrix

EEG neuronal functional connectivity

fMRI metabolic functional connectivity

dMRI structural connectivity

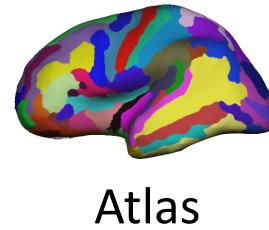
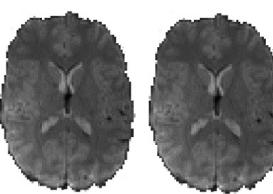
brain dynamics occur on the millisecond timescale → concurrent recordings are needed for the full picture
Trimodal See e.g. [Wirsich 2017, NeuroImage], **structure vs. function later presentations**



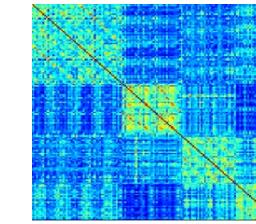
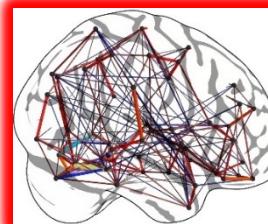
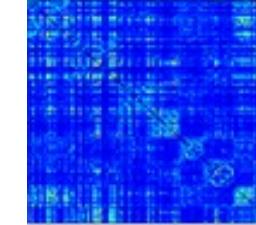
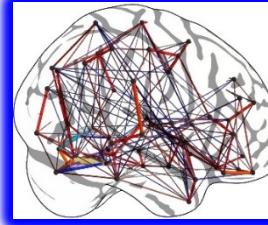
Constructing the multimodal connectome

EEG
1ms
>1cm

Functional
MRI
1-2s
3mm



Connectome Connectivity matrix



regions

Design of network analysis:

- What task?
- Which preprocessing steps?
- What atlas to use?
- What kind of connectivity?
- What properties of the graph?



UNIVERSITÉ
DE GENÈVE

Task Paradigm

resting state



vs.



task

Source: Wikimedia

Task-based see next presentation: Functional Connectivity (FC) and Dynamical FC

Resting state

- **Connections most commonly derived from resting state**



- **Session length most commonly 5-20min**
 - Long enough for multiple cycles of infraslow (<0.1Hz) frequencies (fMRI), faster cycle possible with EEG
 - Short enough to minimize mental state change
 - Shorter term time-varying dynamics (e.g. sliding window)

Preprocessing fMRI

- **Preprocessing: same considerations as any fMRI connectivity study:**
 - Motion correction
 - Slice time correction
 - Physiological nuisance measures (ECG, breathing, etc.)?
 - Compartment signal regression (Gray Matter aka. Global mean, White Matter, Cerebrospinal-Fluid)
 - EEG covered in previous lecture

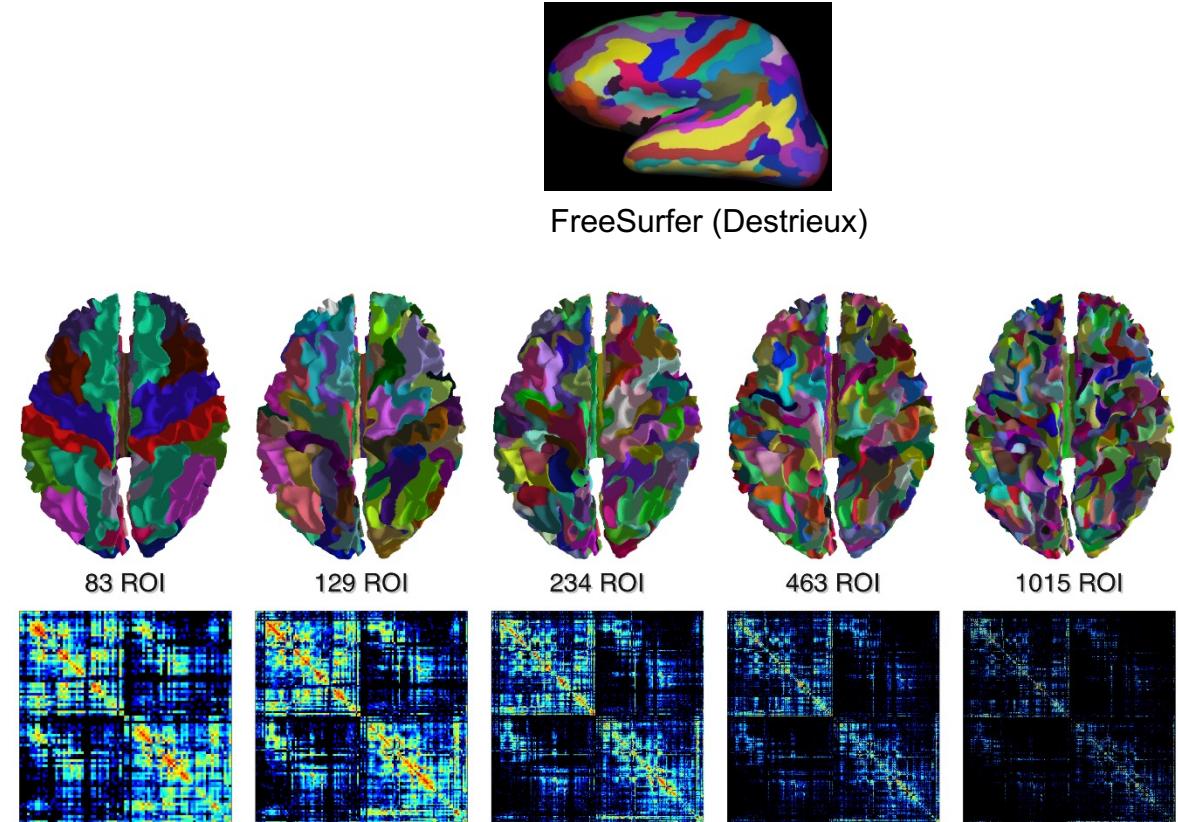
Anatomical atlases

Nodes:

- Internally coherent / homogeneous (connectivity)
- Externally independent

Anatomical atlases

- Talairach & Tournoux
- Automated Anatomical Labeling (AAL) template
- Eickhoff-Zilles (Cytoarchitectonic)
- FreeSurfer (Individual surface-based)
- Lausanne (subdivision of Freesurfer)

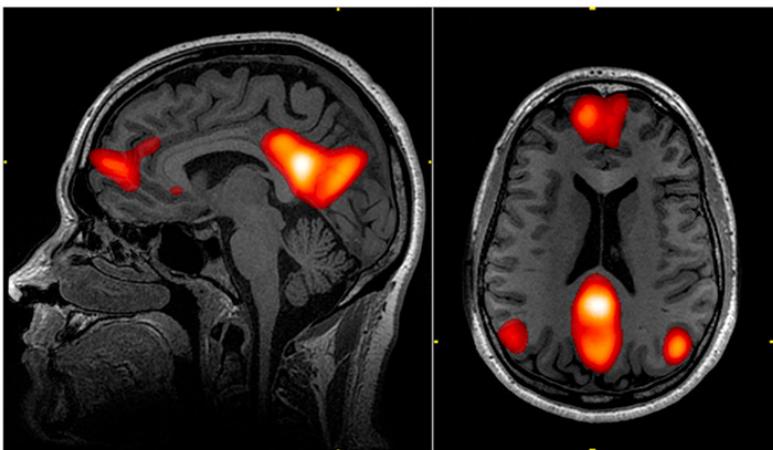


- ☺ Comparability (across subjects and modalities)
- ☹ Highly variable node size
(can be mediated by subdividing such as Lausanne atlas). Not functionally coherent.

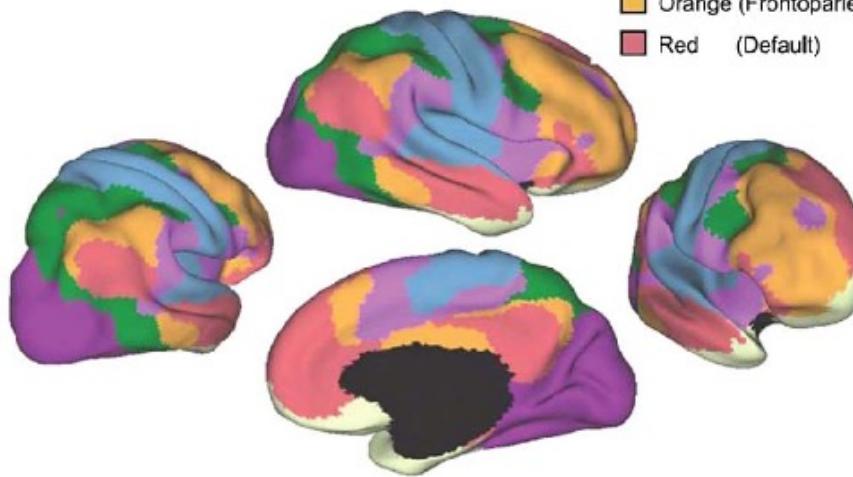
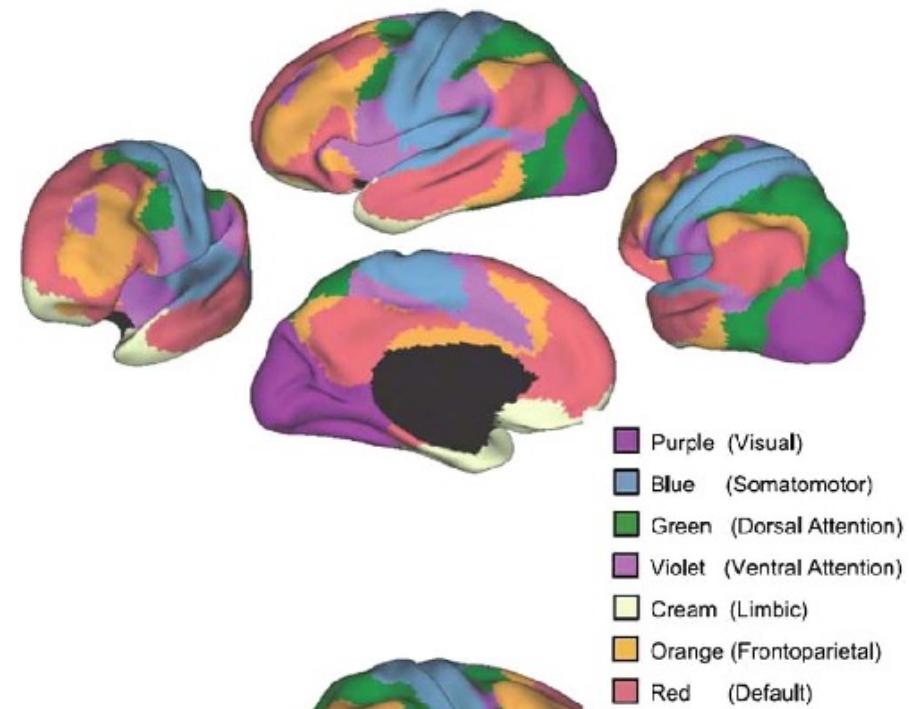
Intrinsic connectivity networks (M/EEG and fMRI)



Resting-state = Default mode network activated



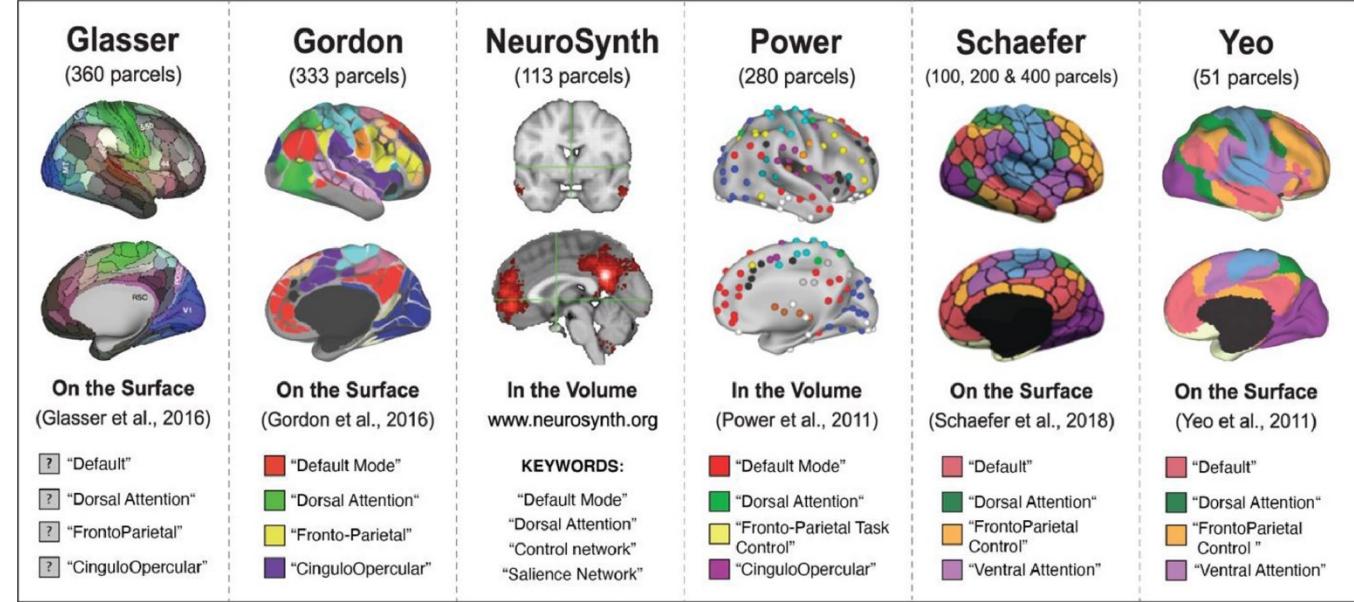
[Graner 2013, Front in NeuroSci]



[Yeo 2011, J Neurophysiol]

Functional atlases

- Functional atlases
 - Comparability across subjects.
 - Functionally coherent
(but suboptimal for individuals)



Functional subject-specific parcellations

- ICA (Seed-based)
- Connectivity homogeneity: Craddock
- Functionally coherent
- Time-intensive

[Bryce 2021, NeuroImage]

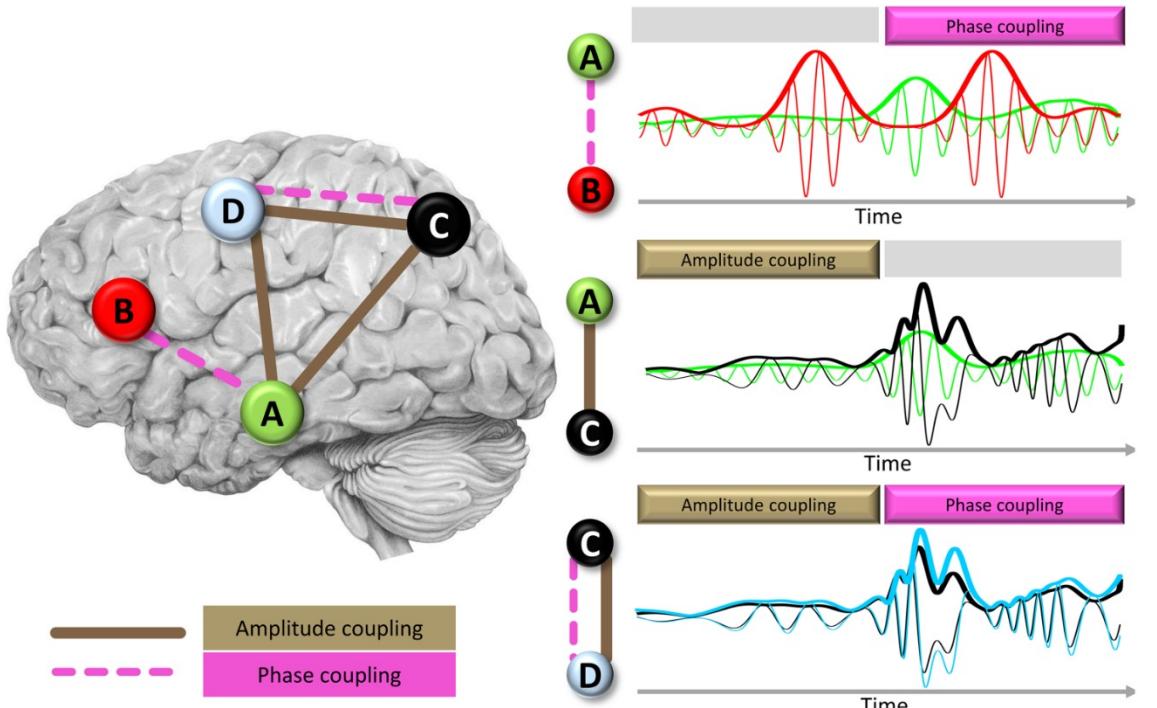
List of atlases: <https://en.wikibooks.org/wiki/SPM/Atlases>



UNIVERSITÉ
DE GENÈVE

Edges in EEG and fMRI

- Based on magnitude of temporal covariation
 - Pearson's cross-correlations (by far most common)
 - Partial correlations
 - Mutual information
 - → symmetric adjacency matrices (undirected graphs)
- Directionality problematic in fMRI (but measures of effective connectivity possible)
- EEG: coherence, amplitude correlations, etc. directionality less problematic but source leakage



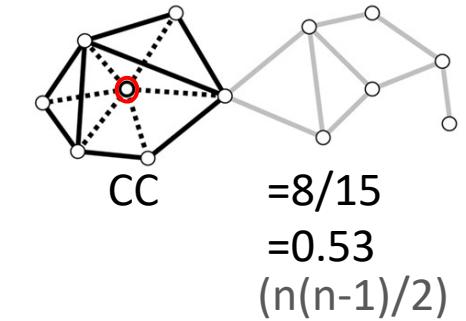
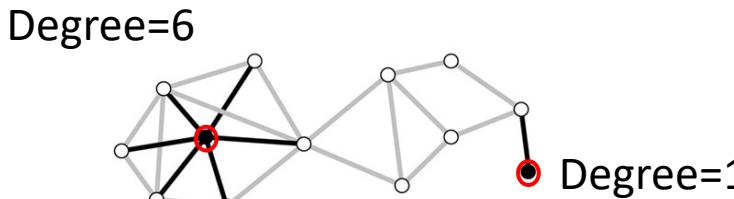
[Sadaghiani and Wirsich 2020, Network NeuroSci]



Nodal Measures

Degree

- Number of edges connected to a node

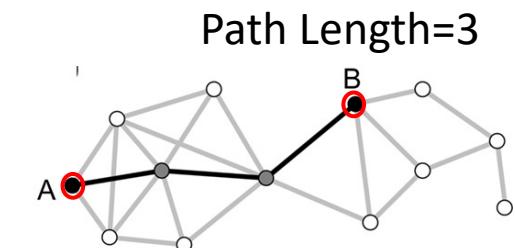


Nodal Clustering Coefficient (→ basis for measure of global segregation)

- Fraction of all possible edges realized among a node's neighbors
= Fraction of all possible triangles around a node

Shortest Path Length (→ basis for measure of global integration)

- Number of edges on shortest geodesic path between two nodes



Global Measures

Small-worldness

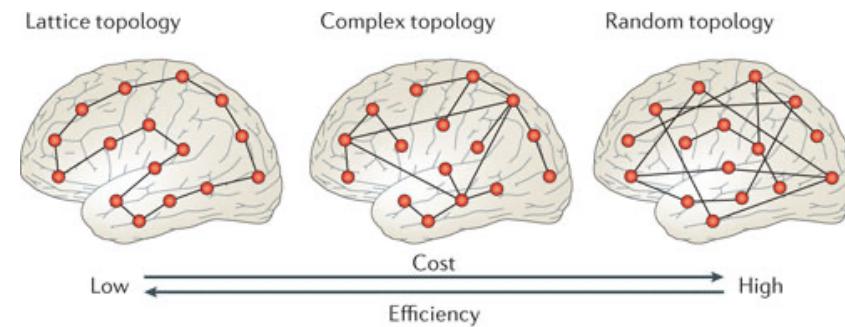
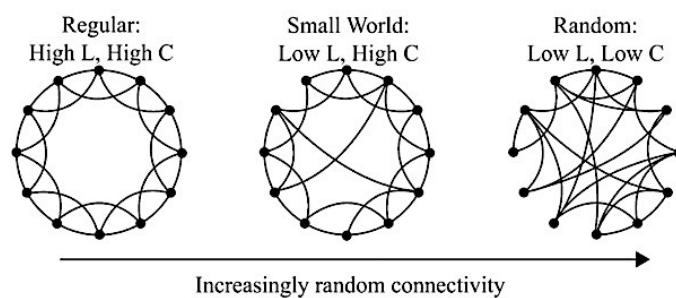
- Optimal balance between functional segregation and integration:

- C (high = segregated):
- L (low = integrated):

Clustering Coefficient_{real} / Clustering Coefficient_{random}

Characteristic Path Length_{real} / Characteristic Path Length_{random}

[Watts and Strogatz 1998, Nature]

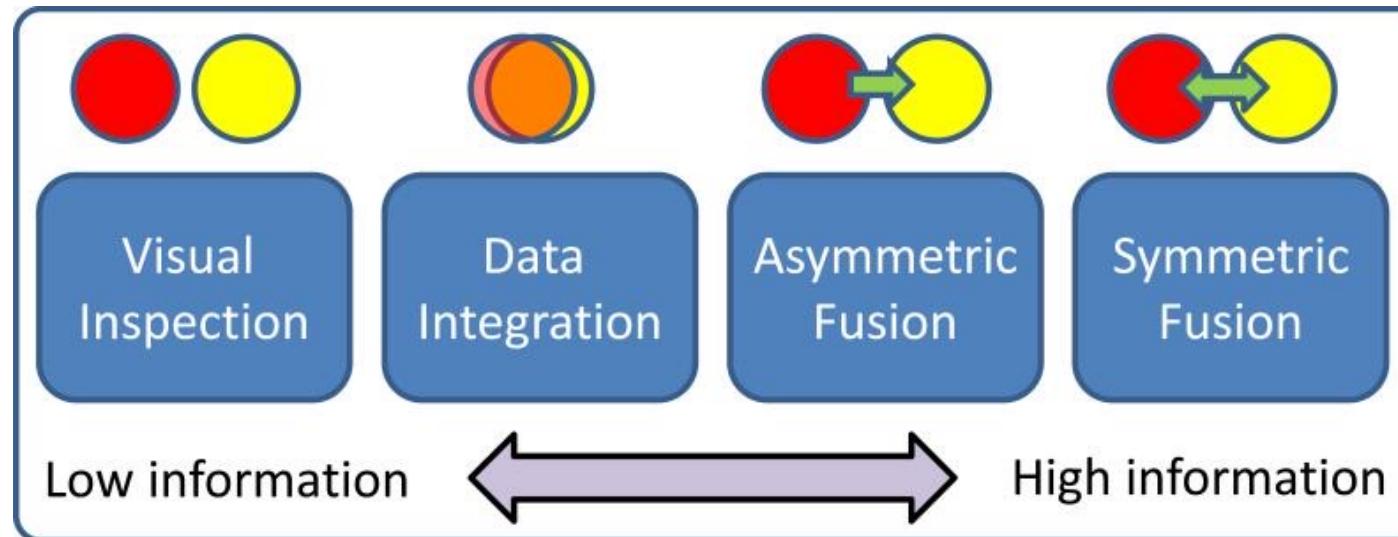


☺Functionally specialized (segregated) modules **AND** intermodular (integrating) edges

- A very easy global parameter is the spatial correlation between sessions subjects or modalities



Fusion and Integration of EEG and fMRI



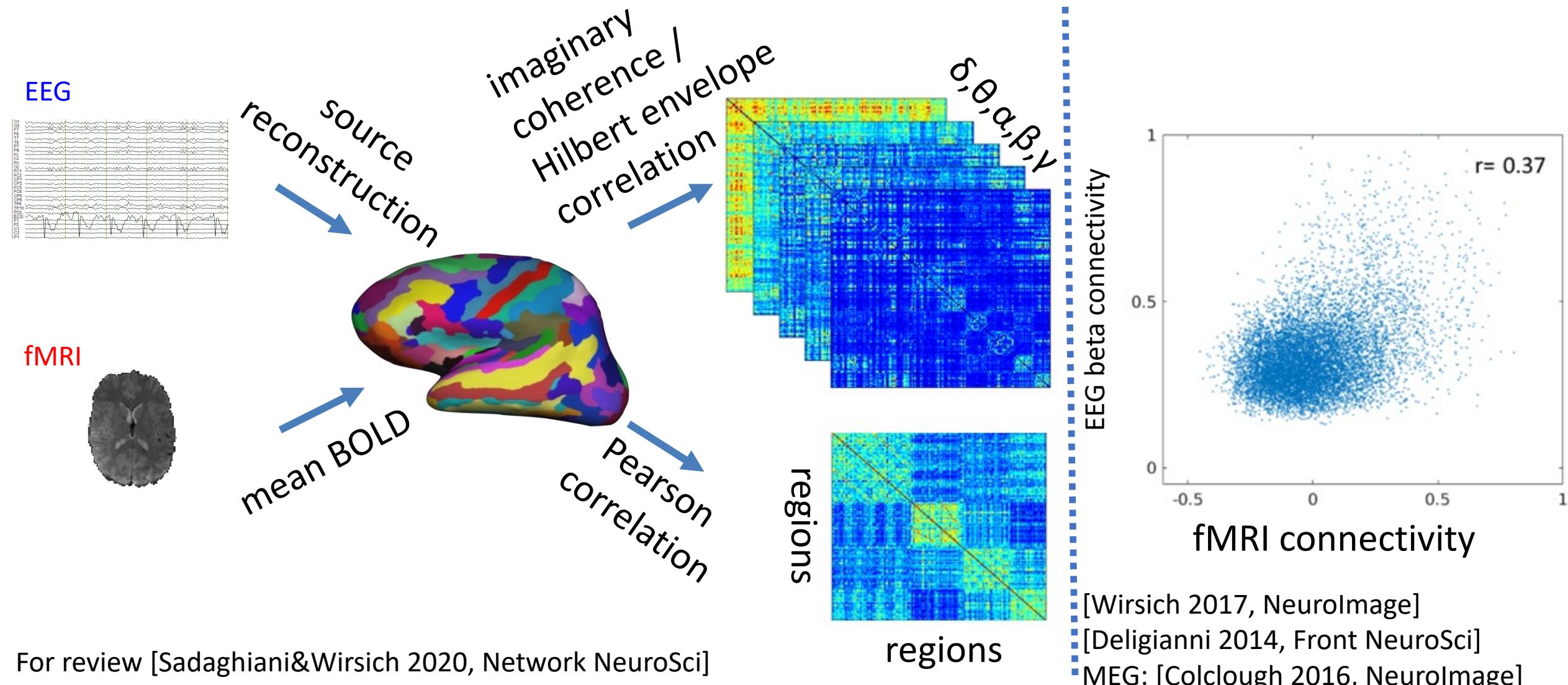
Source: [Calhoun 2016, BPCN Neuro]



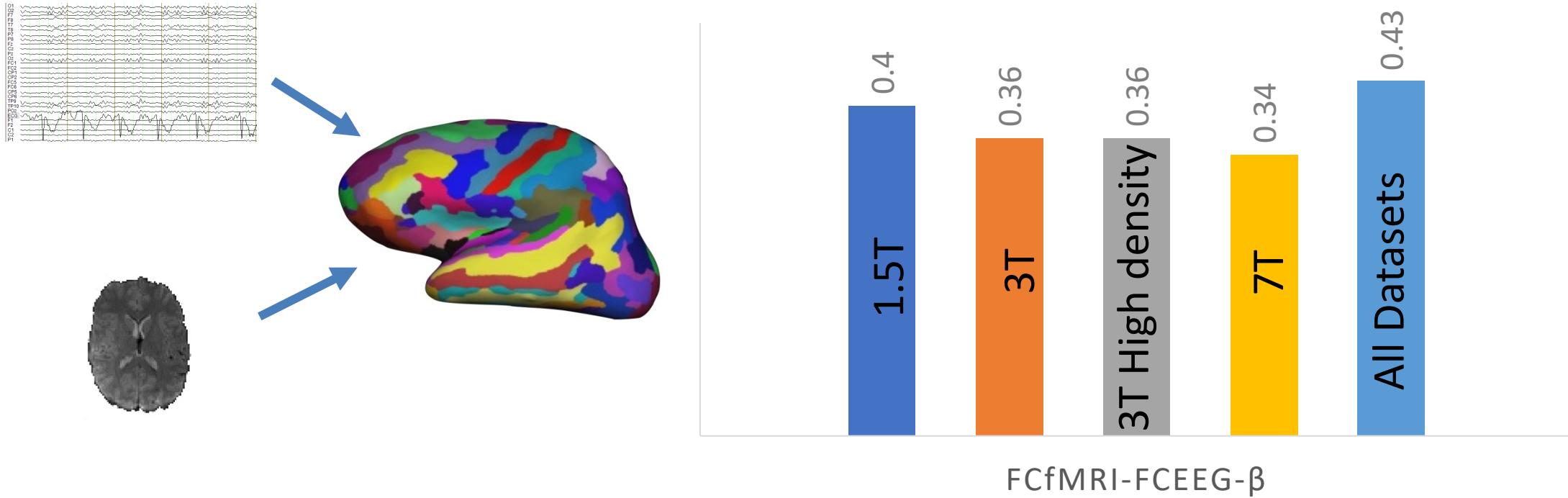
- Low information: Comparison of graph properties (e.g. clustering, small-worldness)
- Fusion: high temporal resolution of EEG and high spatial resolution of fMRI are used: see e.g. [Wirsich 2014, NeuroImage]

Source: xkcd.com

Spatial integration of EEG and fMRI connectomes

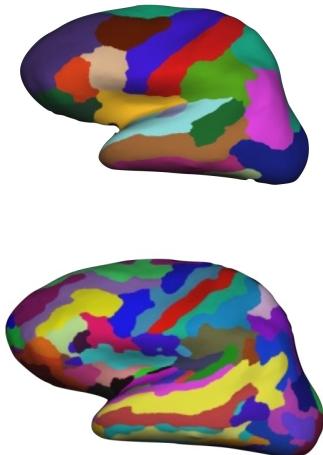


Spatial correlation of group average EEG and fMRI connectomes



Correlation around $r=0.3$, gamma performs well only on 1.5T, averaging all datasets results in best results

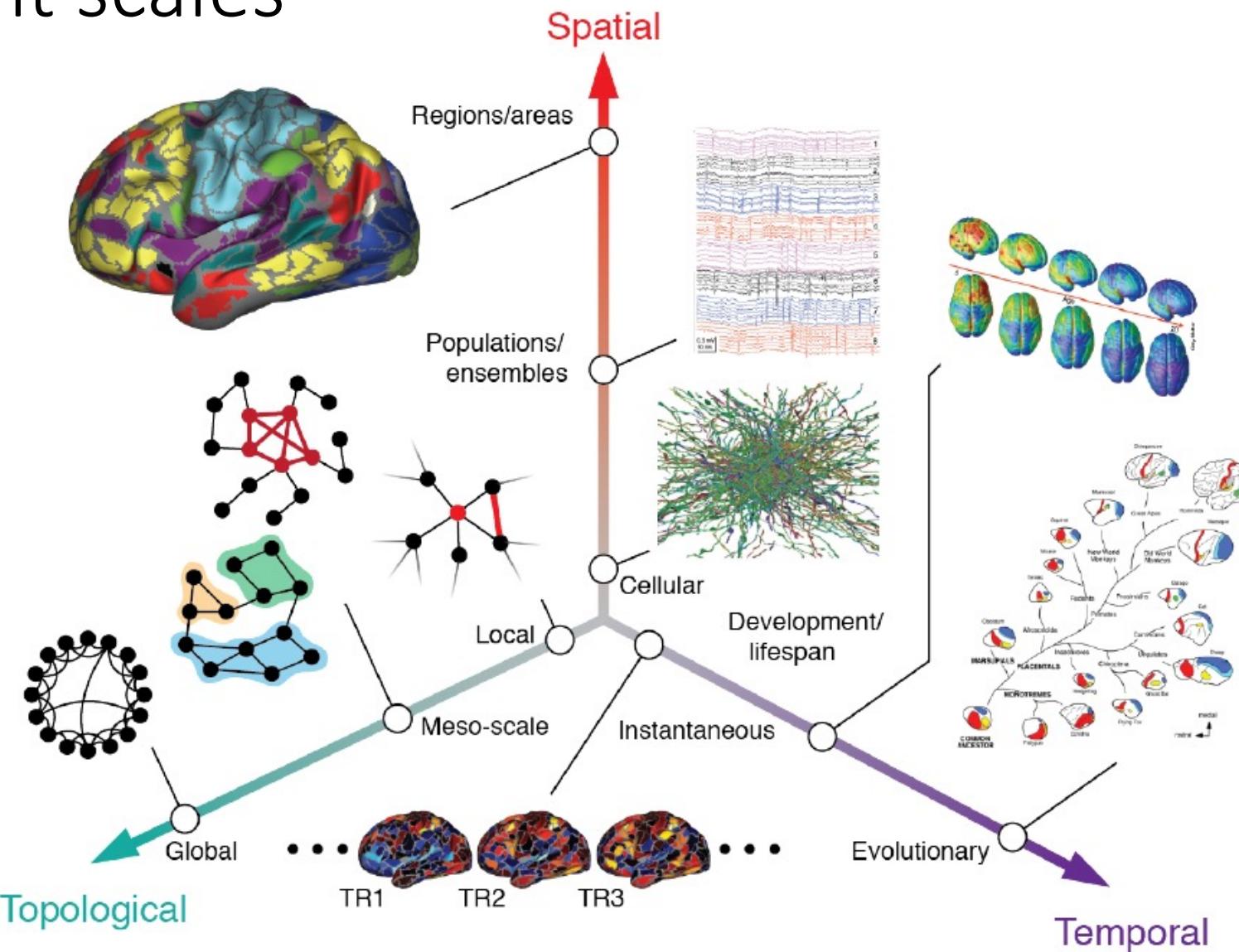
Increased EEG-fMRI correlation in smaller atlas



Imaginary coherence Beta	64Ch-1.5T	64Ch-3T	256Ch-3T	64Ch-7T	All combined
Desikan n=68	0.40	0.36	0.36	0.34	0.43
Destrieux n=148	0.29	0.33	0.26	0.31	0.36

- Quality of Destrieux can be boosted by combining more datasets
→ Different setups seem to do better for some connections

Understanding the brain = know how to link different scales



[Betzel and Bassett 2017, NeuroImage]



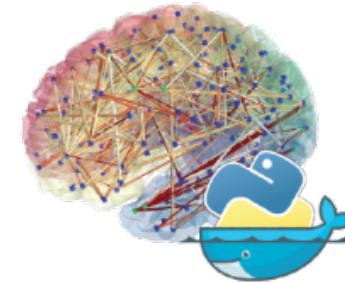
UNIVERSITÉ
DE GENÈVE

Resources

Preprocessing

Connectomemapper

<https://connectome-mapper-3.readthedocs.io/en/latest/>



Graph analysis

MATLAB-based: **Brain Connectivity Toolbox** (Rubinov and Sporns, 2010)

- <https://sites.google.com/site/bctnet/>
 - Python-based: **NetworkX** (Hagberg et al., 2008)
- <https://networkx.github.io>

Reading

- Rubinov M, Sporns O. Complex network measures of brain connectivity: Uses and interpretations. *NeuroImage*. 2010 Sep;52(3):1059–69.
- Bullmore ET, Bassett DS. Brain Graphs: Graphical Models of the Human Brain Connectome. *Annu Rev Clin Psychol*. 2010 Apr;7(1):113–40.
- Sadaghiani S, Wirsich J. Intrinsic connectome organization across temporal scales: New insights from cross-modal approaches. *Network Neuroscience* 2020 4:1, 1-29
- Daducci et al. The Connectome Mapper: An Open-Source Processing Pipeline to Map Connectomes with MRI. *PlosOne* 2012