



Aalto University  
School of Science

# Brain networks Current state and future challenges

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# PART 1

# Brain networks

# ABC

# The Brain according to wikipedia

...The brain is the most **complex** organ in a vertebrate's body...

# The Brain according to wikipedia

...In a typical human the **cerebral cortex** (the largest part) is estimated to contain 15–33 billion ( **$10^9$ !!**) neurons each **connected** by synapses to **several thousand** other neurons...

# Why do we want to study brain networks?

- **The brain is a network with**  
~ $10^{10}$  neurons and ~ $10^4$  connections per neuron
- As for **genomics** in the 20<sup>th</sup> century, many authors are now praising the **connectomics** as the current revolution in neuroscience
- Multi-million projects like the **Human Connectome Project, the BRAIN initiative**
- **Charting the connectome** presents challenges

# What is a connectome?

# The connectome

The connectome is the complete description of the structural connectivity (the physical wiring) of an organism's nervous system.

*Olaf Sporns (2010), Scholarpedia, 5(2):5584.*

# Structure and Function



# Many types of networks

- **Physical networks**
  - *Power grid network*
  - *Physical layer of the internet*
  - *Transportation networks (roads, rails)*
- **Non-physical networks**
  - *Social networks (Facebook, Twitter, etc.)*
  - *Stock Market*
  - *IP layer of the internet*

# Brain networks

- **Structural connectivity**  
(estimating actual connections, the connectome)
- **Functional connectivity**  
(based on temporal “co-variance”)

Craddock, et al. (2013). Imaging human connectomes at the macroscale. *Nature Methods*, 10(6), 524–539. (\*)





**Neural activity is constantly changing. It's like the water of the stream: it never sits still.**

**The connectome is like the bed of the stream: it guides the flow of the water, but over long timescales, the water also reshapes the bed of the stream.**

*Sebastian Seung*

# Connectivity in neuroscience

- **Structural connectivity**  
(estimating actual connections)
  - *Invasive* (tract tracing methods, 2 photon calcium imaging)
  - *Non invasive* (Diffusion Tensor and Diffusion Spectral Imaging)
- **Functional connectivity**  
(based on temporal “co-variance”)
  - *Invasive* (intracranial recordings)
  - *Non invasive* (fMRI, M/EEG, simulated data)

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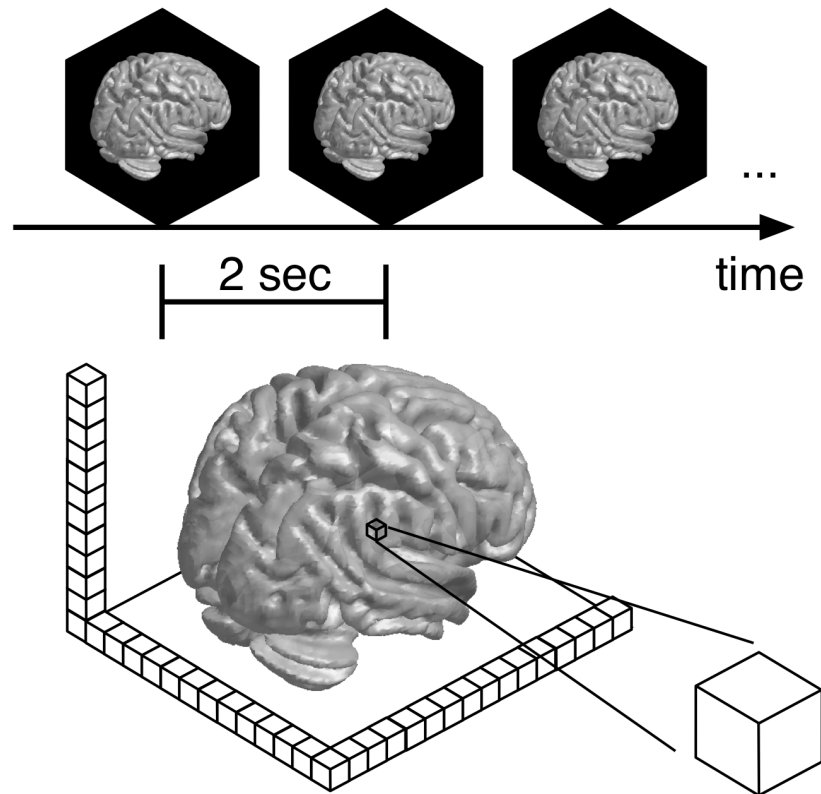
# Brain networks – general principles

- **Geographical networks:** nodes are located in 3D space and each node has specific function
- **Hierarchically structured:** with highly reproducible modules both at structural and functional level
  - Possibly arising from **wiring economy** Rivera-Alba, M. et al (2014). Wiring economy can account for cell body placement across species and brain areas. *Current Biology*, 24(3), R109-R110.
- **Multiple instances:**
  - we can estimate brain networks from **multiple subjects** or from the same subject at **multiple time points**
  - we can estimate brain networks from same subject using **multiple techniques**
- **Computing network-level statistics (p values!):**
  - what is significantly similar across subjects/time-points/techniques
  - What is significantly different (e.g. between groups or conditions)

# Connectomes, spatial scales and imaging techniques

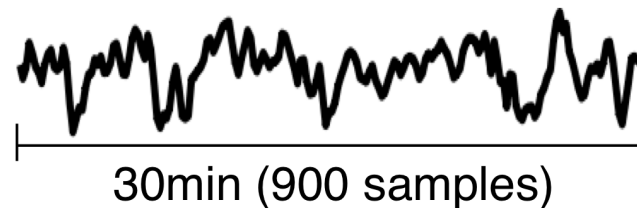
- **Microscopic scale** (i.e. actual neurons): worm, fruit fly, mouse retina [electron microscope or 2 photon imaging]
- **Mesoscopic scale** (based on neuronal tracing): mouse, macaque [usually viral neuronal tracing antero/retrograde transport -> one connection per animal]
- **Mesoscopic/macroscopic scale** (structural non invasive) [MRI, PET, DTI/DSI]
- **Mesoscopic/macroscopic scale** (functional non invasive) [fMRI, M/EEG]

# Functional magnetic resonance imaging (fMRI)



- We measure **multiple time series** at once
- **NOTE:** 1 voxel  $\rightarrow$   $5.5e6$  neurons  $4e10$  synapses (density  $\sim 1/1000$ ) [Logothetis 2008 Nature]

**Blood Oxygen Level signal**

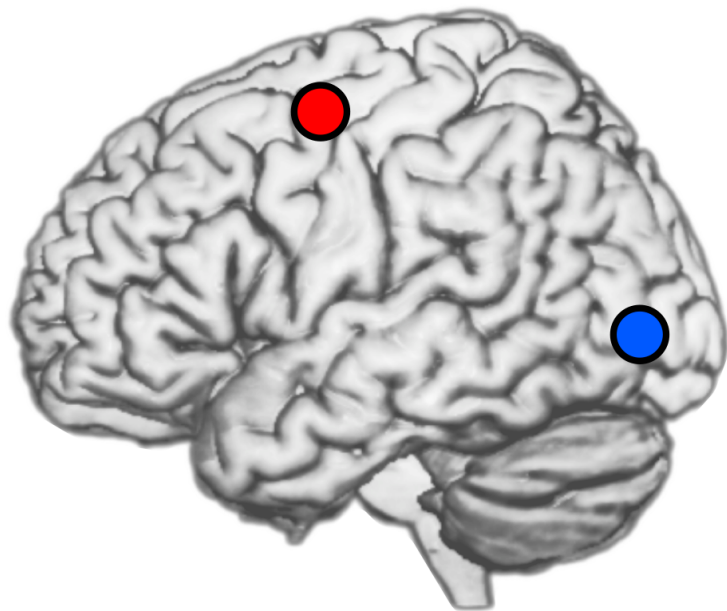




# Building a functional network

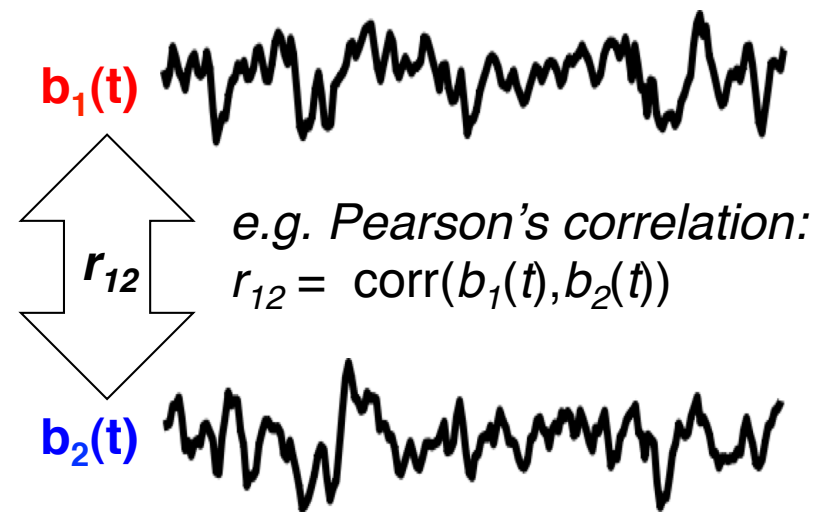
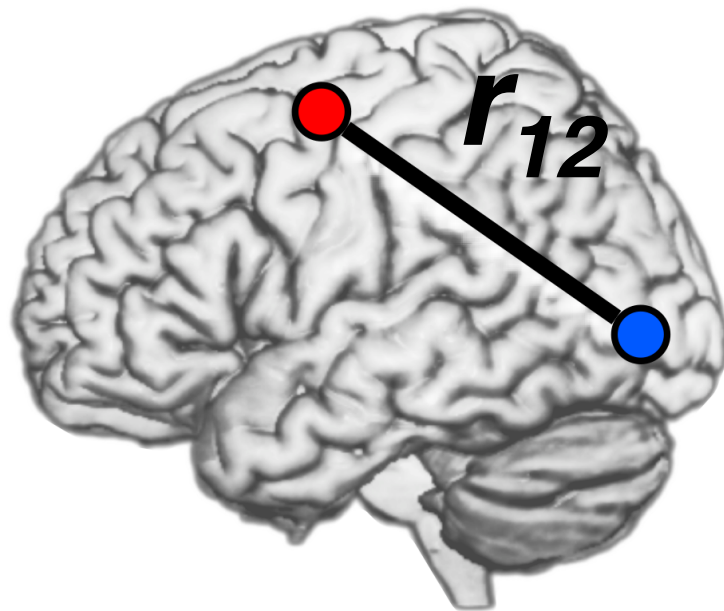
At each **node** we measure a **time series**

We compute their **similarity**



# Building a functional network

Similarity value used as **weight of the edge** between the two nodes. Repeat for each pair of nodes.



# PART 2

# Brain networks

# Current state

# Brain networks in animals

*Why is it important?*

- It's our only chance to estimate **actual connectomes**
- Only way to scale models from **micro/neuronal to macro/functional** and possibly the other way around
- Fascinating painstaking work of labelling each neuron and its synaptic contacts to **understand brain circuits**
- Videos from TED Talk “I am my Connectome”

# Brain networks in animals

## Known connectomes so far

	<i>C. elegans</i> hermaphrodite (or male)	Fruit fly medulla column(s)	Mouse retina IPL segment	Fruit fly brain	Mouse cortical column	Mouse brain	Human brain
Volume in mm <sup>3</sup>	0.0003	0.0001	0.001	0.071	0.12	450	1,400,000
Voxel dimensions X×Y×Z,nm	5×5×70	3.1×3.1×40	16×16×25	~8 <sup>3</sup>	~15 <sup>3</sup>	~15 <sup>3</sup>	~15 <sup>3</sup>
Voxel volume, nm <sup>3</sup>	~1750	380	6800	~500	~3400	~3400	~3400
Voxels, x10 <sup>12</sup>	0.13	0.3	0.15	142	35	130K	280M
Neurons traced	302	379	950	~10 <sup>5</sup>	~10 <sup>4</sup>	~10 <sup>8</sup>	~10 <sup>11</sup>
Synapses annotated	7283	8637	-	~10 <sup>8</sup>	~10 <sup>8</sup>	~10 <sup>12</sup>	~10 <sup>15</sup>
Proof time (person-year)	>10	10	10	4700	1200	4.5M	14G

Plaza, S. M., Scheffer, L. K., & Chklovskii, D. B. (2014). Toward large-scale connectome reconstructions. *Current opinion in neurobiology*, 25, 201-210.

# Brain networks in animals & network science – Resources

- All connectomes known so far stored in **multiple online databases**
  - *Open Connectome* <http://openconnectome.org/graph-services/download/>
  - *Allen Institute Brain Atlases* <http://connectivity.brain-map.org/>
  - *NIH Blueprint non-human primate atlas* <http://www.blueprintnhpatlas.org/>
- Oh, S. W., Harris, J. A., Ng, L., Winslow, B., Cain, N., Mihalas, S., ... & Mortrud, M. T. (2014). A mesoscale connectome of the mouse brain. *Nature*, 508(7495), 207-214.
- Shih, C. T., Sporns, O., Yuan, S. L., Su, T. S., Lin, Y. J., Chuang, C. C., ... & Chiang, A. S. (2015). Connectomics-based analysis of information flow in the Drosophila brain. *Current Biology*, 25(10), 1249-1258.
- Rubinov, M., Ypma, R. J., Watson, C., & Bullmore, E. T. (2015). Wiring cost and topological participation of the mouse brain connectome. *Proceedings of the National Academy of Sciences*, 112(32), 10032-10037.

# Brain networks across species/scales (universality?)

A



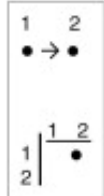
*Drosophila melanogaster*



*Mus musculus*



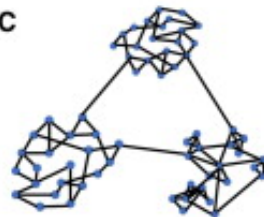
*Macaca mulatta*



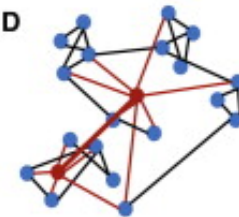
B



C



D



Current Biology

Kaiser, M. (2015).  
Neuroanatomy:  
Connectome Connects Fly  
and Mammalian Brain  
Networks. *Current Biology*,  
25(10), R416-R418.

- Goulas, A., et al. (2014).  
Comparative analysis of the  
macroscale structural connectivity in  
the macaque and human brain.  
*PLoS Comput Biol*, 10(3), e1003529.

- van den Heuvel, MP, et al. (2015).  
Bridging cytoarchitectonics and  
connectomics in human cerebral  
cortex. *The Journal of Neuroscience*,  
35(41), 13943-13948.

# From animals to humans



# Connectivity in neuroscience... non invasively

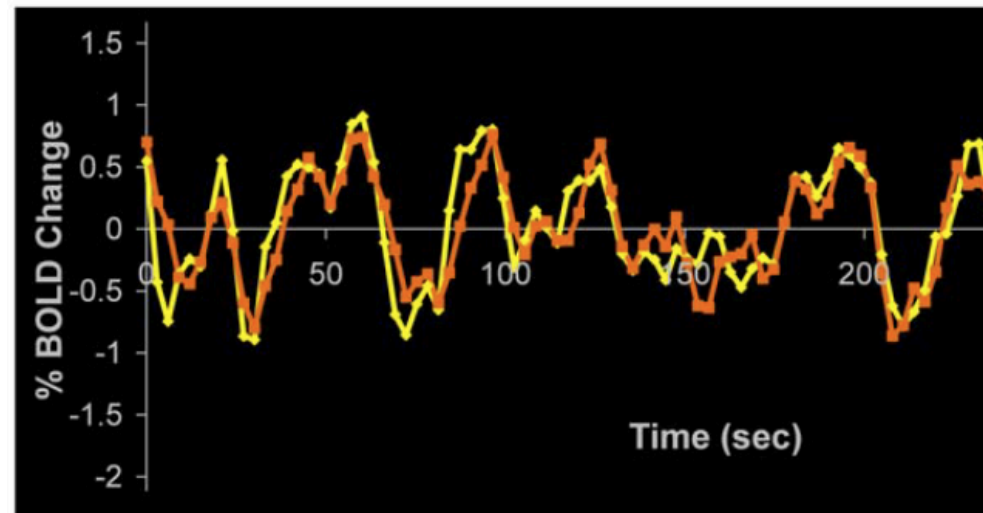
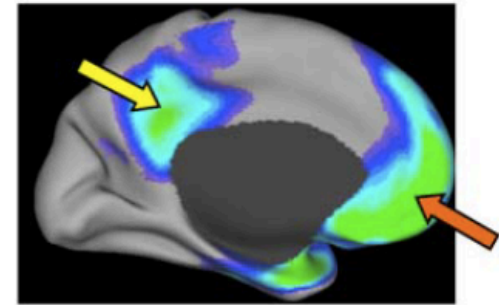
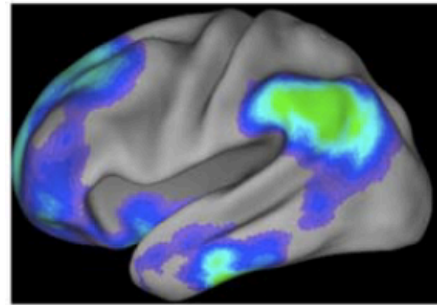
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Craddock, et al. (2013). Imaging human connectomes at the macroscale. *Nature Methods*, 10(6), 524–539. (\*)

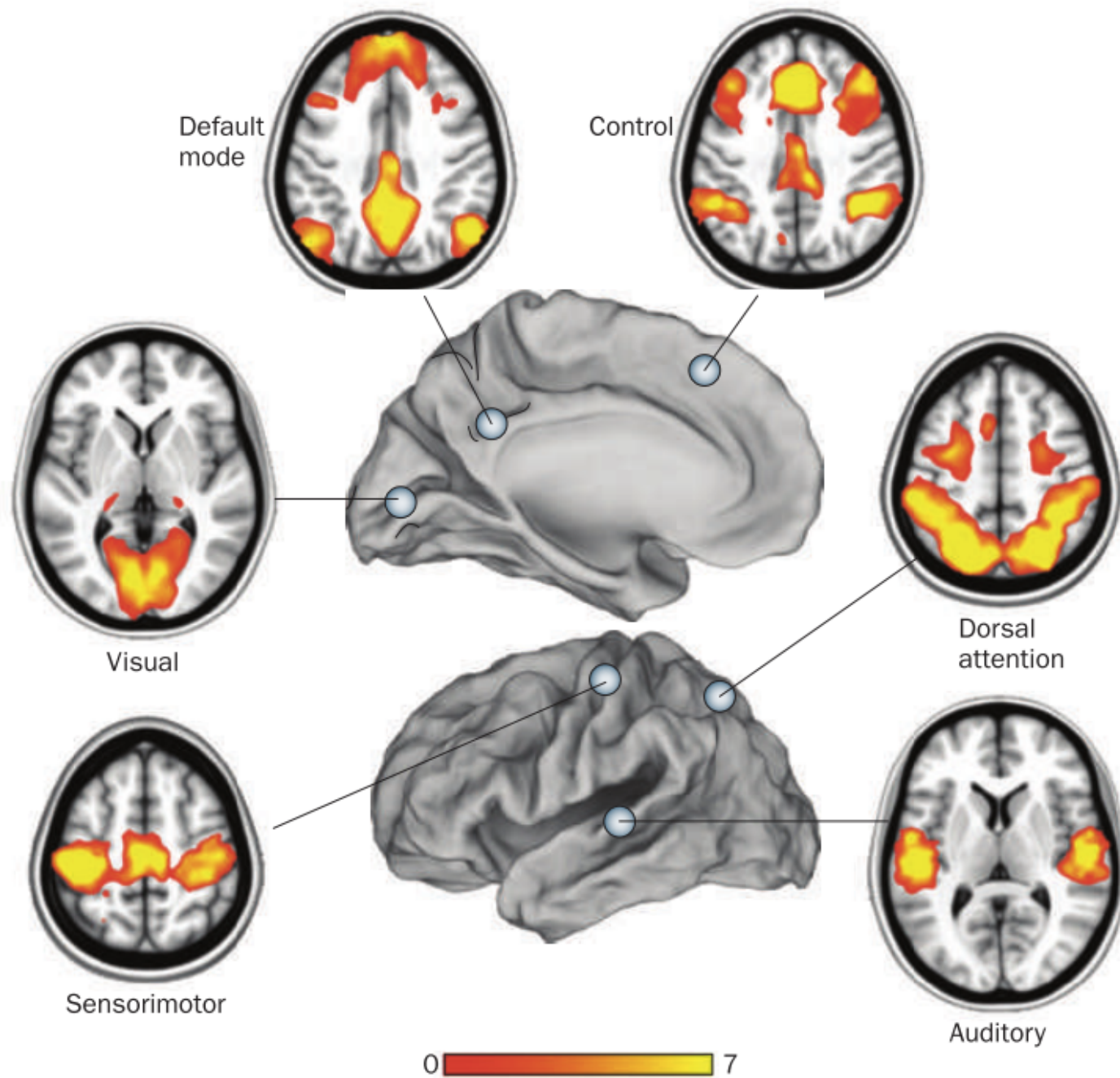
# The brain at rest

# The activity of the brain at rest is ideal for estimating the connectome

By looking at regions that change together in time we can **estimate their connectivity**



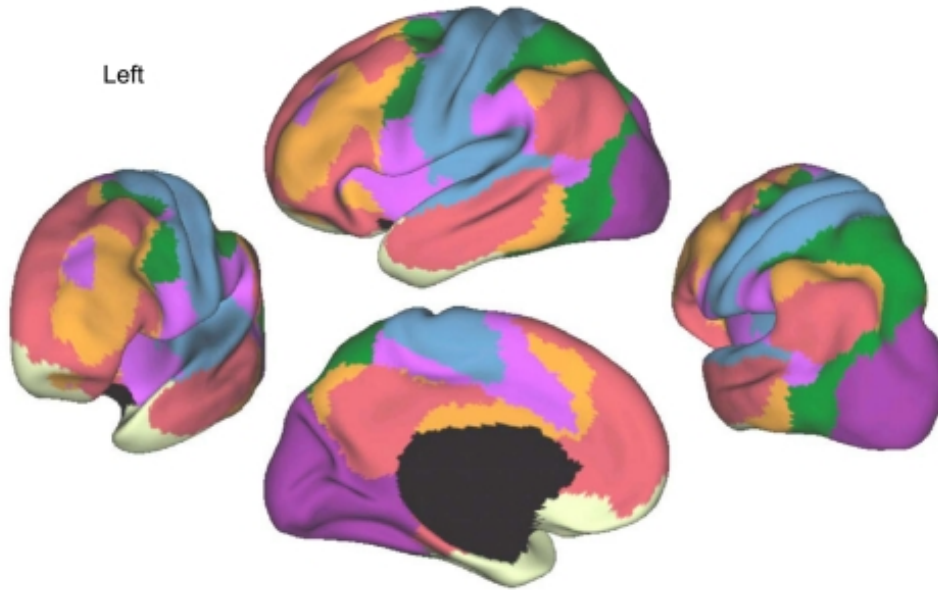
Raichle, M. E. (2010). Two views of brain function. *Trends in Cognitive Sciences*, 14(4)



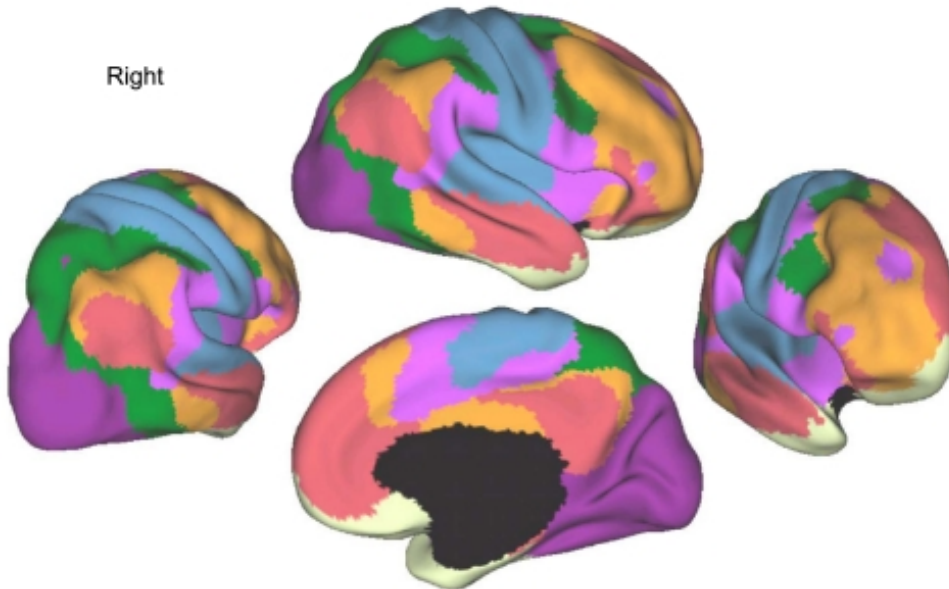
Zhang, D., & Raichle, M. E. (2010). Disease and the brain's dark energy. *Nature reviews. Neurology*, 6(1), 15–28.








7-Network Parcellation (N=1000)

Left



Right

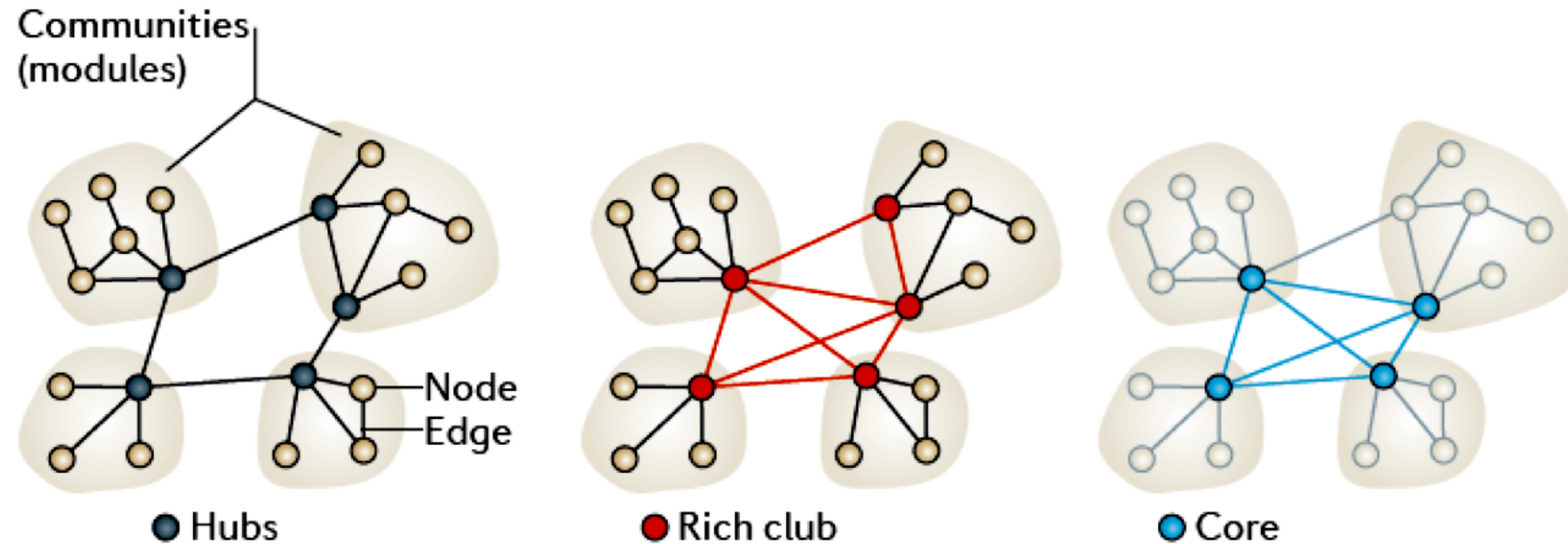


-  Purple (Visual)
-  Blue (Somatomotor)
-  Green (Dorsal Attention)
-  Violet (Ventral Attention)
-  Cream (Limbic)
-  Orange (Frontoparietal)
-  Red (Default)

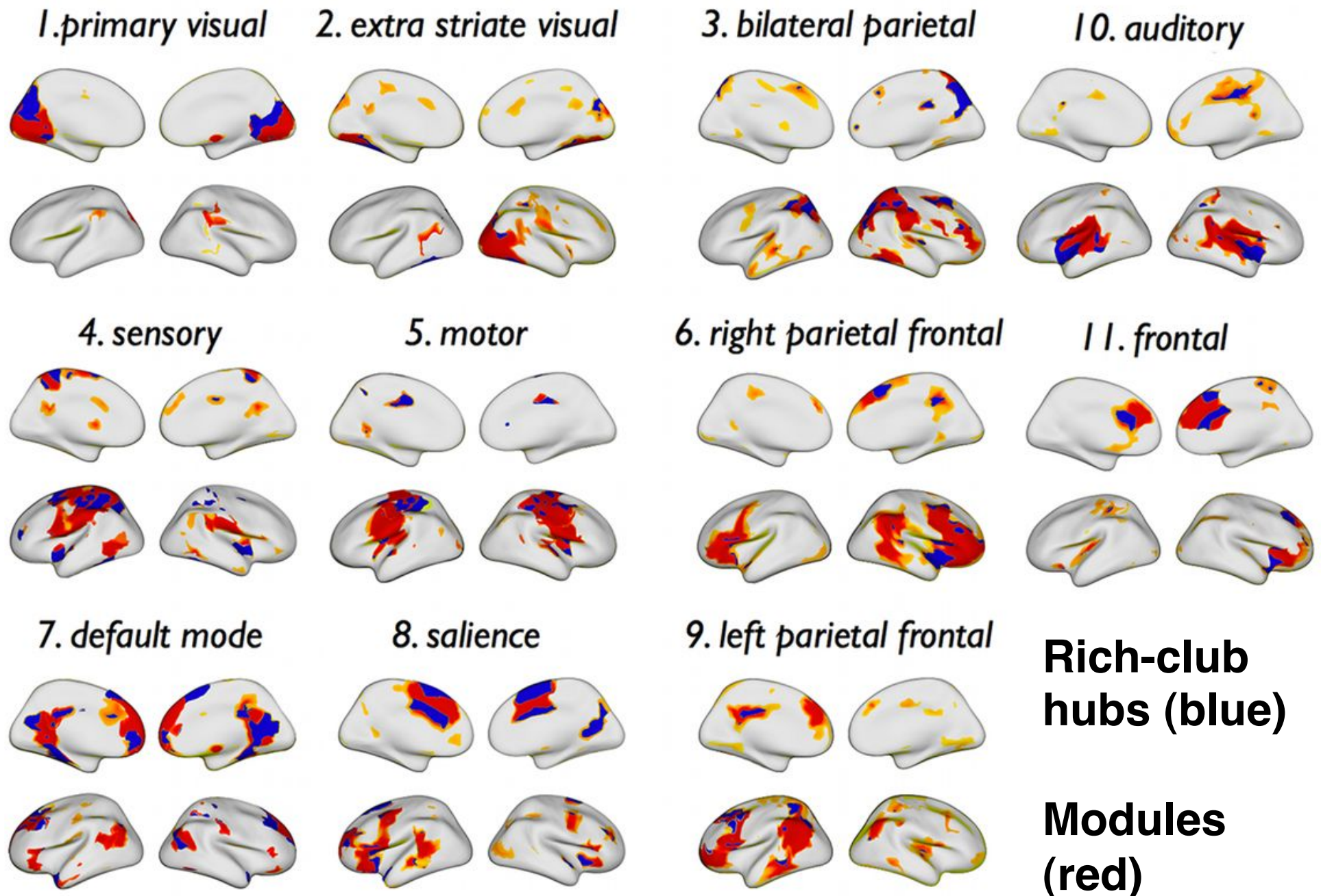
Yeo et al. (2011)

The organization of the human cerebral cortex estimated by intrinsic functional connectivity  
J Neurophysiol. 106(3):1125-65.

# *A rich club* of strong hubs in multiple modules is at the core of the human brain



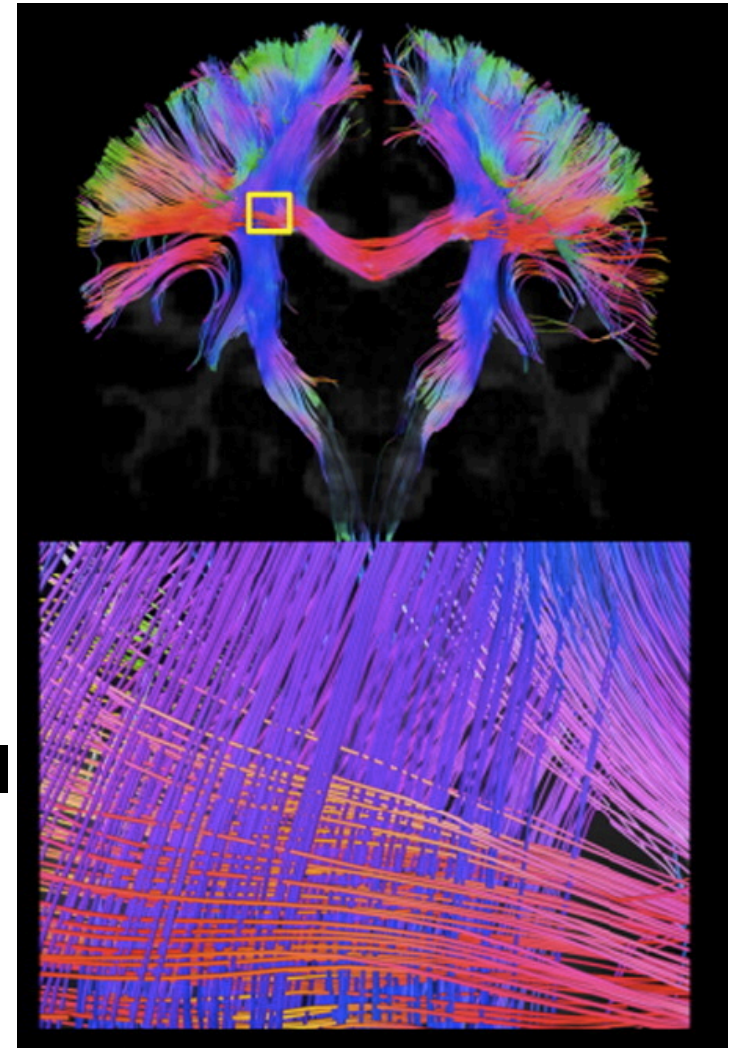
Bullmore, E., & Sporns, O. (2012). The economy of brain network organization. *Nature reviews. Neuroscience*, 13(5), 336–49



Van den Heuvel & Sporns (2013). JNeurosci.

# Brain networks in humans – data

- Multiple data-sharing efforts using **large subjects**
- Functional data: **resting state/task fMRI & MEG** (e.g. human connectome project: 900 subjects with fMRI and 500 with MEG)  
<http://www.humanconnectome.org/data/>
- Structural data DSI:  
<http://www.humanconnectomeproject.org/data/>
- Many initiatives also for various **mental disorders** (Autism, Alzheimer's Disease, Schizophrenia, ADHD)  
[usually MRI and fMRI]





# PART 3

# Brain networks

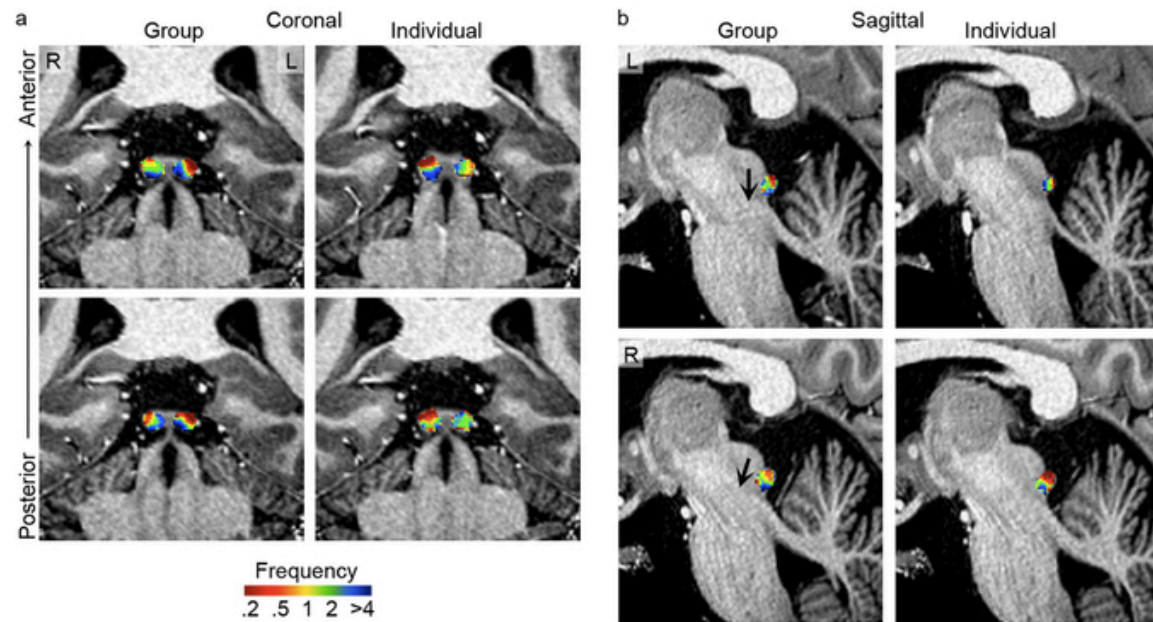
# Challenges

# 1D – Space

# What is a node?

- Many are happy to assume that a node is
  - A region of interest identified histologically or from another dataset
  - A region where the signal of neighbour voxels is more similar (i.e. dimensionality reduction e.g. via ICA)
  - A single voxel

More powerful scanners (7T) -> higher SNR -> single voxels will be more and more important



Moerel et al (2015) Processing of frequency and location in human subcortical auditory structures SciRep

# The spatial resolution challenge

- How to build the network?
- More nodes -> exponential growth of the network
  - *6x6x6mm voxels -> ~6000 nodes -> ~18e6 undirected connections [manageable]*
  - *4x4x4mm voxels -> ~17000 nodes -> ~130e6 undirected connections [huge memory]*
  - *2x2x2mm ??? -> needs distributed computing approach*
- Novel methods are using distributed computing:
  - *Freeman, J., Vladimirov, N., Kawashima, T., Mu, Y., Sofroniew, N. J., Bennett, D. V., ... & Ahrens, M. B. (2014). Mapping brain activity at scale with cluster computing. Nature methods, 11(9), 941-950.*
- How to deal with storage? Storing all the links? Where to draw the line?
- Network scientists should work with neuroscientists starting from the modelling of the time series/network.

# 2D – Individuals

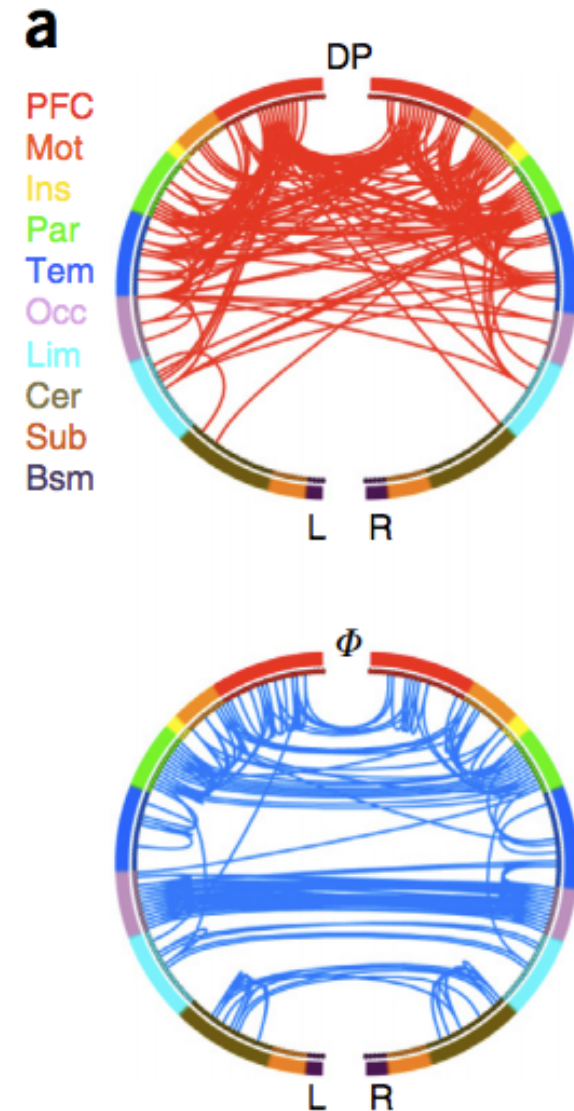
# The average brain

- As a first step, it is informative to look at the **average brain**
- However the **average brain does not actually exist**
- **Variability** between individuals and on the same individual over time
- First efforts
  - *Machine learning: fingerprinting of brains*
  - *Large scale modelling (canonical correlation) between individual behavioural variables and connectivity patterns*
  - *Comparing network of individuals at brain level vs behavioural vs phenotypical etc*

# Connectivity fingerprints **a**

- Red links: highly unique
- Blue links: highly consistent

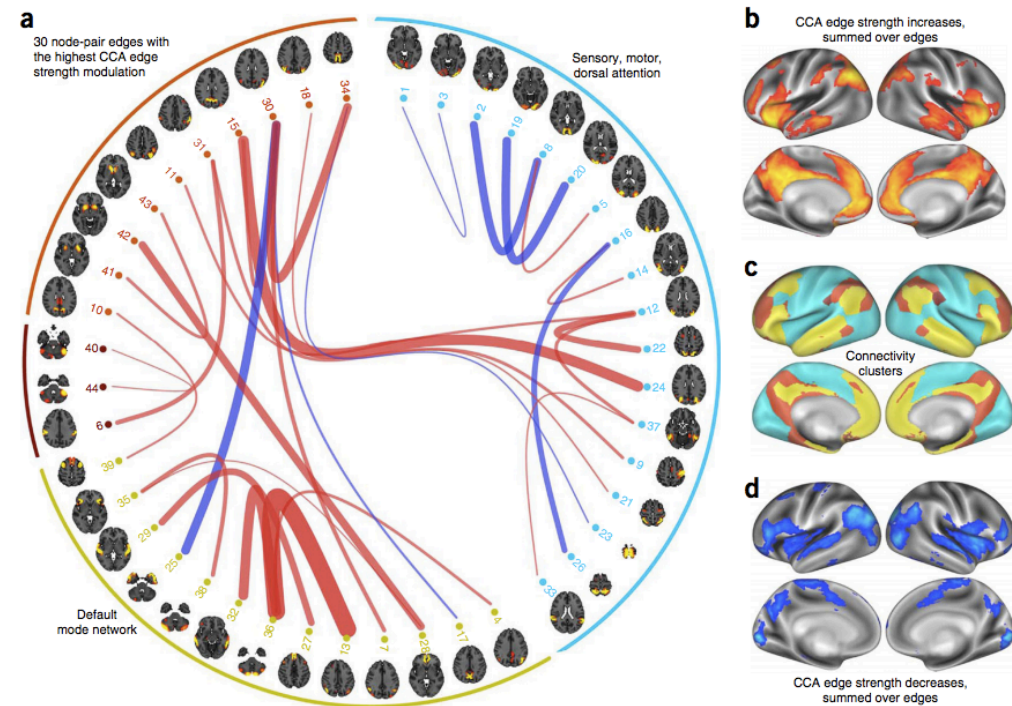
Finn, E. S., Shen, X., Scheinost, D., Rosenberg, M. D., Huang, J., Chun, M. M., ... & Constable, R. T. (2015). **Functional connectome fingerprinting: identifying individuals using patterns of brain connectivity.** Nature neuroscience.



# Matching individual traits with connectivity

- 461 resting subjects
- 158 variables
  - (age, gender, income, IQ, smoking habits, etc)
- ‘Good’ and ‘bad’ humans

Smith SM, Nichols TE, Vidaurre D, Winkler AM, Behrens TE, Glasser MF, Ugurbil K, Barch DM, Van Essen DC, & Miller KL (2015). **A positive-negative mode of population covariation links brain connectivity, demographics and behavior.** Nature Neuroscience

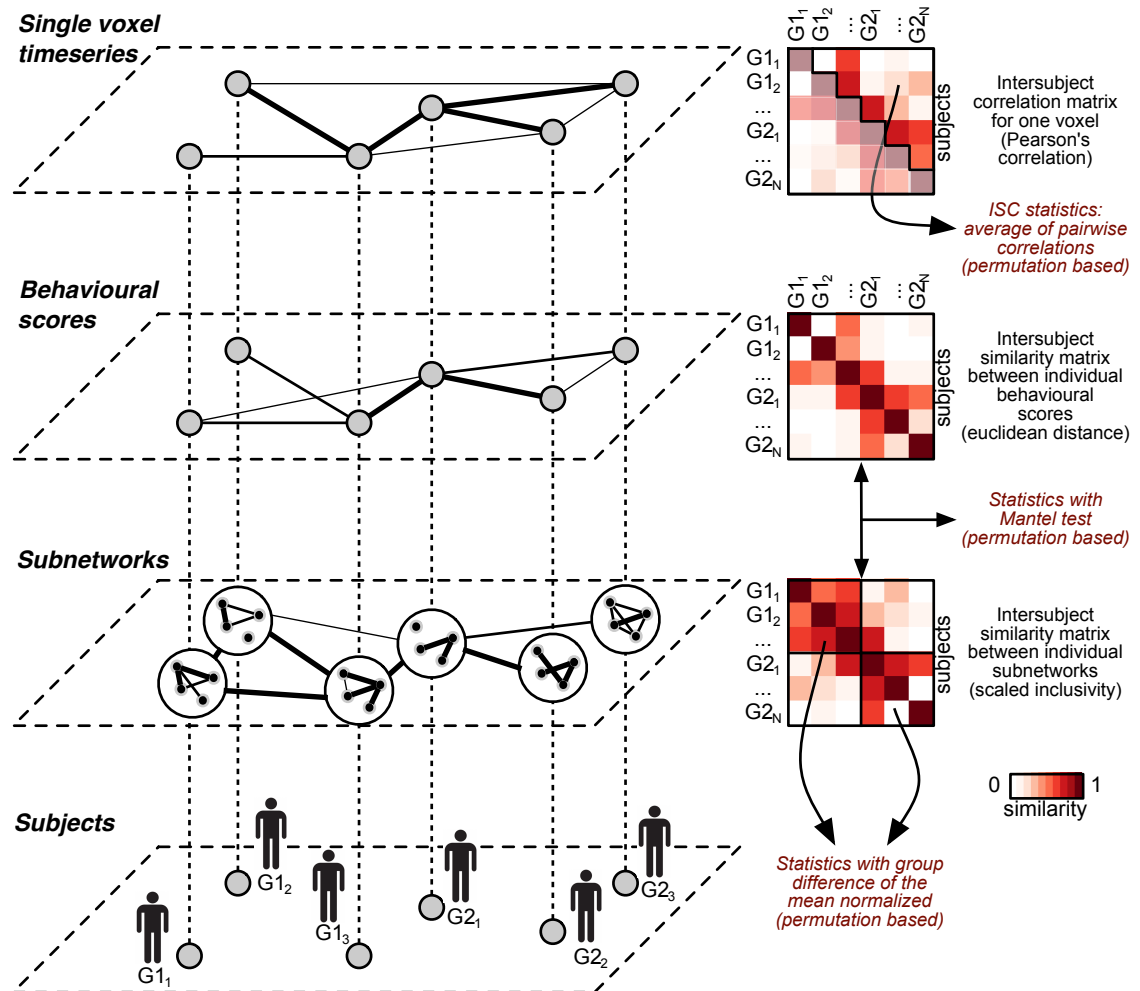




# Intersubject analysis framework

- Networks of subjects compared across dimensions
- **Mantel test** (comparison between similarity matrices)

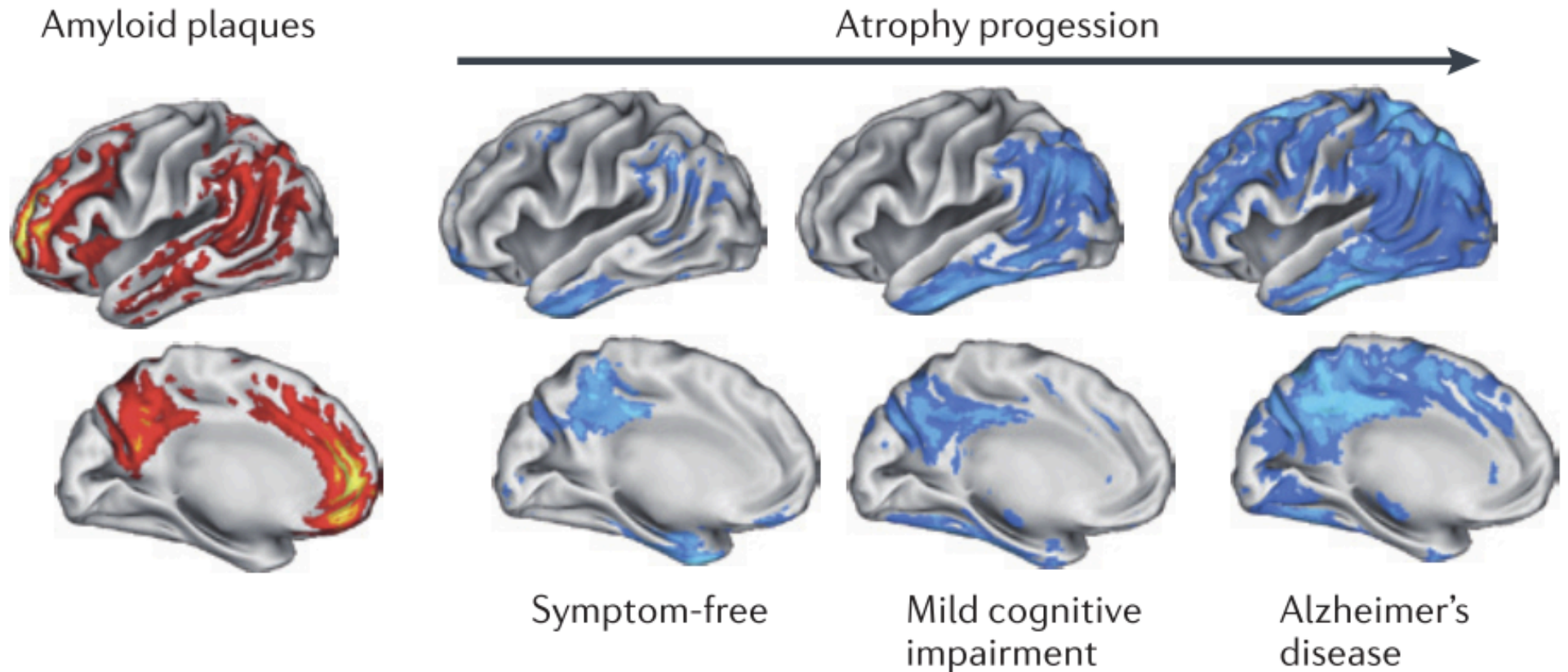
Glerean, E., et al. (in press)  
**Reorganization of functionally connected brain subnetworks in high-functioning autism**  
 Human Brain Mapping.



# 2D – individuals: mental health

# Alzheimer's disease

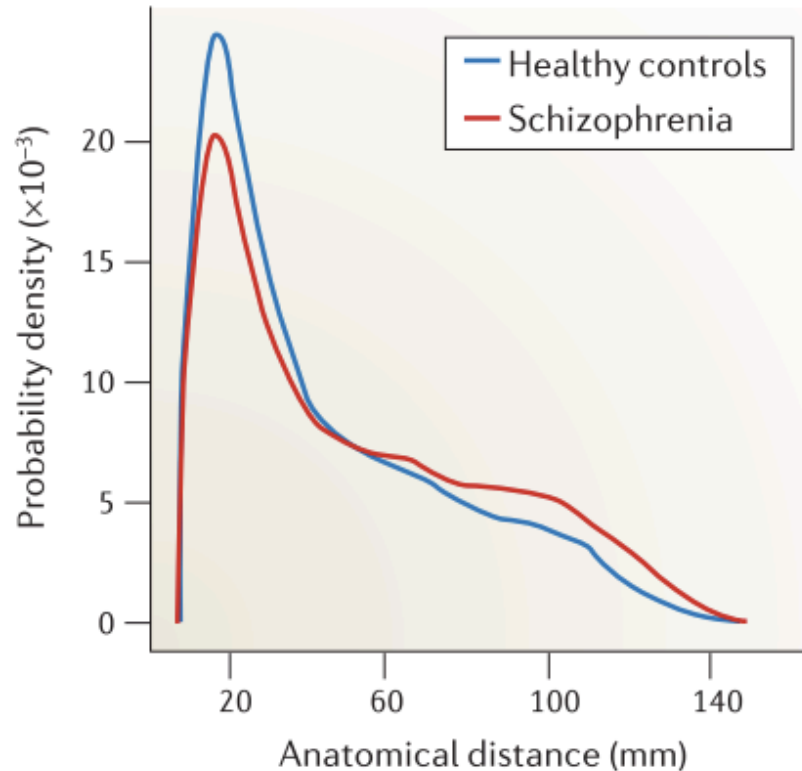
- The most expensive hubs are attacked by the disease



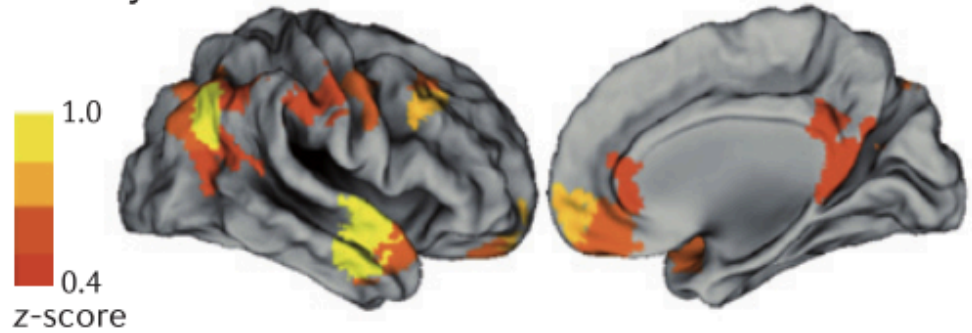
# Schizophrenia

Bullmore, E., & Sporns, O. (2012).  
The economy of brain network  
organization.

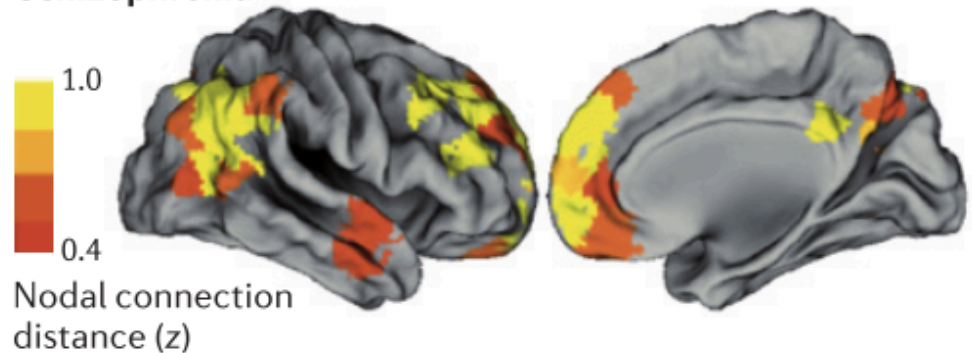
- **Unbalanced small-worldness**



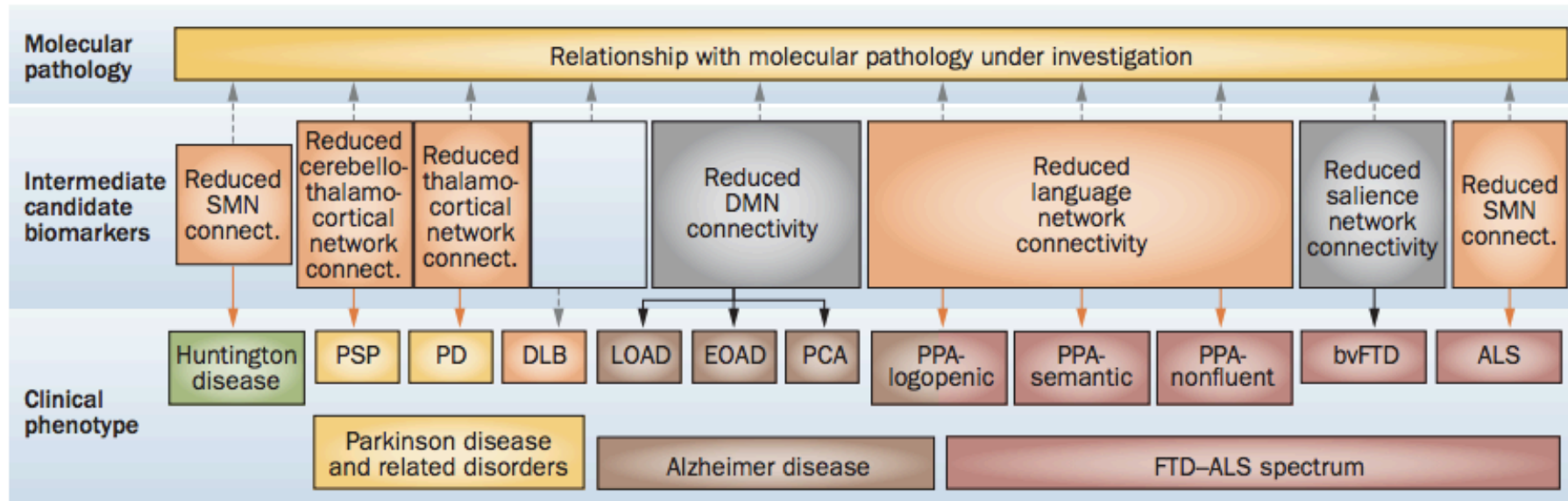
Healthy volunteers



Schizophrenia



# Brain networks as biomarkers



Pievani, M., Filippini, N., van den Heuvel, M. P., Cappa, S. F., & Frisoni, G. B. (2014). Brain connectivity in neurodegenerative diseases [mdash] from phenotype to proteinopathy. *Nature Reviews Neurology*, 10(11), 620-633. Chicago

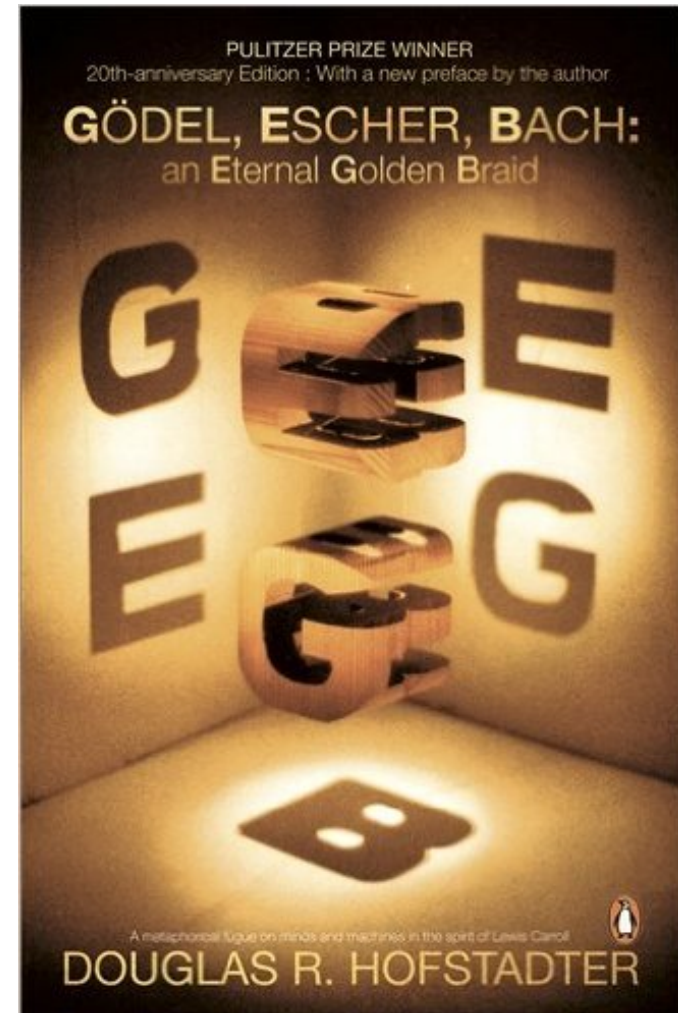
# How to deal with individuality?

- Other **approaches**?
- **Dimensionality scaling** (machine learning, ICA, subnetworks) is still a big limiting decision
- Modelling **noise vs signal**?
- It adds on top of the **other challenges** (especially on how to build the network)
- Healthy vs **clinical** groups

# 3D – Imaging modalities

# Brain networks across modalities

- Multiple imaging modalities
- Is it the same object projected over different planes?
- **Integrating modalities**





# Brain networks across modalities

- Successful efforts of integrating
  - *fMRI*
  - *M/EEG*
  - *DTI/DSI*
  - *MRI*
  - *Computational models*

*Reference list is long, I can email*

- Upcoming efforts of linking **PET** with DTI/DSI and functional connectivity

# Different modalities present different assumptions and artefacts

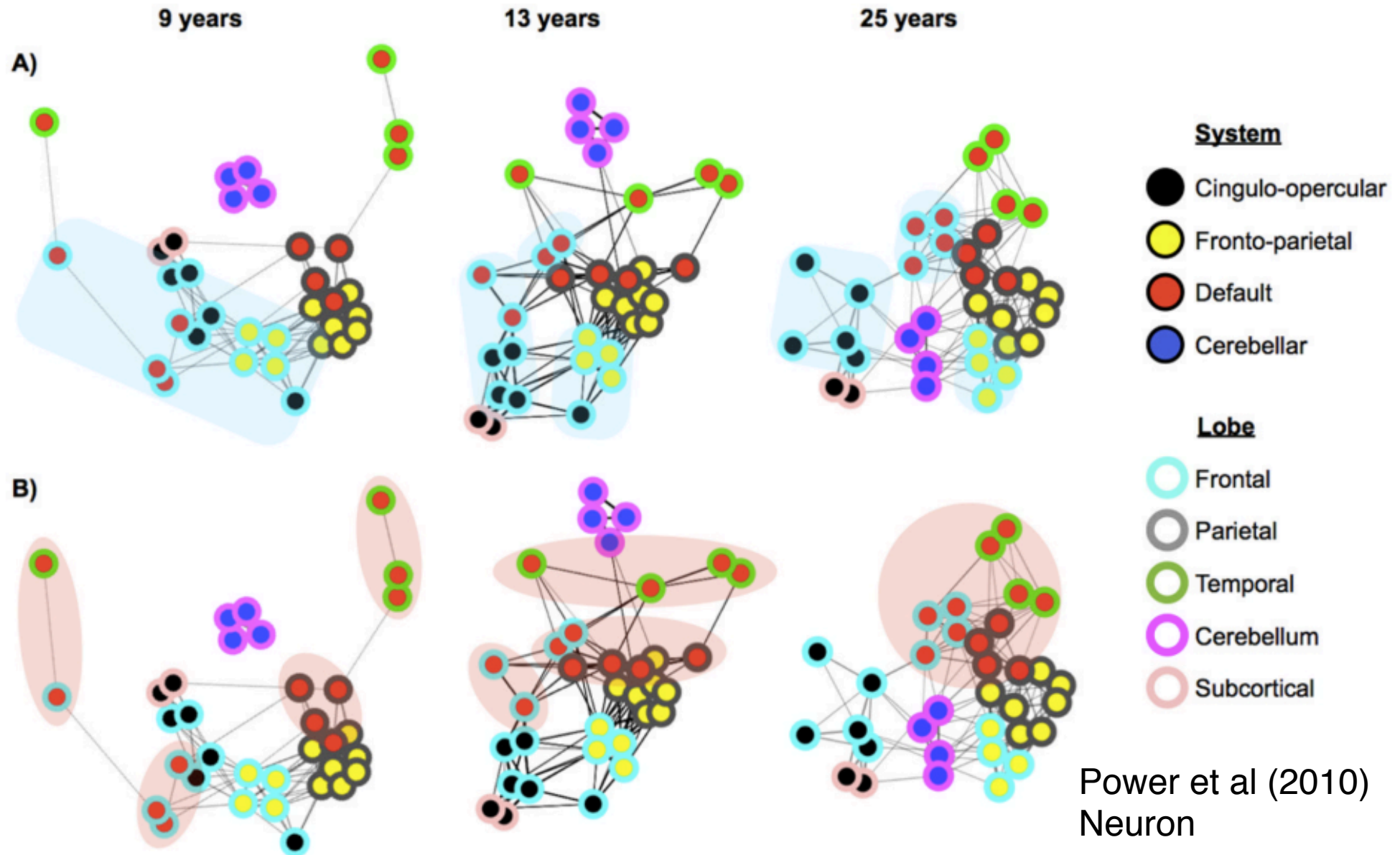
- How to **combine** all the types of networks from **same subject**?
- How to use one modality as ‘prior’ knowledge for another modality?
- How to summarize properties that generalize across modalities (or not!) [multiplex networks?]

# 4D – Time

# Brain networks in time

- At ms scales: interesting works works with MEG
- At second scales: sliding window connectivity and other measures
  - *Lots of disagreement in the field*
- Repeated measures for a single subject
  - *My connectome project <http://myconnectome.org/wp/> same subject scanned ~100 times*
  - *Some measures are stable some other covary with time of the day, time of the year etc.*
- Work across age groups (e.g. the developing brain)

# Network changes from childhood to adulthood



# Brain networks in time

- First we need **meaningful methods to estimated meaningful temporal networks**
- Can novel **advances in the network science of temporal networks** help neuroscientists?
- How to integrate temporal measures **across imaging techniques**

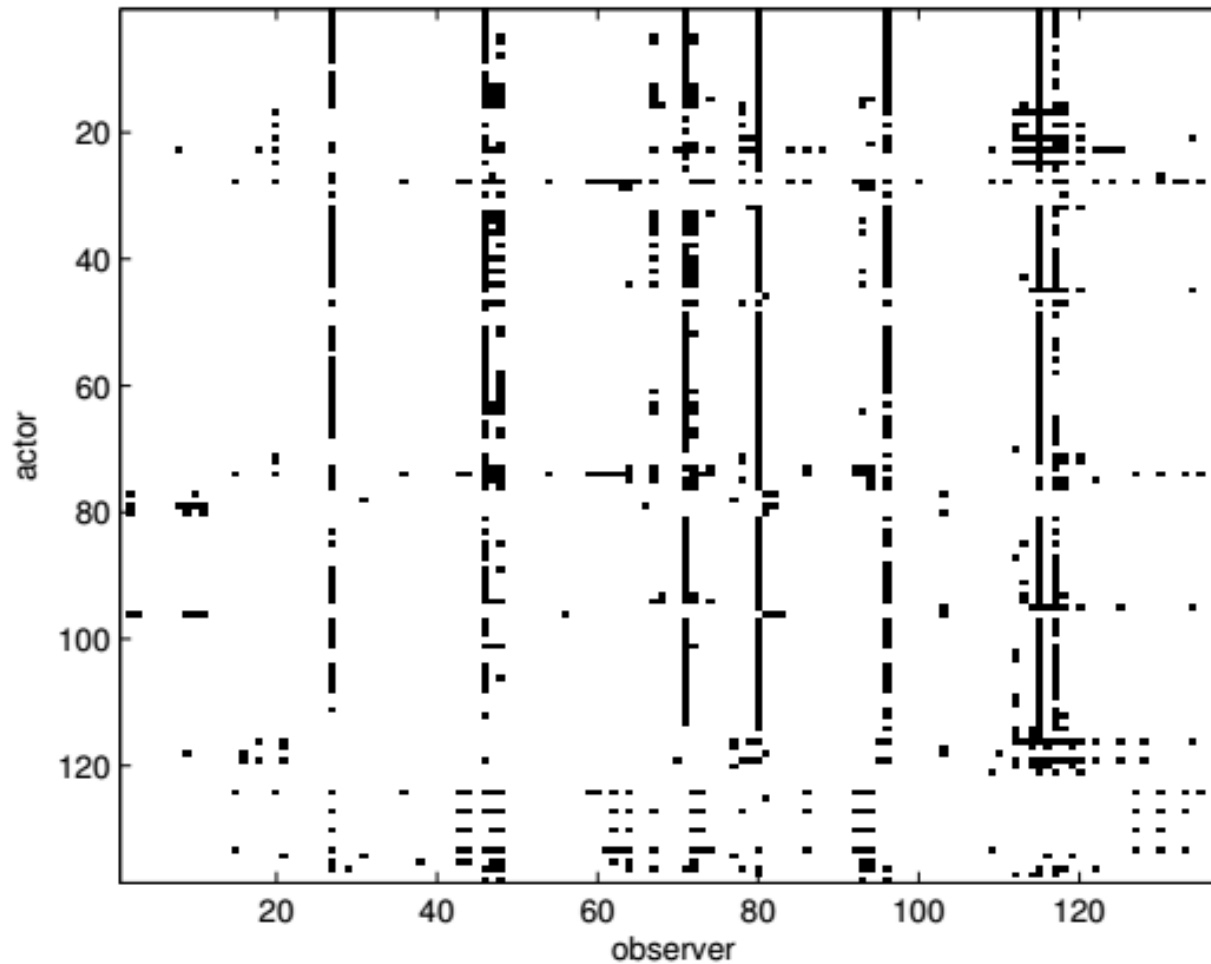
# 5D – Networks of networks

# Two persons neuroscience

- We interact with each other and our brain network activity influences others' brain networks activity
- Modelling a network between the two networks
- This network will not have usual properties (e.g. lower significance of strength of connections)
- How to model this? Bi-partite network? [so far I have been using permutation based approaches]



# Two persons neuroscience



# Conclusion

# Summary of challenges

- 1) **space** (increasingly larger networks due to higher quality of data)
- 2) **individual variability** (individual subjects with individual phenotypical/behavioral/genetic/clinical variables versus group averages)
- 3) **imaging modalities** (integration of multiple modalities/ types of data such as structural MRI, functional MRI, M/ EEG, PET, DTI/DSI, modeling, etc)
- 4) **time** (networks evolving in time at ms/seconds/hours/ years temporal scales)
- 5) **networks of networks** (two-persons neuroscience)



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# Brain networks Current state and future challenges

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