

Brain networks & Functional Connectivity

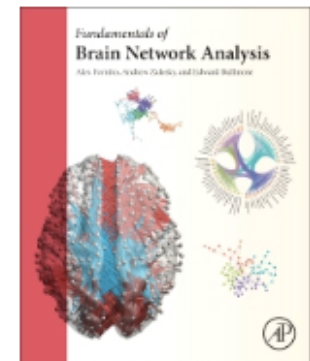
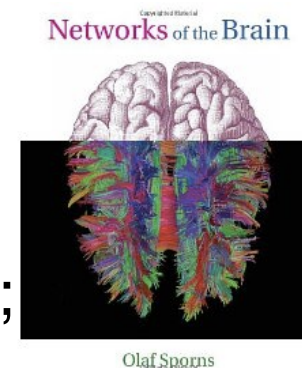
Enrico Glerean – web: www.glerean.com – twitter: @eglerean

Some fundamental references

- **Bullmore, E., & Sporns, O. (2012). The economy of brain network organization.** Nature reviews. Neuroscience, 13(5), 336–49.
- **Craddock, et al. (2013). Imaging human connectomes at the macroscale.** Nature Methods, 10(6), 524–539.

- **Networks of the Brain**
Sporns, O; 2010, MIT Press.
- **Fundamentals of Brain Network Analysis.** Fornito, Zalesky, Bullmore; 2017, Elsevier

(They can be taken as book exams <http://www.brain-mind.fi/courses.html>)



...and something in Finnish about network science

<https://www.researchgate.net/publication/242719764> *Kompleksisten verkostojen fysiikkaa*

Brain networks – Outline

- **Part 1 – Brain connectivity: ABC**
- **Part 2 – Brain network science**
- **Part 3 – Impact of this research**

Feel free to ask any question.

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...

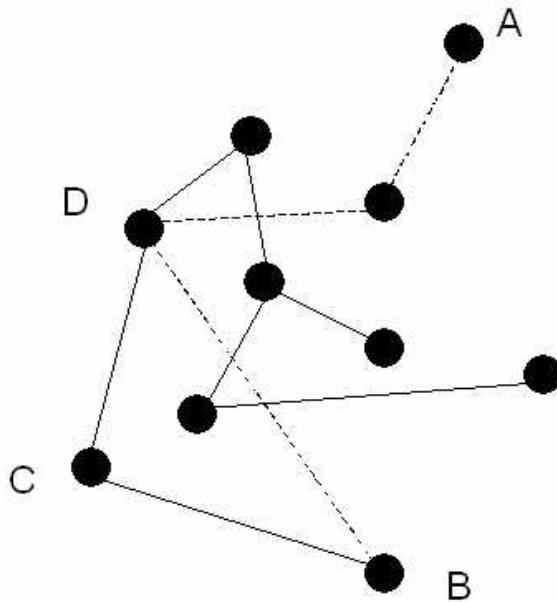
Why do we want to study brain networks?

- **The brain is a network with**
~ 10^{10} neurons and ~ 10^4 connections per neuron
- As for **genomics** in the 20th 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

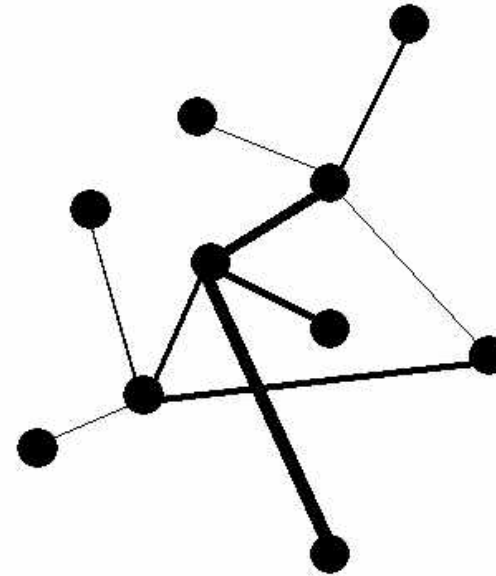
What is a network?

A (complex) network, a graph

Unweighted graph



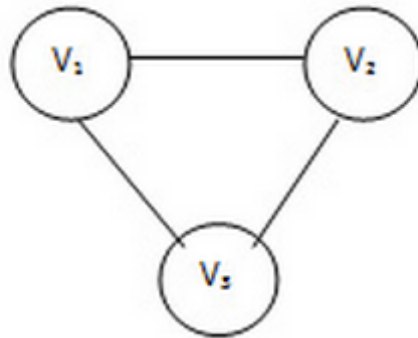
Weighted graph



Newman, M. E. J., **Networks: An introduction**. Oxford University Press, Oxford, March 2010.

Directed and undirected graphs

Undirected Graph



Directed Graph

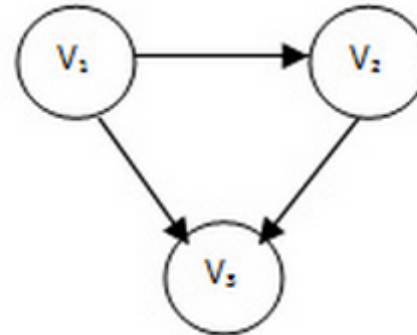


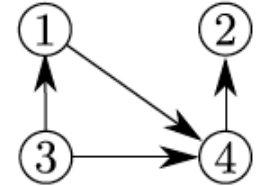
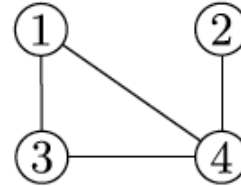
Figure 1: An Undirected Graph

Figure 2: A Directed Graph

Newman, M. E. J., **Networks: An introduction**. Oxford University Press, Oxford, March 2010.

Representation of networks

Source: Jari Saramäki's course slides



Adjacency list

i : neighbors

1: 3, 4

2: 4

3: 1, 4

4: 1, 2, 3

i : neighbors

1: 4

2:

3: 1, 4

4: 2

Adjacency matrix

$$a_{ij} = \begin{cases} 1 & \text{if } (j, i) \in E, \\ 0 & \text{if } (j, i) \notin E \end{cases}$$

$$\begin{bmatrix} 0 & 0 & 1 & 1 \\ 0 & 0 & 0 & 1 \\ 1 & 0 & 0 & 1 \\ 1 & 1 & 1 & 0 \end{bmatrix}$$

$$\begin{bmatrix} 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 \end{bmatrix}$$

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*

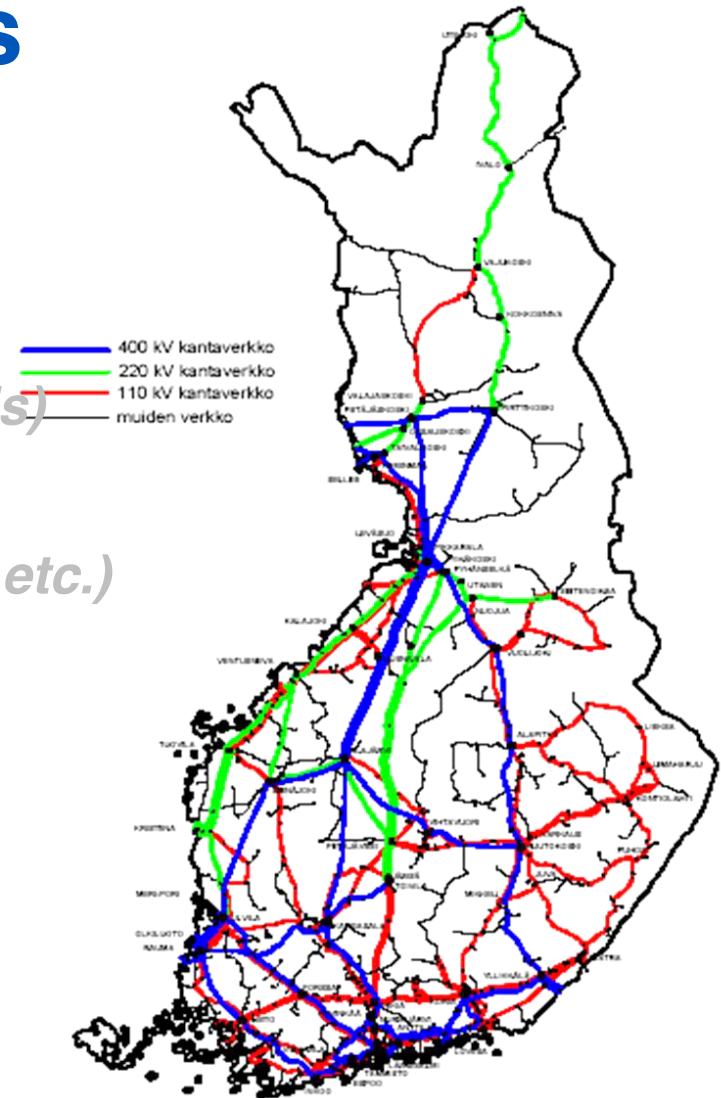
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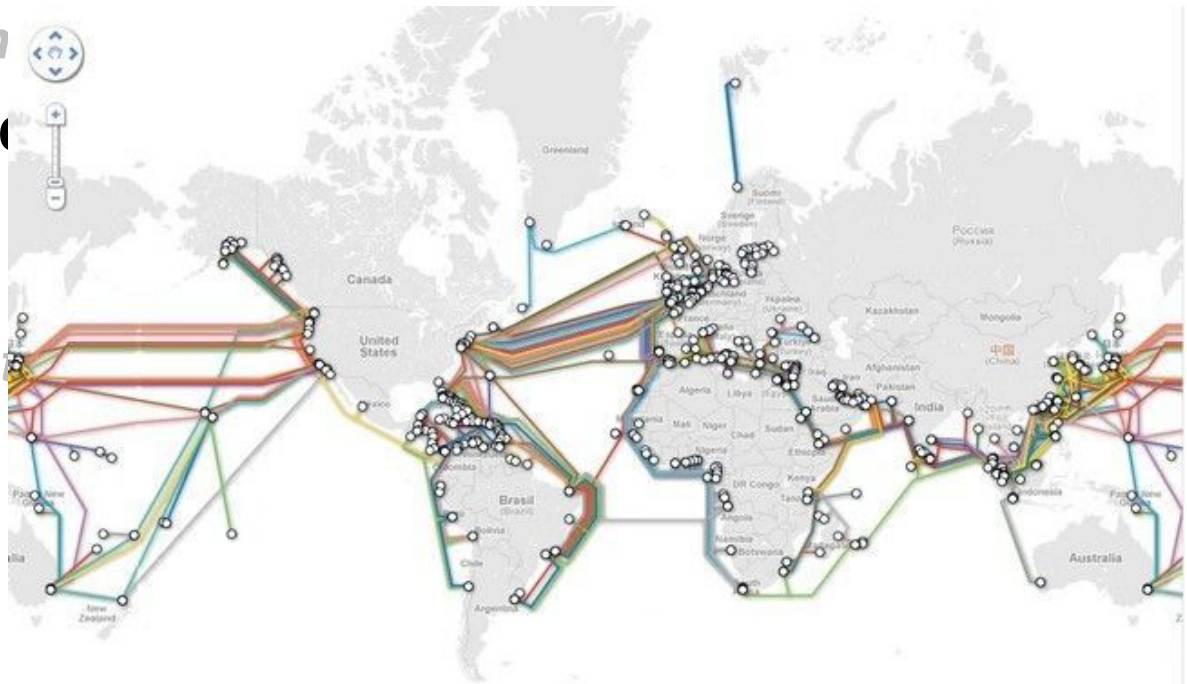
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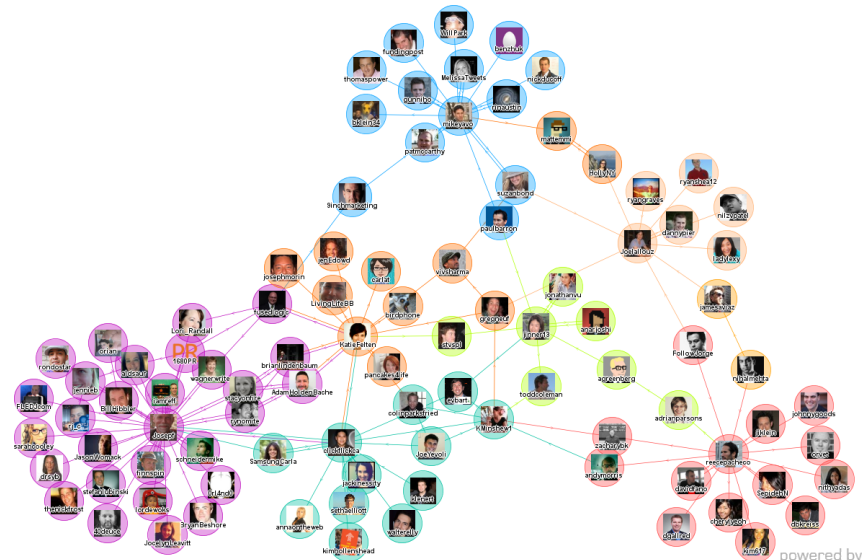
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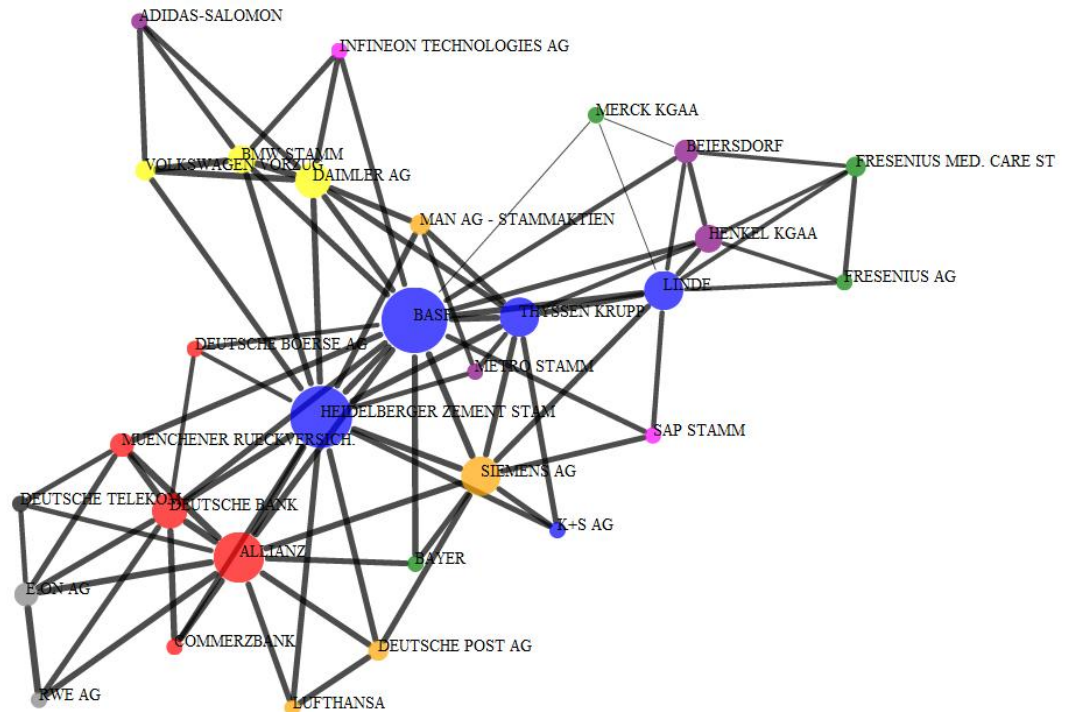
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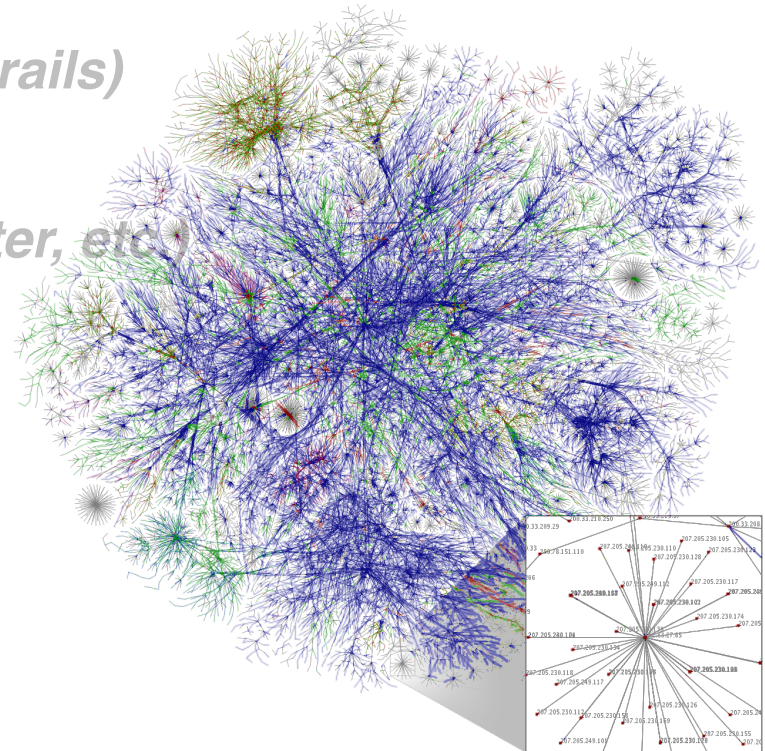
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Complex network

From Wikipedia, the free encyclopedia

In the context of [network theory](#), a **complex network** is a [graph](#) (network) with non-trivial [topological](#) features—features that do not occur in simple networks such as [lattices](#) or [random graphs](#) but often occur in real graphs. The study of complex networks is a young and active area of scientific research inspired largely by the empirical study of real-world networks such as [computer networks](#) and [social networks](#).

Contents [\[hide\]](#)

- 1 Definition
- 2 Scale-free networks
- 3 Small-world networks
- 4 See also
- 5 Books
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Definition [\[edit source | edit ^{beta} \]](#)

Most [social](#), [biological](#), and [technological networks](#) display substantial non-trivial topological features, with patterns of connection between their elements that are neither purely regular nor purely random. Such features

Network science



[Theory](#) · [History](#)

- [Graph](#) · [Complex network](#) · [Contagion](#)
- [Small-world](#) · [Scale-free](#) · [Community structure](#) · [Percolation](#) · [Evolution](#) · [Controllability](#) · [Topology](#) · [Graph drawing](#) · [Social capital](#) · [Link analysis](#) · [Optimization](#) · [Reciprocity](#) · [Closure](#) · [Homophily](#) · [Transitivity](#) · [Preferential attachment](#) · [Balance](#) · [Network effect](#) · [Influence](#)

Types of Networks

- [Information](#) · [Telecommunication](#)
- [Social](#) · [Biological](#) · [Neural](#)
- [Interdependent](#) · [Semantic](#)
- [Random](#) · [Dependency](#) · [Flow](#)

**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.

**What is brain
connectivity?**

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

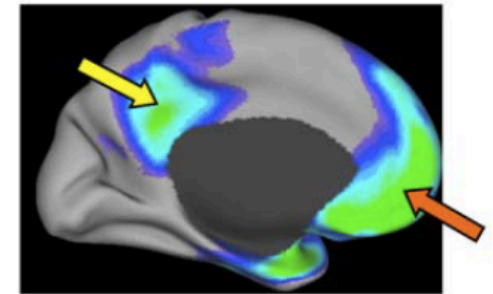
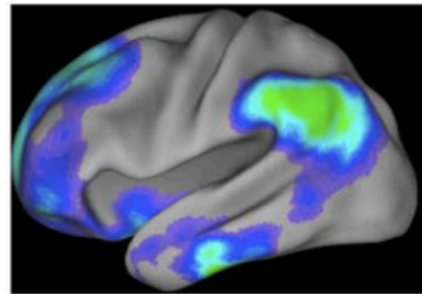
Connectivity in neuroscience

- **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)

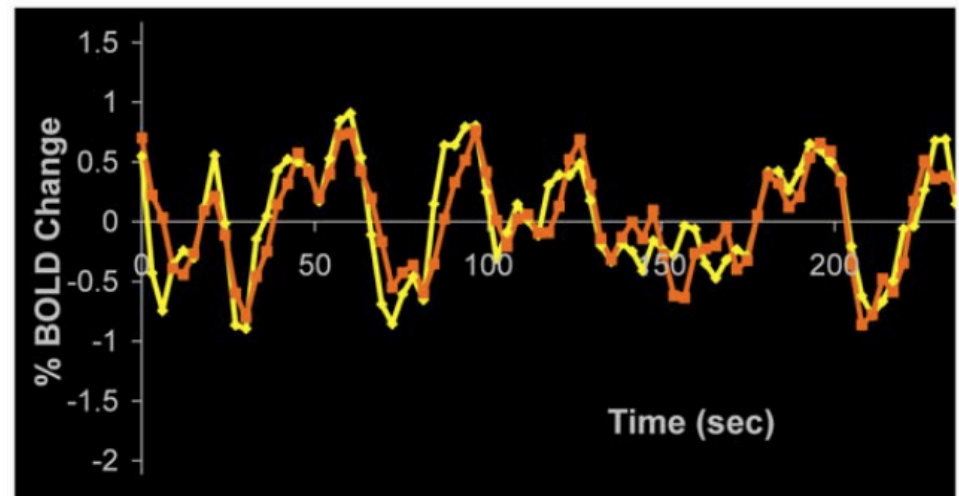
Craddock, et al. (2013). Imaging human connectomes at the macroscale. *Nature Methods*, 10(6), 524–539. (*)

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**

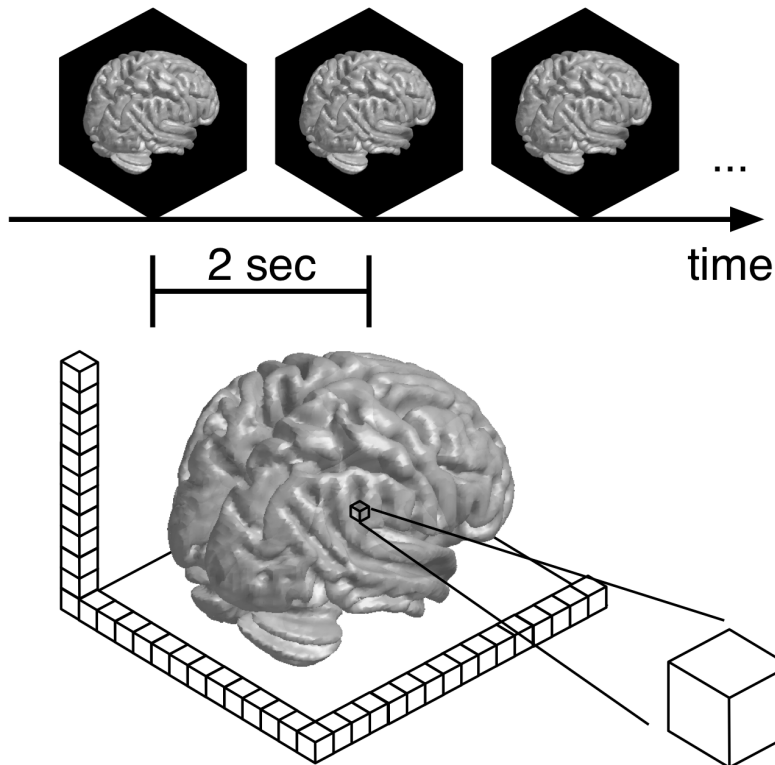


Raichle, M. E. (2010). Two views of brain function. *Trends in Cognitive Sciences*, 14(4)



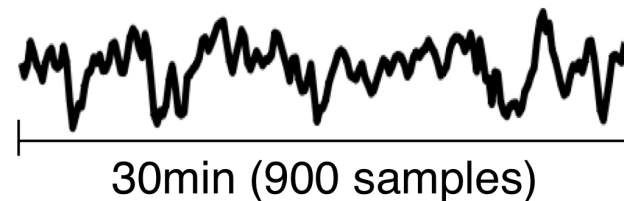
**How do we
compute a
functional brain
network?**

Functional magnetic resonance imaging (fMRI)



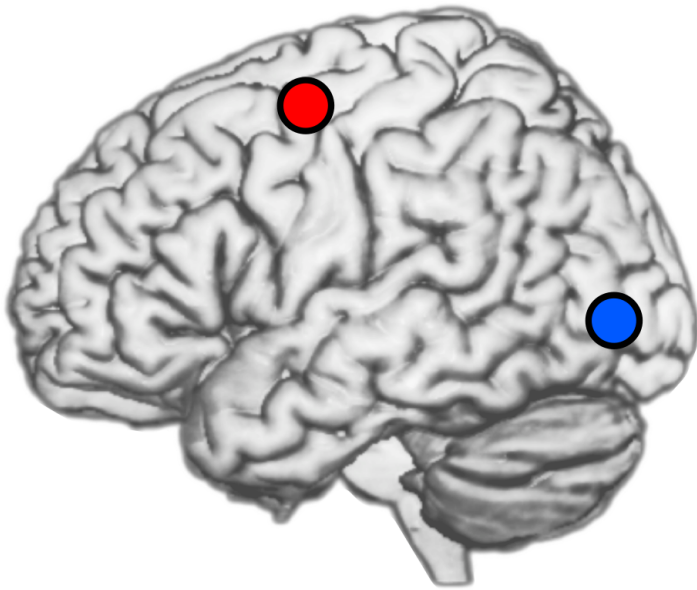
- We measure **multiple time series** at once
- We can consider them **independently** (e.g. GLM) or we can look at **mutual relationships**

Blood Oxygen Level signal



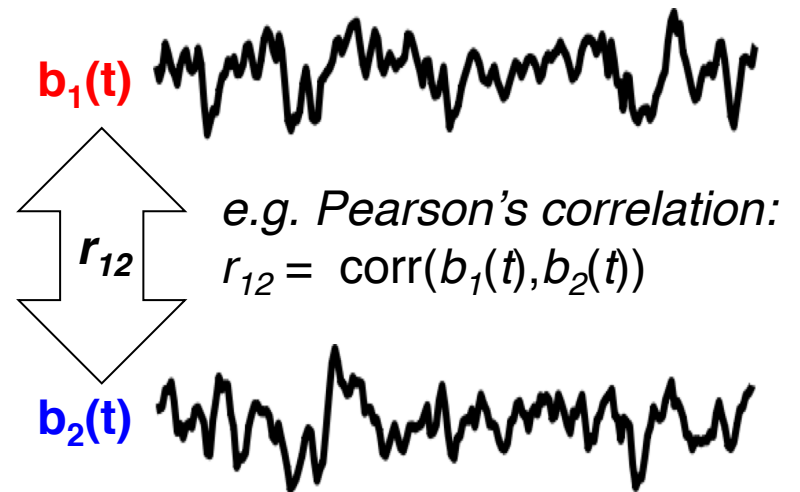
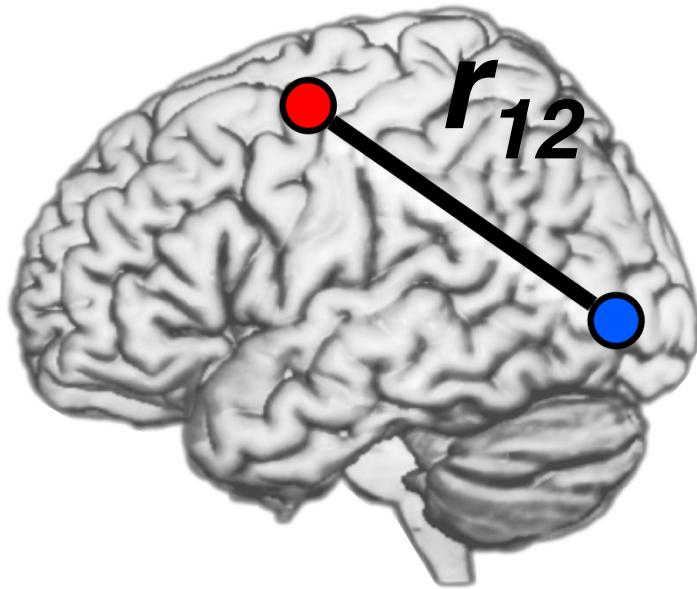
Building a functional network

At each **node** we measure a **time series**
We compute their **similarity**



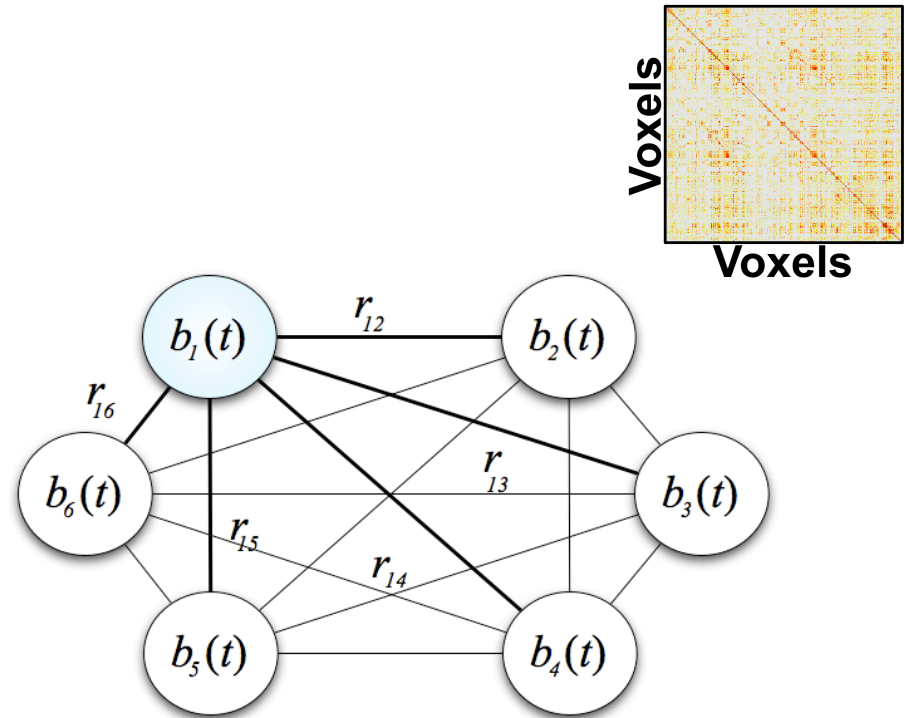
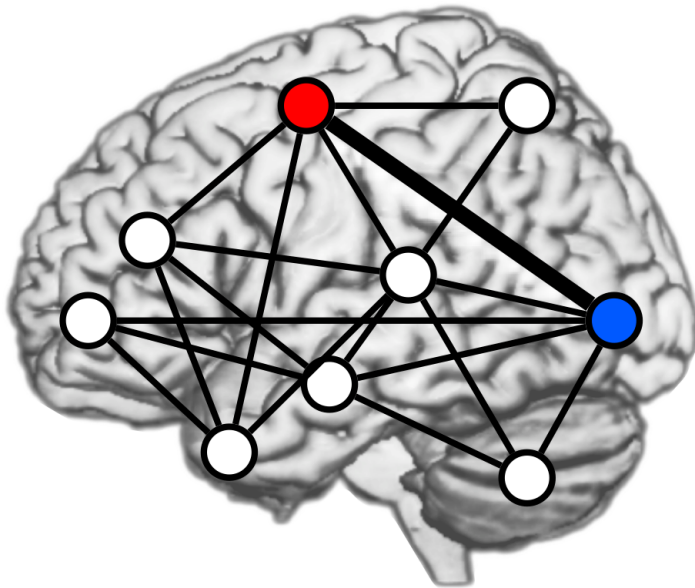
Building a functional network

Similarity value used as **weight of the edge** between the two nodes.



Building a functional network

Repeat for all pairs of nodes and we get the full functional network



What is a node in a functional brain network?

Nodes in fMRI FC

- **A node is a voxel**

- At 2mm isotropic voxels we have ~160K nodes, i.e. **12.8e9 links!**
- At 6mm isotropic voxels we have ~6K nodes, i.e. 18e6 links

- **A node is a region of interest (ROI)**

- We consider multiple voxels that are anatomically defined and derive one time series (using average or first PC) [e.g. atlas based: AAL atlas, Harvard Oxford atlas, UCLA atlas, Brainnetome]
- We consider a *seed*: a sphere centred at a specific location (usual size of diameter is 1cm) [based on literature, or nodes templates e.g. “Functional network organization of the human brain” Power JD, et al. Neuron. 2011 Nov 17; 72(4):665-78.
- WARNING: selection of ROIs can introduce bias

What is a link in a functional brain network?

Methods for similarity between time series

- **Pearson's correlation:** simple correlation
- **Partial correlation:** choose a pair of nodes, regress out all other nodes (more towards a multivariate than bivariate)
- **Regularised inverse covariance:** useful for short sess.
- **Mutual information:** (non)linear share of information
- **Coherence:** looking at cross-spectral similarity between a frequency representation of the time series
- Other methods related to task (**gPPI, beta series**)

Which one is the best method?

- The answer is: **it depends.**
- If you are looking for subtle differences e.g. between groups or between conditions, some more refined measures could perform better (Smith et al. showed **partial correlation, inverse covariance** and **Bayes-net** methods as winners)
- However, in most cases simple linear correlation is enough, see Hlinka, J., et al (2011). **Functional connectivity in resting-state fMRI: is linear correlation sufficient?** NeuroImage, 54(3), 2218–25.
doi:10.1016/j.neuroimage.2010.08.042

**Material not
covered**

Definitions

Functional and effective connectivity

- **Functional connectivity** = statistical dependencies among remote neurophysiological events
 - *Pairwise and “data driven”*
 - *No “direction” in the estimated connections*
- **Effective connectivity** = the influence that one neural system exerts over another
 - *Estimates the direction of influence between nodes in the network*
 - *Lag based methods (Granger causality)*
 - *Model based (Bayesian methods such as Dynamic Causal Modelling)*
 - *Higher order statistics via ICA (e.g. LiNGAM)*

Paradigms for functional connectivity

- **Resting state FC**

Looking at spontaneous BOLD activity while the subject is in the scanner

Correlated with anatomy

- **Task related FC**

The subject is performing a task with multiple conditions (usually block design or *naturalistic design*, i.e. a block design with longer blocks)

Task

1. The subject is doing a task

1. Task structure

1. *In **Blocks***

2. *As **Events** separated in time*

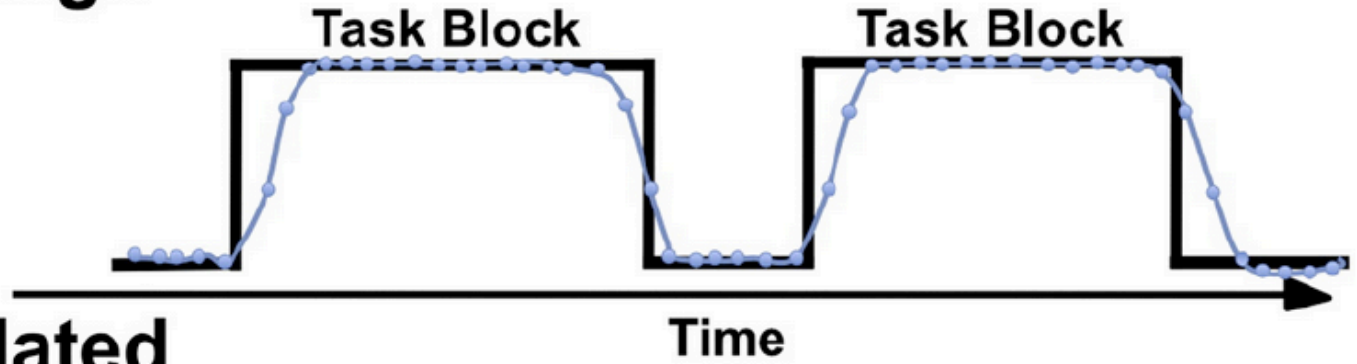
3. *As a **stream of events (naturalistic)***

2. Passive vs Active

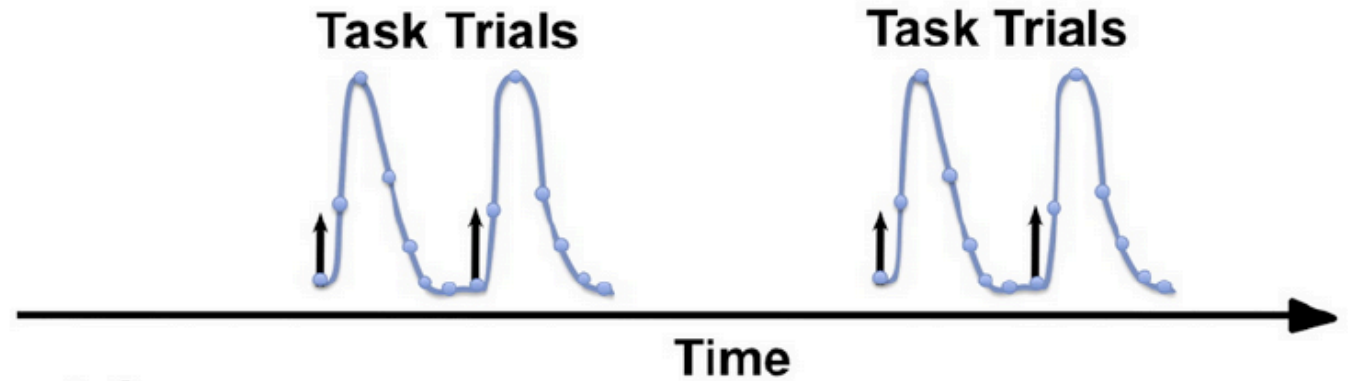
1. *Pressing a **button, etc***

2. ***Just watching and mentalizing***

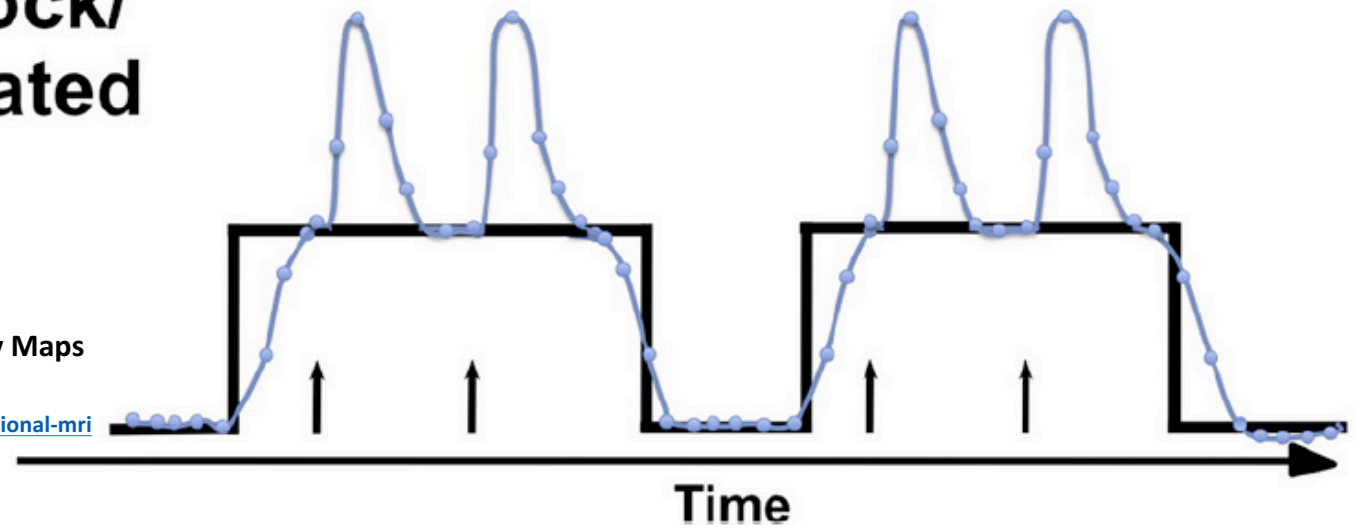
A) Block Design



B) Event-Related



C) Mixed Block/Event-Related



How to analyze task connectivity given task structure

- The more structured the task, the less you can use the time series (and viceversa)
- With **block** and with (not too fast) **event related** design we use the **general linear model GLM** to abstract from the time series into “**activations**”

$$\begin{array}{l} y_1 = b_0 + b_1X_{11} + \dots \dots \dots + b_pX_{1p} + e_1 \\ y_2 = b_0 + b_1X_{21} + \dots \dots \dots + b_pX_{2p} + e_2 \\ y_3 = b_0 + b_1X_{31} + \dots \dots \dots + b_pX_{3p} + e_3 \\ \vdots \qquad \qquad \qquad \qquad \qquad \qquad \vdots \qquad \qquad \qquad \qquad \qquad \qquad \vdots \\ y_n = b_0 + b_1X_{n1} + \dots \dots \dots + b_pX_{np} + e_n \end{array}$$

$y =$ Observed
BOLD

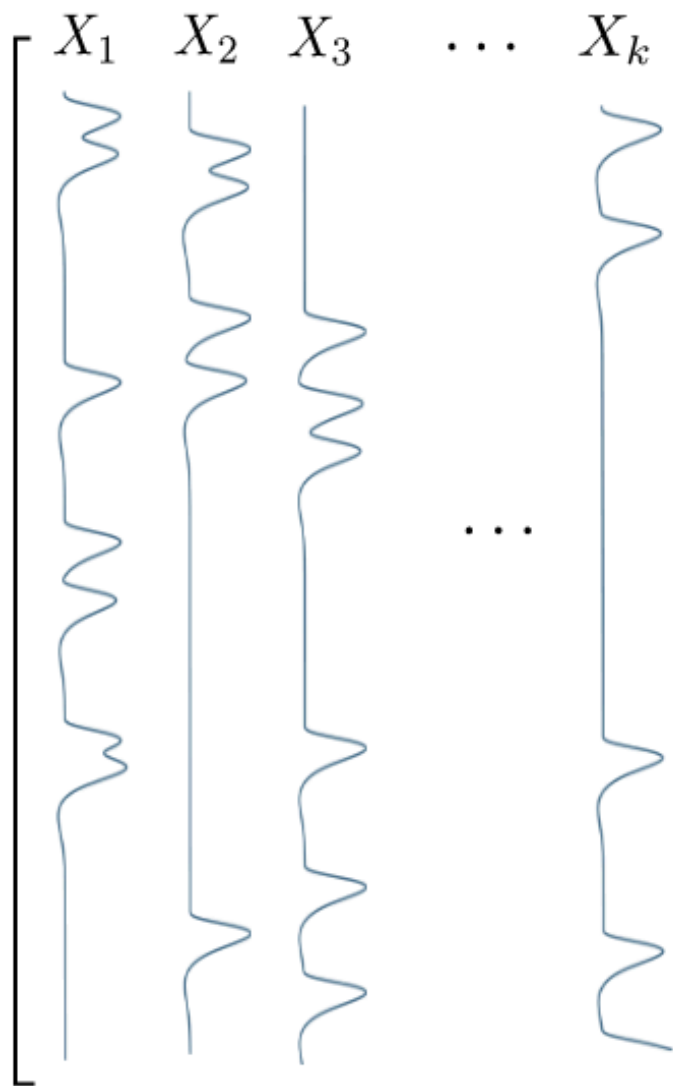
$X =$ Design Matrix

$\beta =$ Activation
coefficients

$\epsilon =$ Noise



=



...



+

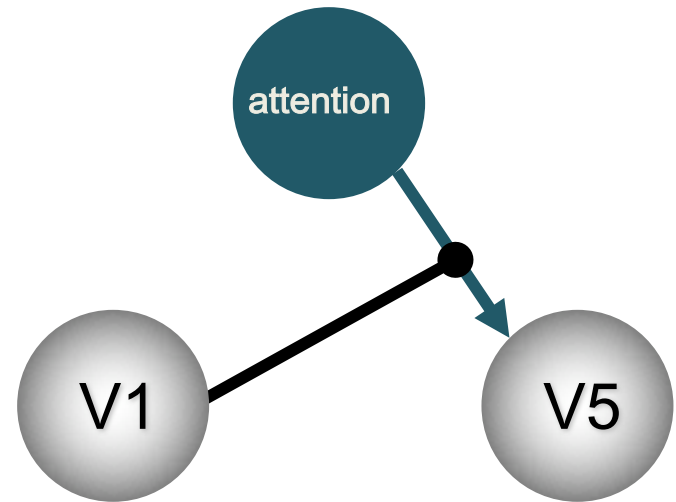
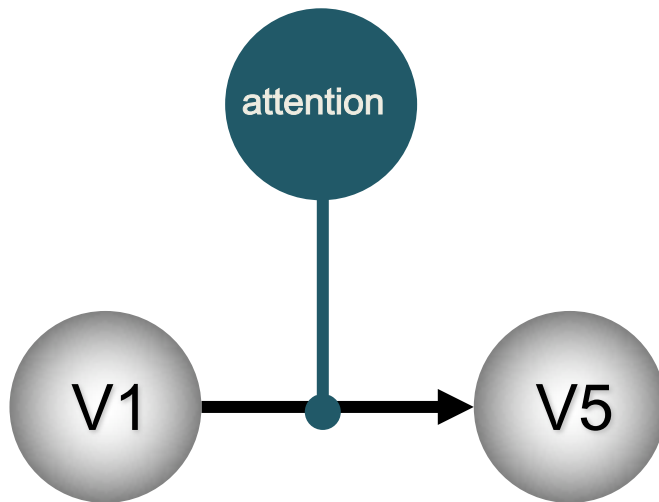


How to analyze task connectivity given task structure

- **With 20s blocks, the best is PPI***

$$Y = (\text{Att-NoAtt}) \beta_1 + \underline{\text{V1}} \beta_2 + (\text{Att-NoAtt}) * \text{V1} \beta_3 + e$$

- **Modeling** signal **Y**, given **task**, given **another signal** (V1), given an **interaction between task and signal**



Source: http://www.fil.ion.ucl.ac.uk/mfd_archive/2011/page1/mfd2011_connectivity_PPI_SEM.pptx

*PPI = psychophysiological interaction

How to analyze task connectivity given task structure

- **Resources for PPI**

- SPM (matlab)
- FSL (stand alone)
- gPPI (generalized PPI,
<https://www.nitrc.org/projects/gppi>)

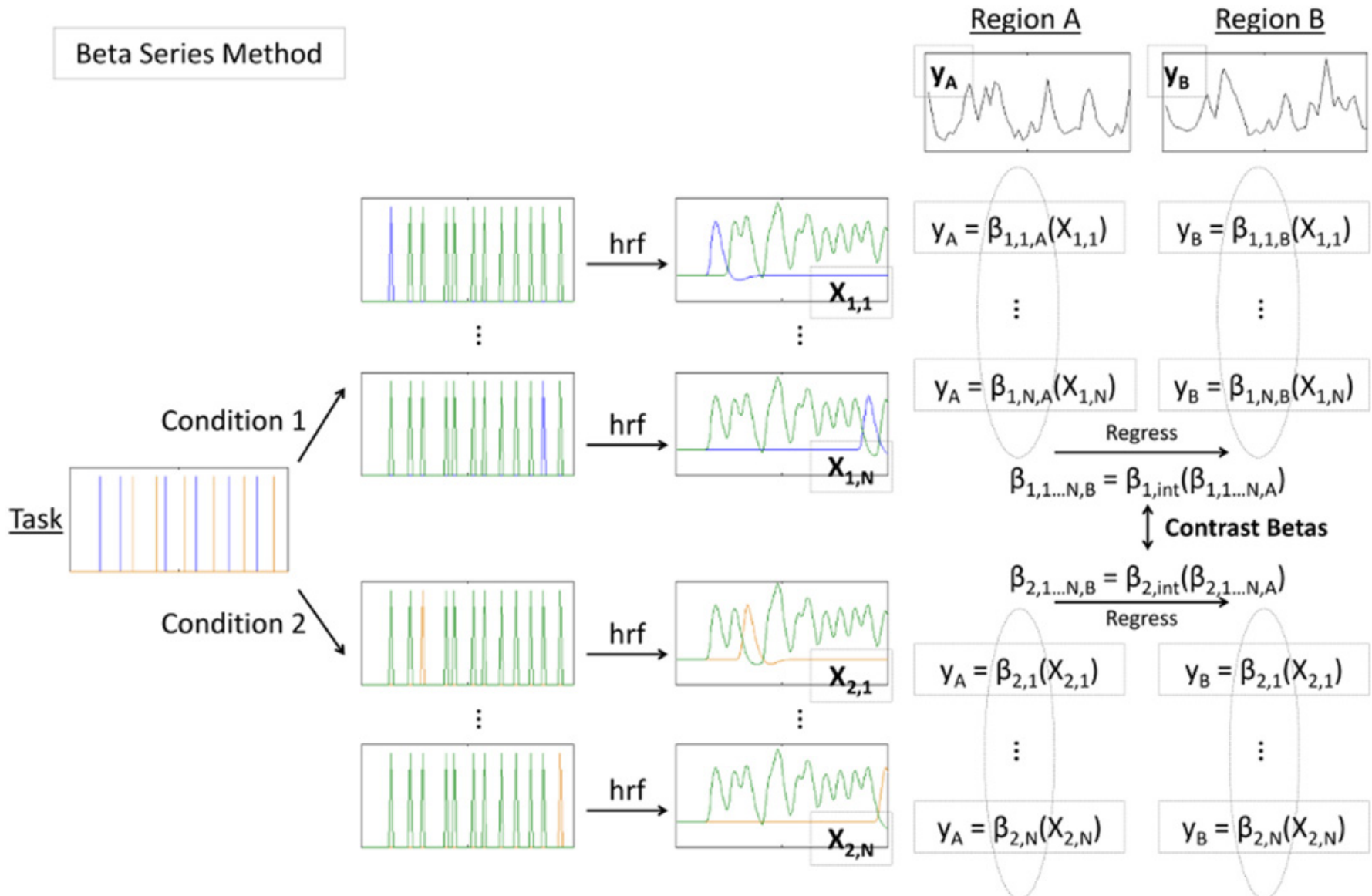
*PPI = psychophysiological interaction

How to analyze task connectivity given task structure

- **Event related, the best is beta series**
- **For every event we compute a beta weight in the GLM sense**
- **We replace BOLD time series with beta time series**
- **We correlate beta time series between regions**

How to analyze task connectivity given task structure

Beta Series Method



How to analyze task connectivity given task structure

- **Resources for beta series**
 - <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4019671/>
 - **BASCO toolbox:**
<https://www.nitrc.org/projects/basco/>
 - Mini function I made:
https://version.aalto.fi/gitlab/BML/bramila/blob/master/bramila_betaserie.m

Task

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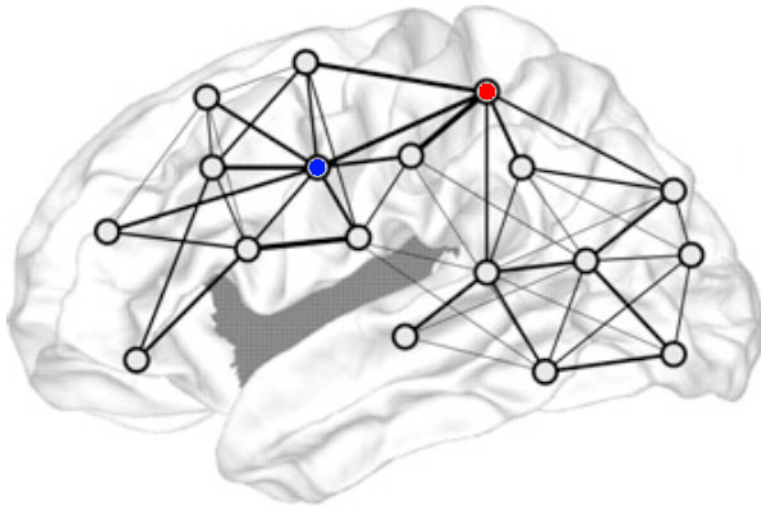
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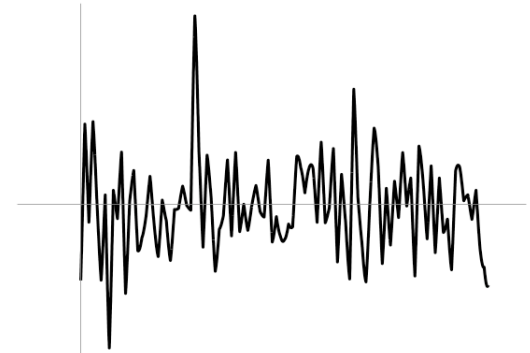
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Correlation approaches

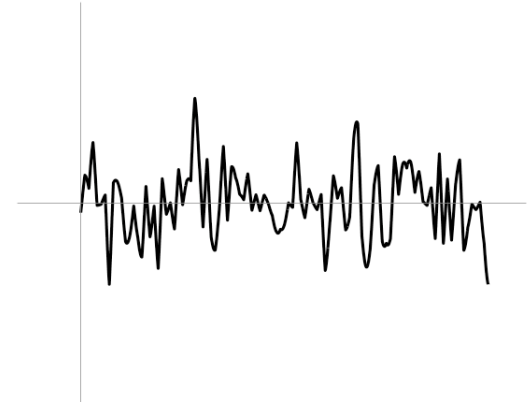
- Let's consider two time series for two voxels



$b_1(t)$

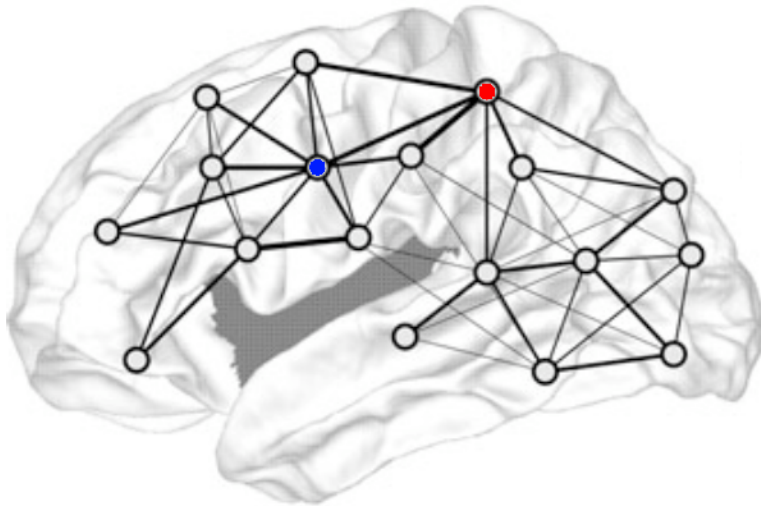


$b_2(t)$

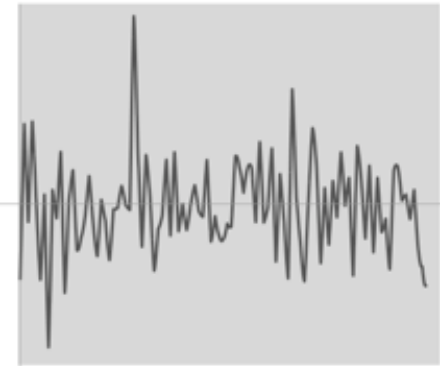


Correlation approaches

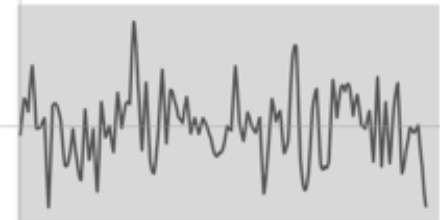
- Let's take all time points



$b_1(t)$

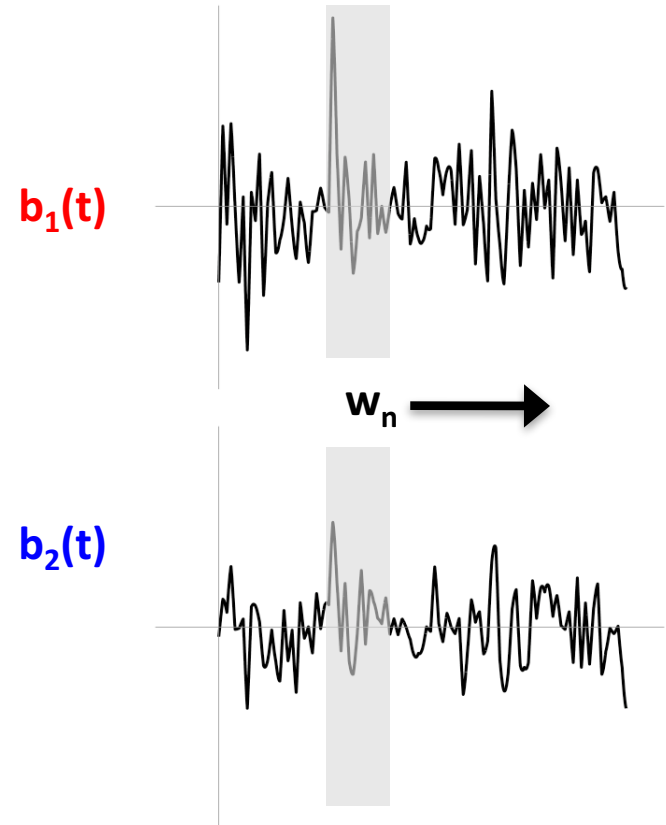
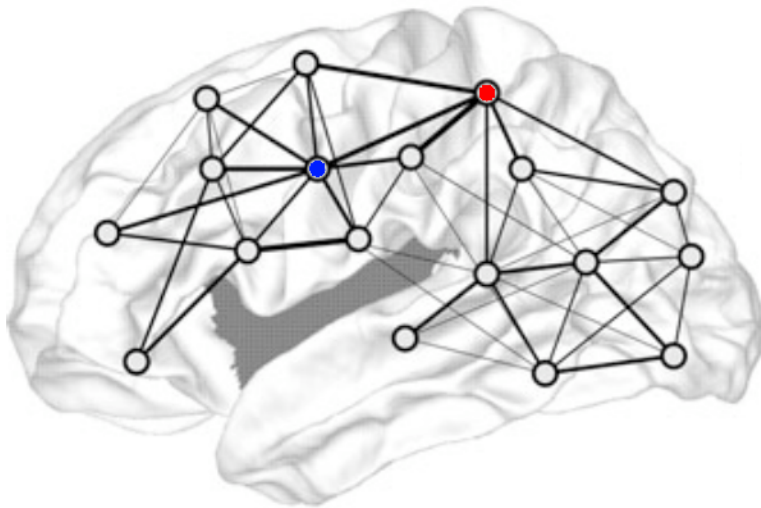


$b_2(t)$



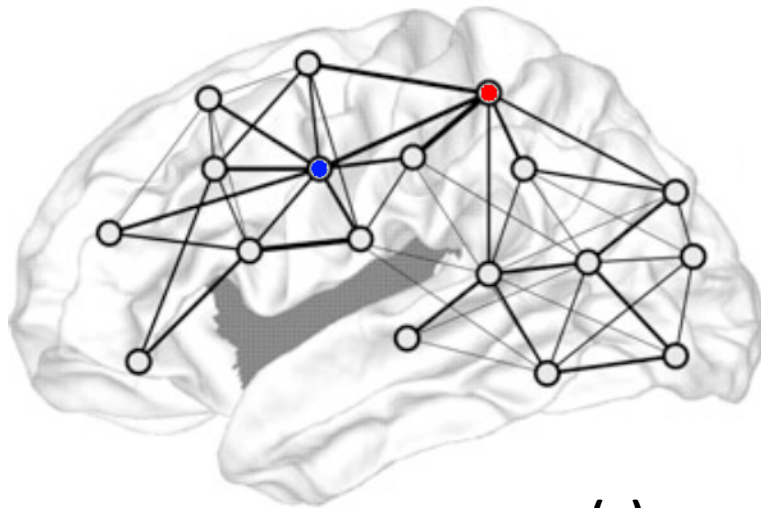
Functional connectivity in time

- Sliding window correlation

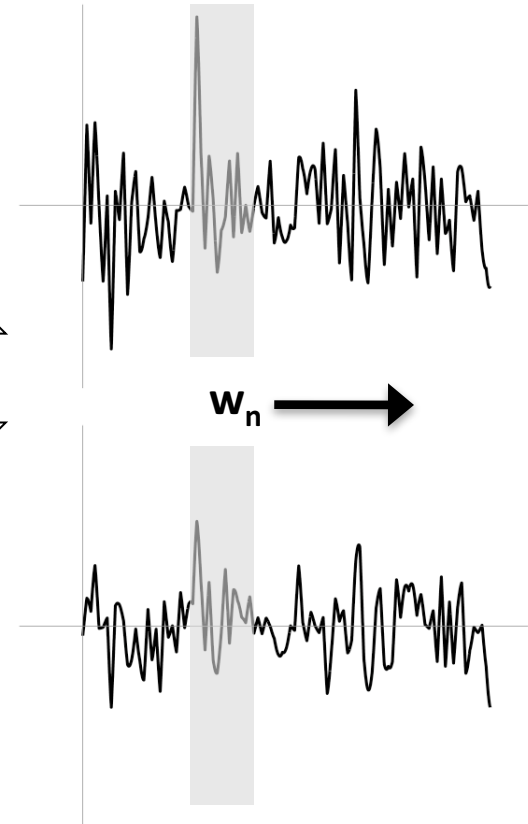
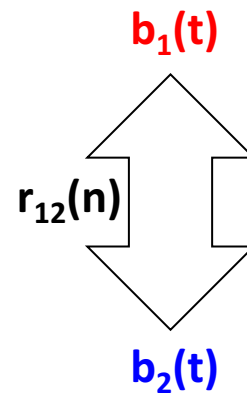


Functional connectivity in time

- Sliding window correlation for functional connectivity produces link time-series



$$e.g. r_{12}(n) = \text{corr}(b_1(w_n), b_2(w_n))$$

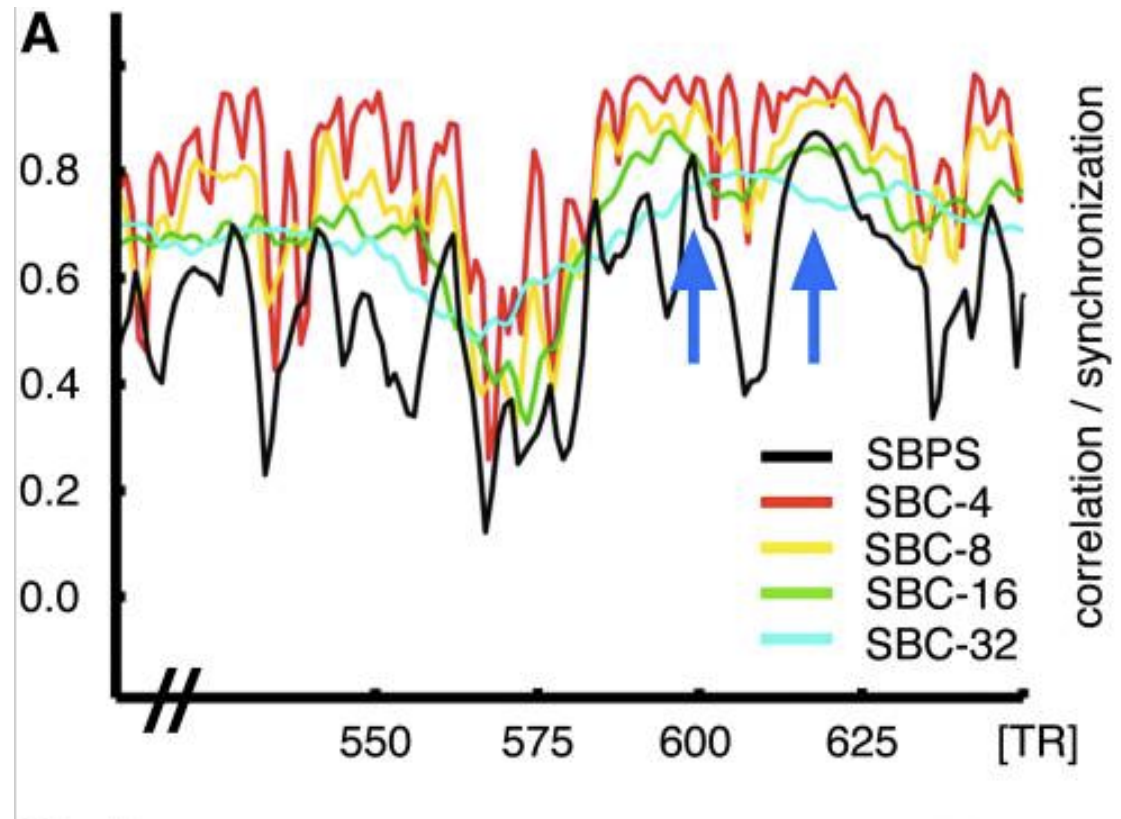


Problems with sliding window connectivity

- Field is still arguing what **Dynamic Functional Connectivity** means
- **Size of window depends on the temporal frequencies of the signal**
 - <http://www.sciencedirect.com/science/article/pii/S1053811914007496>
 - <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4758830/>

Functional connectivity in time: other approaches

- **Wavelet decomposition** <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2827259/>
- **Multiplication of derivatives** <http://www.sciencedirect.com/science/article/pii/S1053811915006849>
- **Phase synchronisation (Glerean et al 2012)** <https://www.ncbi.nlm.nih.gov/pubmed/22559794>



**How do we
compare networks
at the link level?**

Network statistics

- We have computed **links**, so you can think that links are what the voxels were in usual statistical parametric mapping and apply the same logic
- We have a **multiple comparisons problem** as we run as many test as many links (10^3 - 10^6)
- If links are **correlations** (i.e. in a range -1,1) then they are usually **z-transformed (atanh)** so they become more gaussianly distributed
- **The best way is to use permutation based approaches**

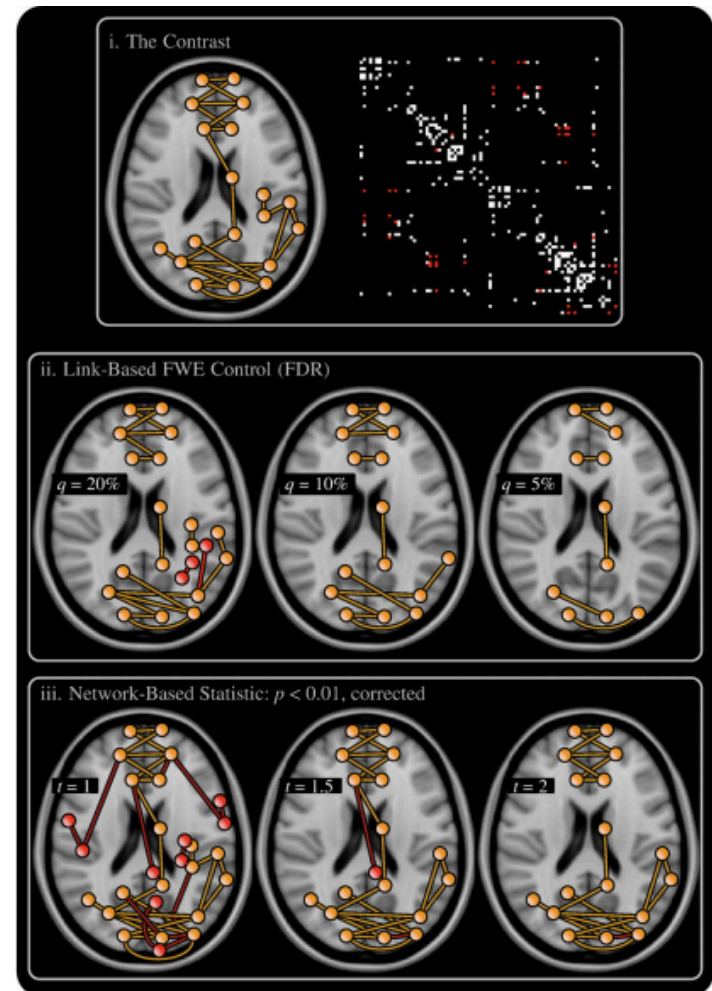
Solving the multiple comparison problem for networks

Network based statistics is the cluster correction applied to the links

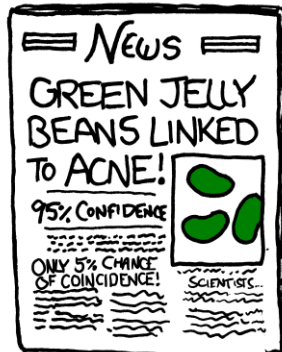
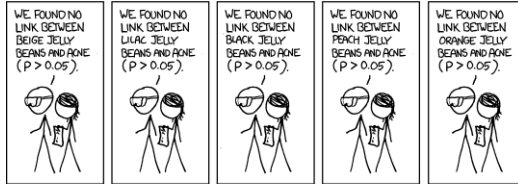
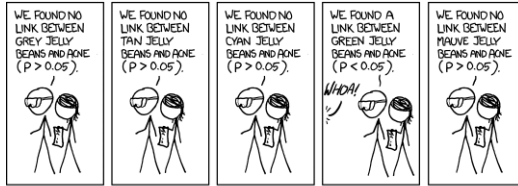
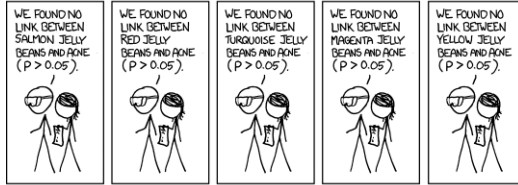
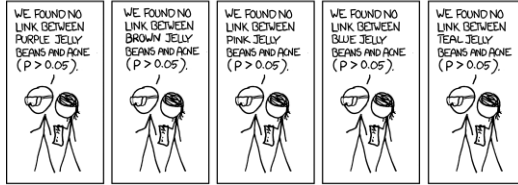
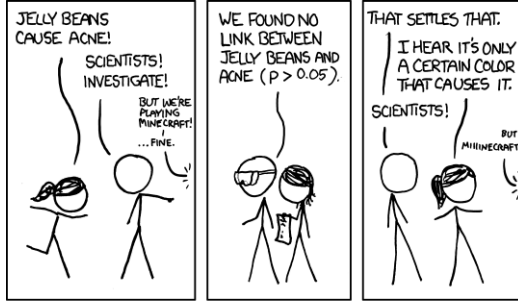
Check

<http://www.sciencedirect.com/science/article/pii/S10538119120008>

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Understanding the multiple comparison problem



Controlling for multiple comparisons – frequentist approach

- **Family of methods also called “Family wise error rate” control**
- **Classic example: Bonferroni correction. $\alpha = 0.05/NC$**
- **Ok, for smallish NC, but it’s not going to work with networks (264 nodes, 34716 links $\rightarrow \alpha \approx 10^{-6}$)**

Controlling for multiple comparisons – better approaches

- **False Discovery Rate (FDR)**
- **Based on distribution of p-values**
- **Procedure:**
 - Smallest p value $< \alpha/NC$
 - Second smallest p value $< \alpha/(NC-1)$
 - Third smallest p value $< \alpha/(NC-2)$
 - Etc etc...

Book by Efron “Large-scale statistics”

Controlling for multiple comparisons – using permutations

- **Do permutation simultaneously for all multiple variables (e.g. all links) to generate at once many surrogate values**
- **Pick the strongest (max statistics)**
- **The null distribution will look more skewed towards the maximum**

Controlling for multiple comparisons – using permutations

- **Cluster approaches (with fMRI)**
- **Non-parametric:**
 - At each permutation set a cluster forming threshold
 - Count how many voxels in the largest connected cluster
 - Compare number of connected voxels in the un-permuted cluster

🏠 > Current Issue > vol. 113 no. 28 > Anders Eklund, 7900–7905, doi: 10.1073/pnas.1602413113

 Check for updates

Cluster failure: Why fMRI inferences for spatial extent have inflated false-positive rates

Anders Eklund^{a,b,c,1}, Thomas E. Nichols^{d,e}, and Hans Knutsson^{a,c}

Author Affiliations 


Edited by Emery N. Brown, Massachusetts General Hospital, Boston, MA, and approved May 17, 2016 (received for review February 12, 2016)

This Issue



July 12, 2016
vol. 113 no. 28
[Masthead \(PDF\)](#)
[Table of Contents](#)

 PREVIOUS ARTICLE

NEXT ARTICLE 

Don't Miss

<http://www.pnas.org/content/113/28/7900.full>

According to many of the headlines that greeted “Cluster failure”, the paper is a devastating bombshell that could demolish the whole field of functional magnetic resonance imaging (fMRI):

Bug in fMRI software calls 15 years of research into question (Wired)

A bug in fMRI software could invalidate 15 years of brain research. This is huge. (ScienceAlert)

New Research Suggests That Tens Of Thousands Of fMRI Brain Studies May Be Flawed (Motherboard)

<http://blogs.discovermagazine.com/neuroskeptic/2016/07/07/false-positive-fmri-mainstream/#.WRq28SN97-m>

Controlling for multiple comparisons – using permutations

- **Cluster approach (with fMRI) recently re-tested**
- **Fake task using resting state data**
- **Comparing cluster approaches: parametric (RFT) and non parametric**
- **Permutations was the only one closest to the “truth”**

**How about
networks?**

How to compute differences between networks

- **At each level (node/link/global) you can test for a difference between two groups or from a baseline prior knowledge**
- **Links are correlations -> they can be mapped to p-values but degrees of freedom must be estimated**

How to compute differences between networks

- **Node properties are coming from very long tailed/weirdly shaped distributions -> permutation approaches or build null models with networks**
- **Network null models have problems (suboptimal)**

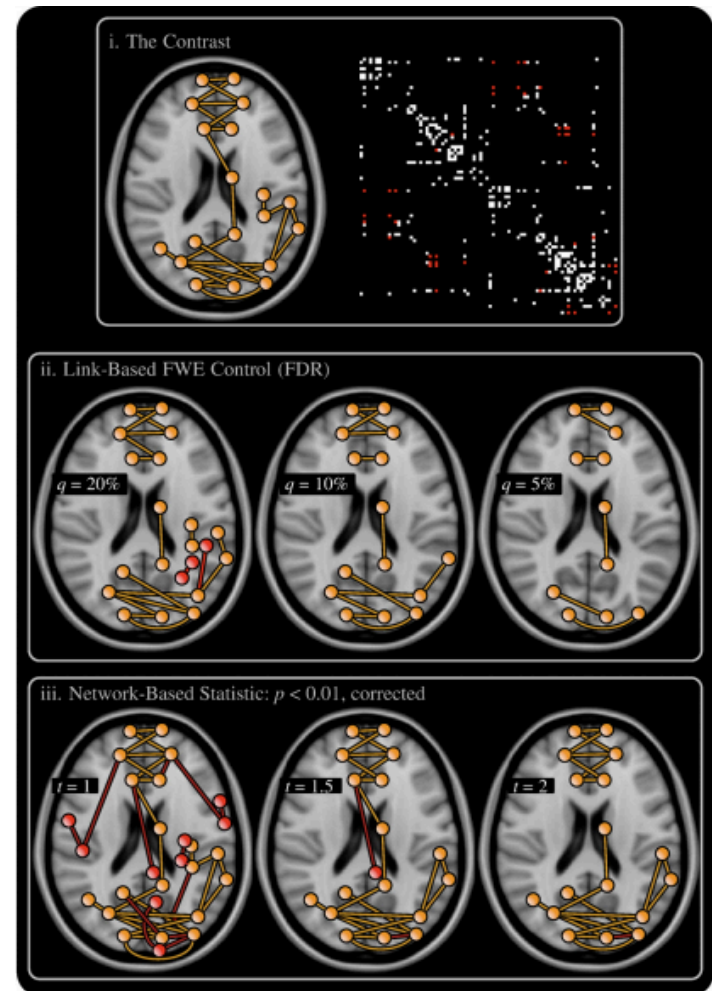
Solving the multiple comparison problem for networks

Network based statistics is the cluster correction applied to the links

Check

<http://www.sciencedirect.com/science/article/pii/S10538119120008>

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PART 2

Brain network properties

Network topology

NETWORK LEVEL FEATURES

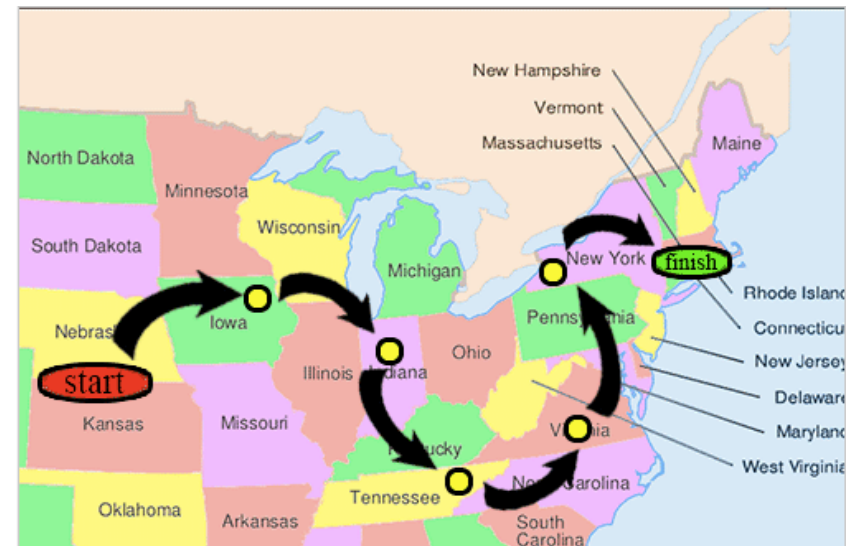
What?

WHAT IS A SMALL WORLD NETWORK?

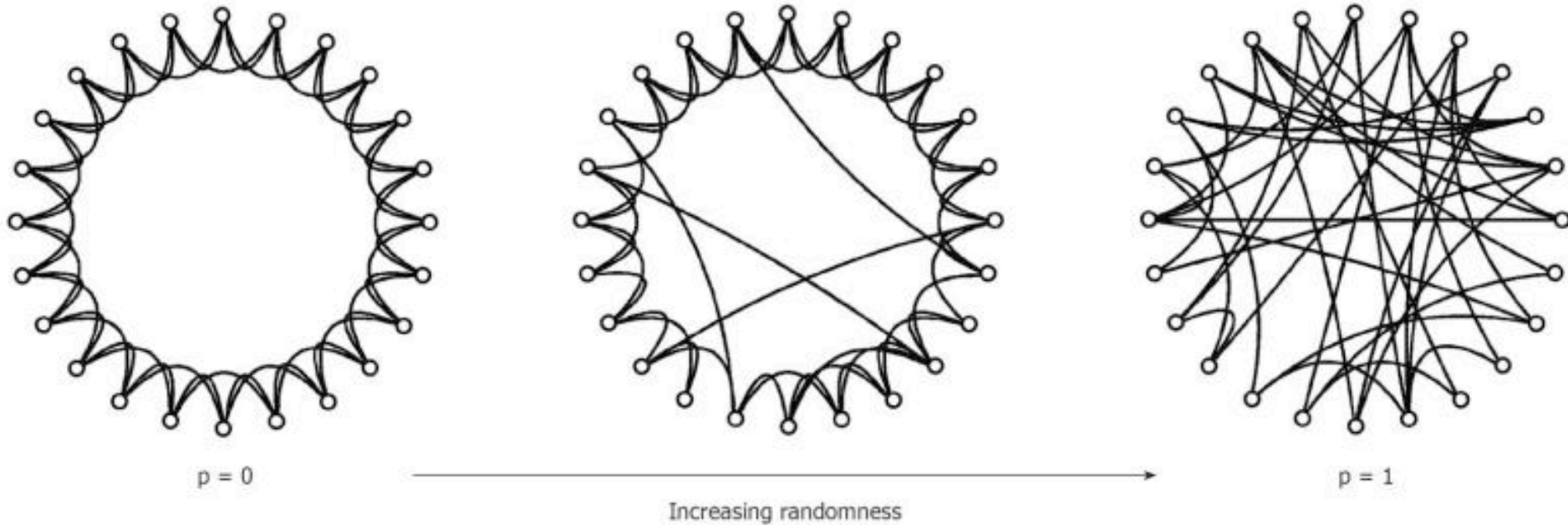
The small world experiment

Stanley Milgram (1969)

- Try to send a letter to Boston through a **chain of people** by only forward it to a friend who might know the final recipient
- **Six degrees of separation**
i.e. an *average path* of 6 links in the network



Small world networks



Watts, D. J., & Strogatz, S. H. (1998). Collective dynamics of “small-world” networks. *Nature*, 393(6684), 440–2. doi:10.1038/30918

Small world networks

Small world networks are present in biological system as an efficient way to keep the average path low and limit connection cost.

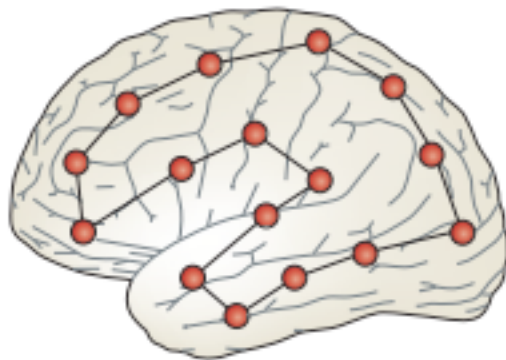
The brain is a small world network.

Why?

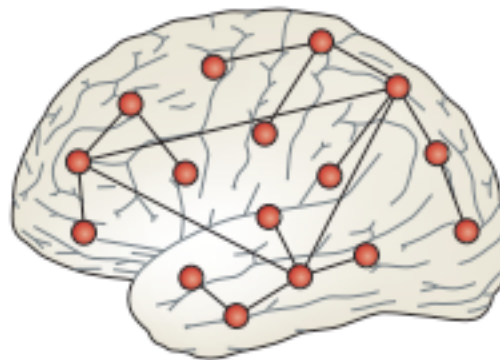
WHY IS THE BRAIN A SMALL WORLD NETWORK?

The small-world configuration is the optimal to optimize communication cost and efficiency

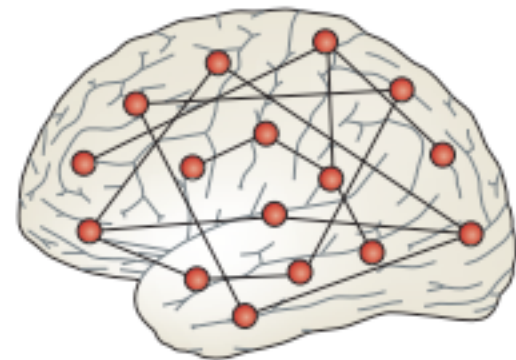
Lattice topology



Complex topology



Random topology



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

Small world topology implies segregation and integration

- **Small world topology implies high clustering:**

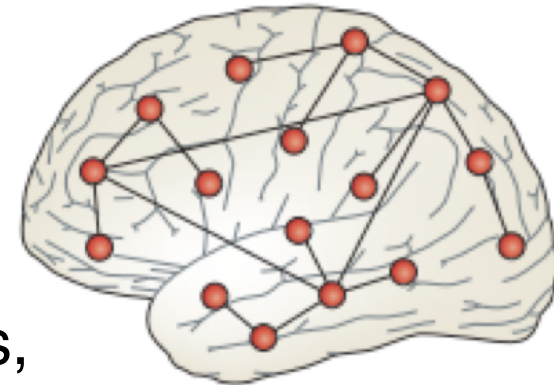
within a region we have more connections, regions are specialized (e.g. visual cortex, auditory cortex)

- **Small world topology implies short path:**

densely connected regions are joined together by long-range links

- **Clustering -> Segregation**

- **Short path -> Integration**



Network topology

NODE LEVEL FEATURES

What?

WHAT IS A HUB?

What is a hub?

A hub is the effective center of an activity, region, or network...

i.e. an important node in the network

What is a hub?

A hub is the effective center of an activity, region, or network...

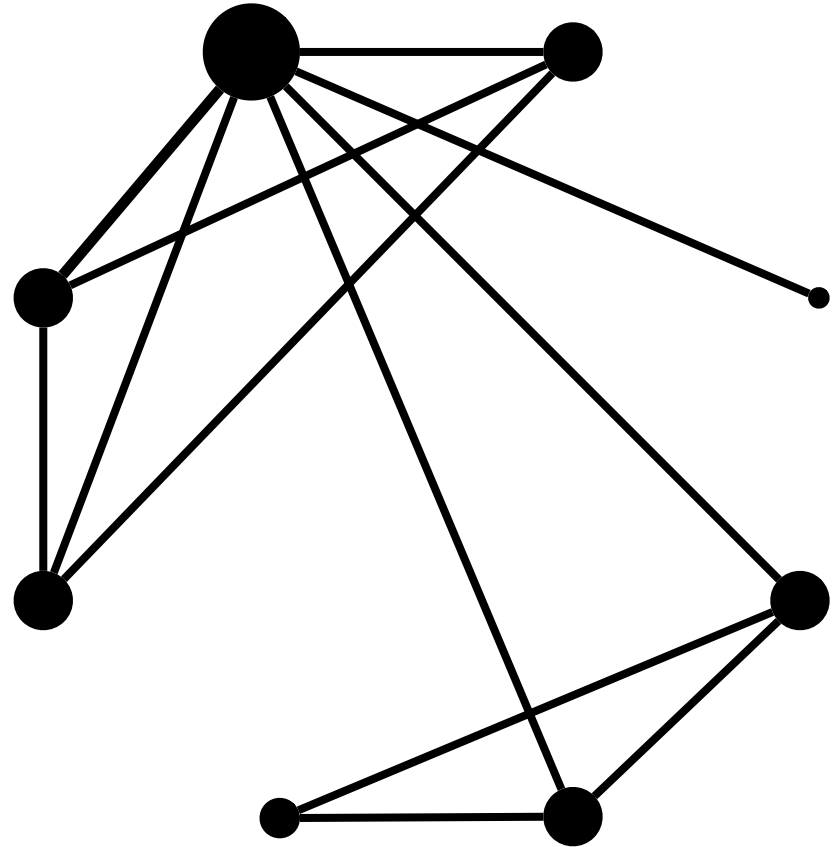
i.e. an important node in the network

How?

HOW CAN WE QUANTIFY A HUB?

Microscopic (node level) measures

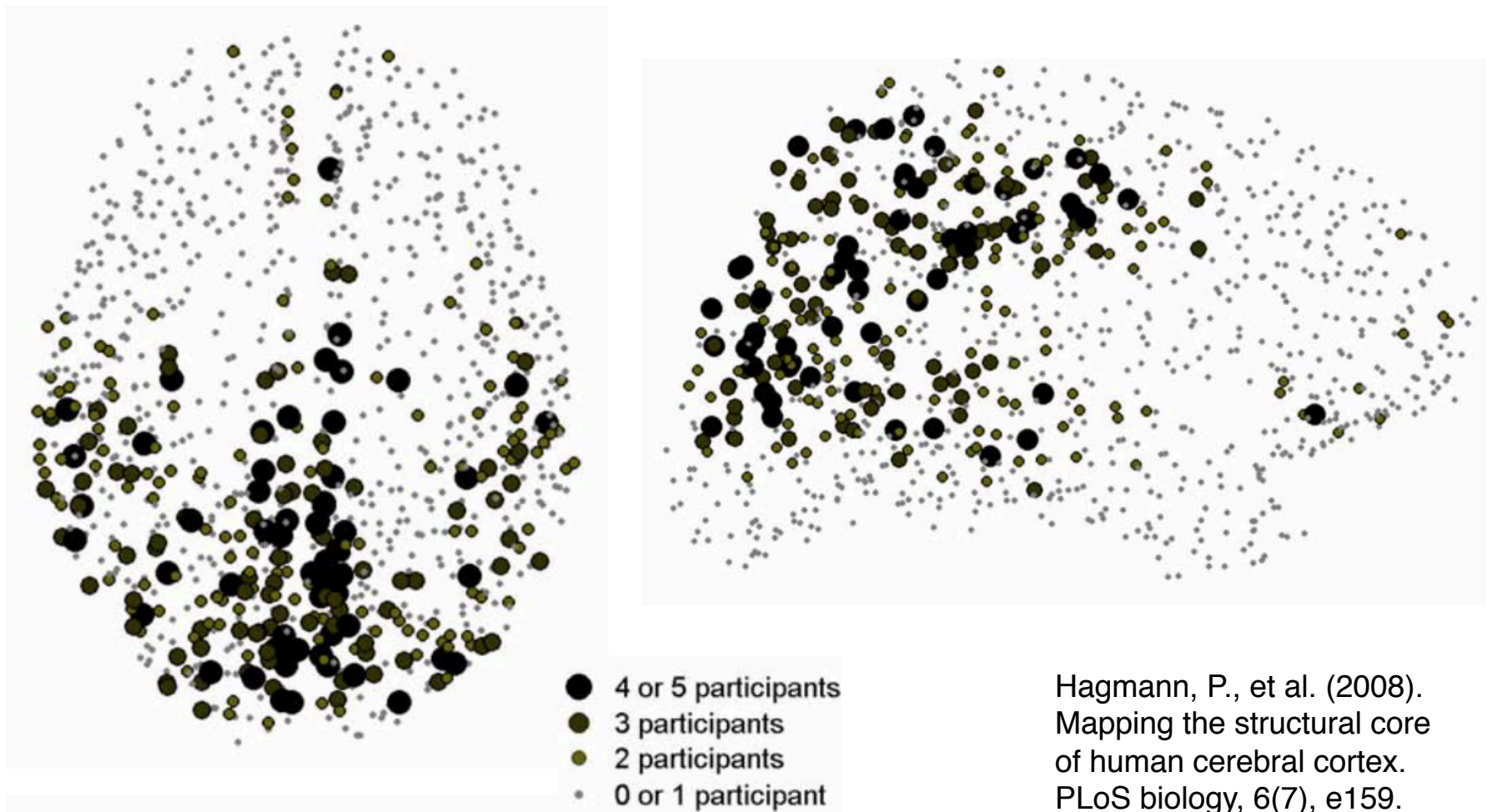
- **Node degree/strength**
How strong is a node?
- **Clustering**
How close is the node with the neighbours?
- **Closeness centrality**
How distant is the node?
- **Betweenness centrality**
How many shortest paths through the node?



What?

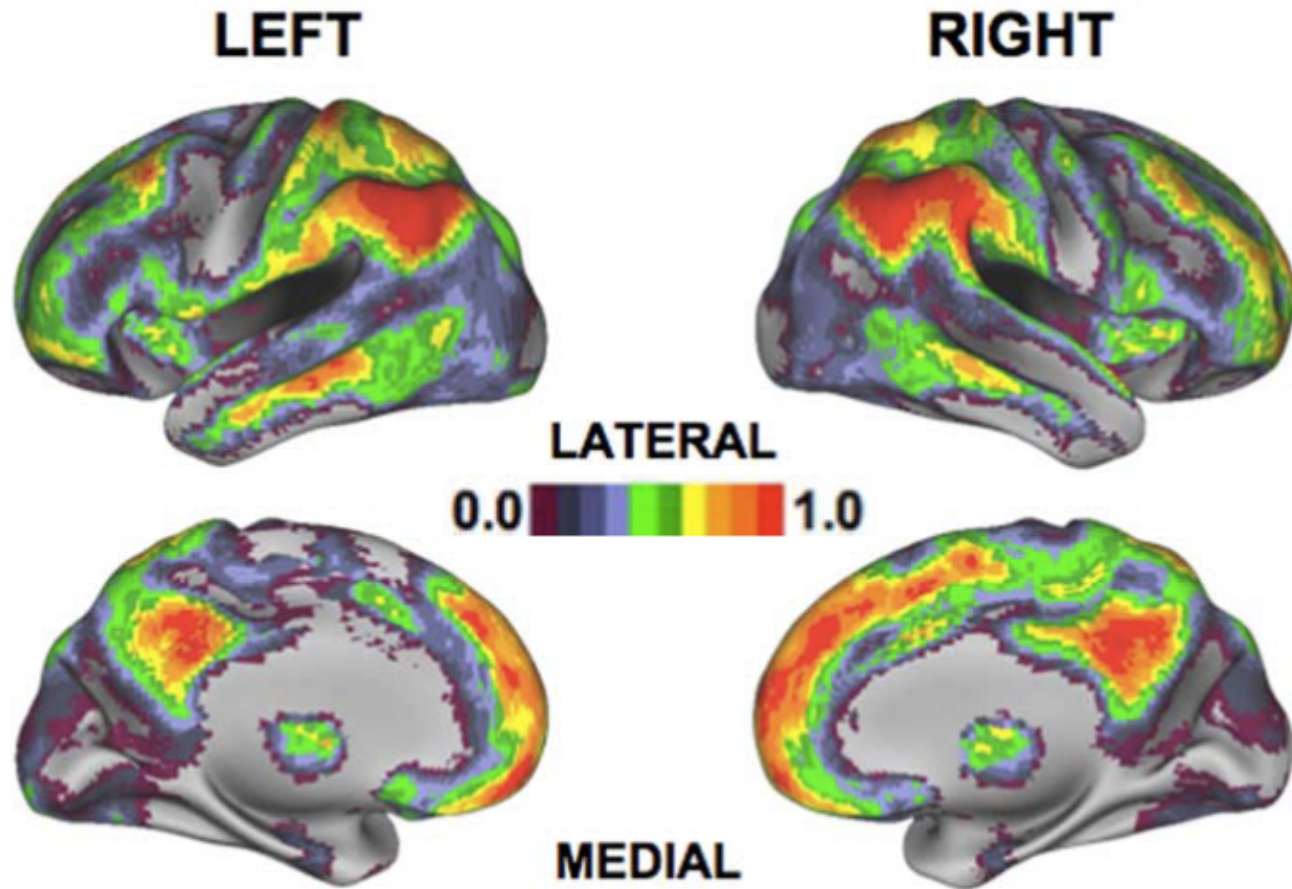
WHAT ARE THE HUBS IN THE BRAIN?

Cortical hubs in the human brain



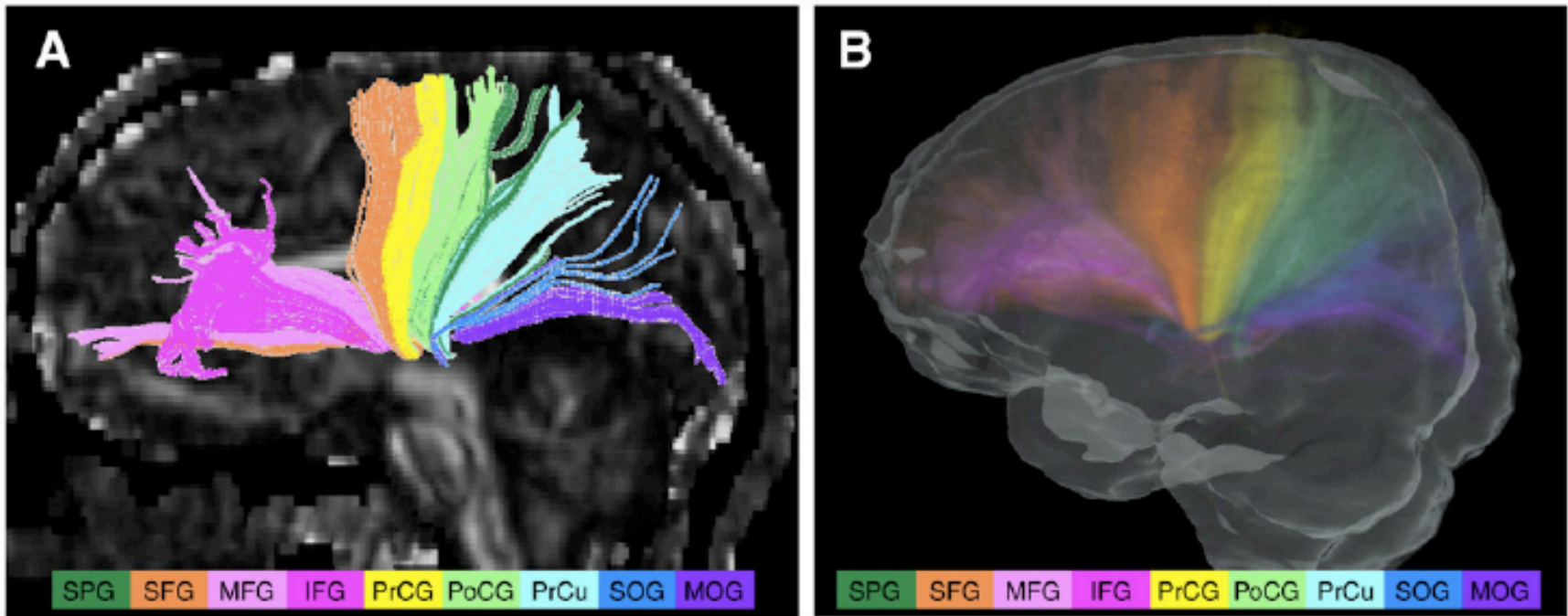
Hagmann, P., et al. (2008).
Mapping the structural core
of human cerebral cortex.
PLoS biology, 6(7), e159.

Cortical hubs in the human brain



Buckner, R. L., et al. (2009). Cortical hubs revealed by intrinsic functional connectivity. *The Journal of neuroscience* 29(6), 1860–73.

Sub-cortical hubs in the human brain: the thalamus

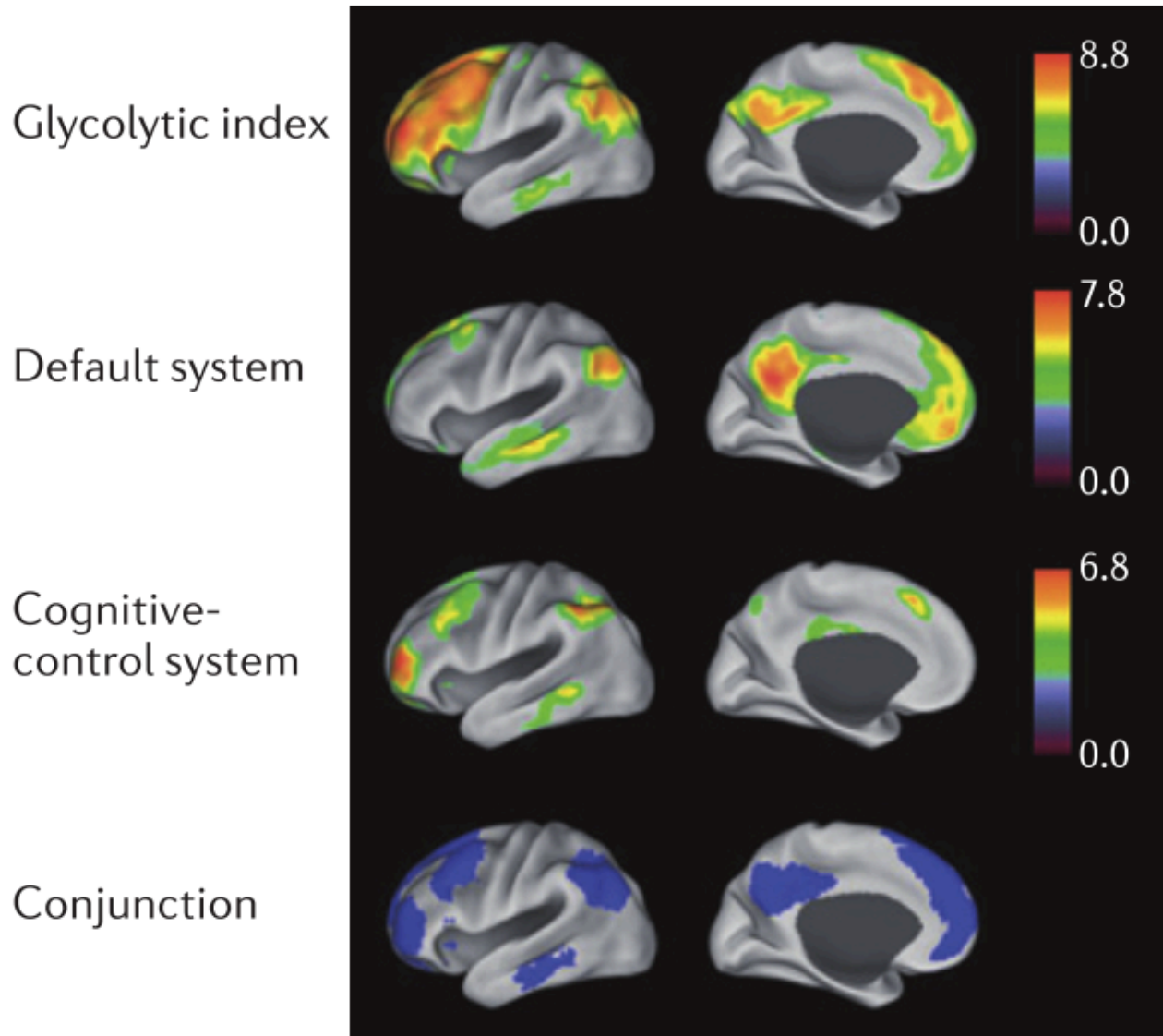


Zhang et al. (2010) Atlas-guided tract reconstruction for automated and comprehensive examination of the white matter anatomy. *Neuroimage*. 2010 Oct 1;52(4):1289-301.

What?

**WHAT IS THE RELATIONSHIP BETWEEN
HUBS AND BRAIN ACTIVITY?**

Energy consumption in the brain



The most important (central) hubs are those with higher glycolytic index, i.e. higher metabolic cost.

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

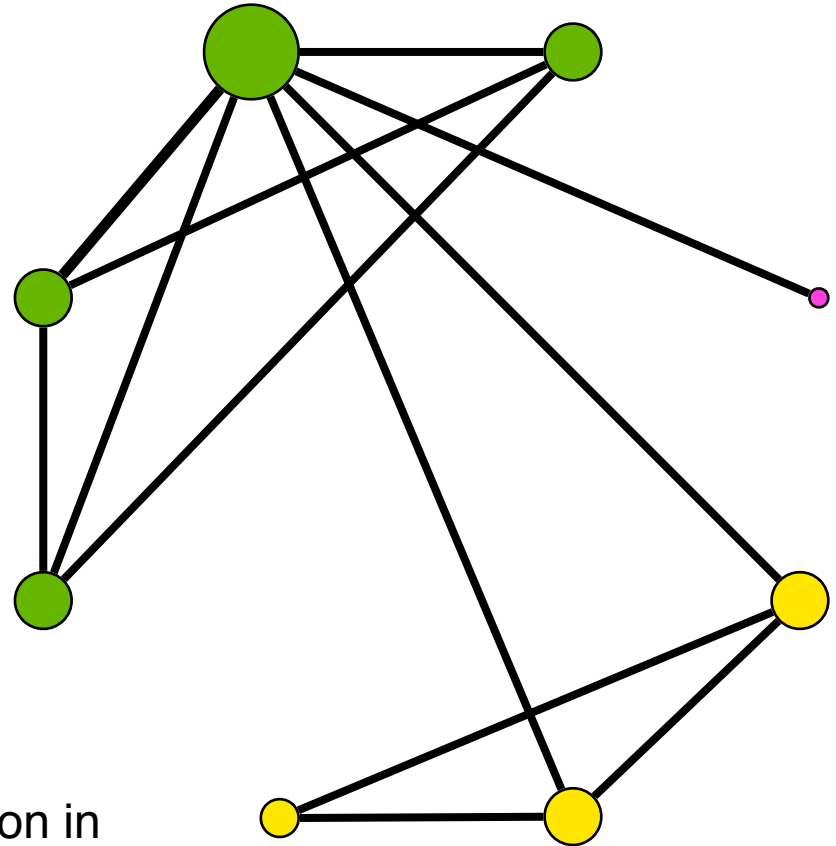
What?

WHAT IS A NETWORK MODULE?

Quantifying modules in networks

Communities/clusters

Finding subsets of nodes that are forming a module, i.e. they are more connected with each other than with other parts of the network



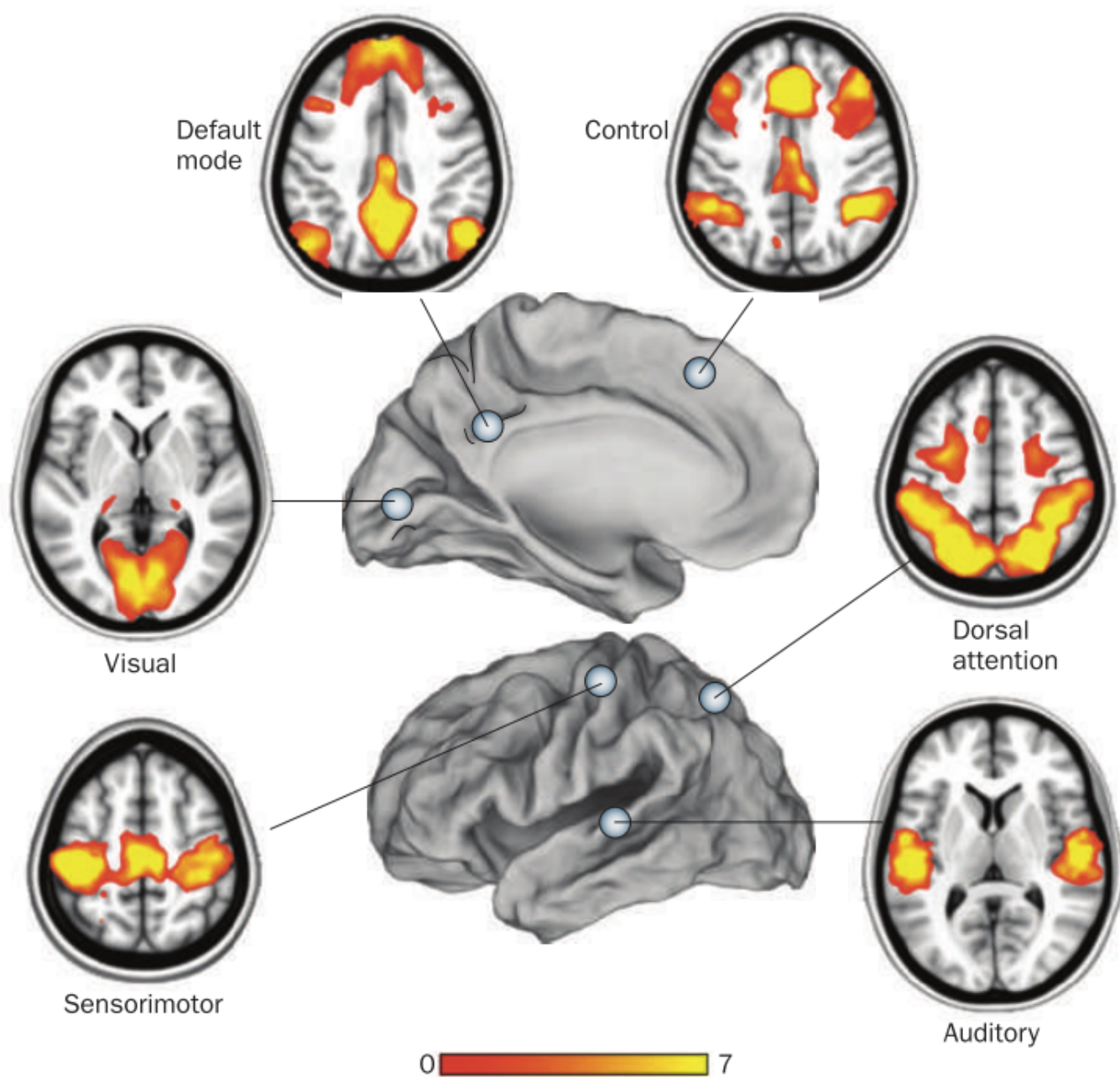
Fortunato, S. (2010). Community detection in graphs. *Physics Reports*, 486(3-5), 75–174

What?

WHAT ARE THE MODULES IN THE BRAIN?

The networks of the human brain

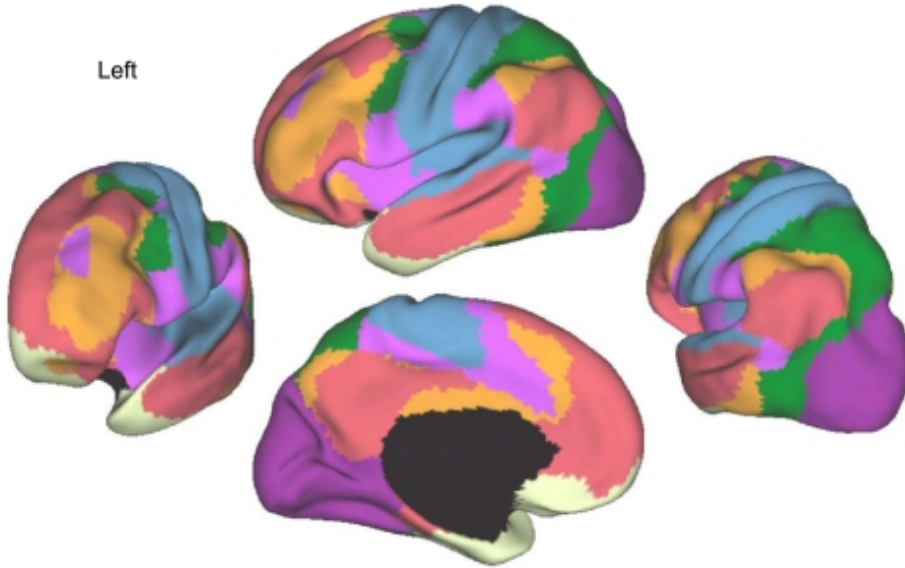
- We look at **which regions are more connected with each other (clustering)**
- We identify **~6 main modules** in the human cortex that corresponds to important cognitive functions
- They are often called “**networks**” although they are technically sub-networks



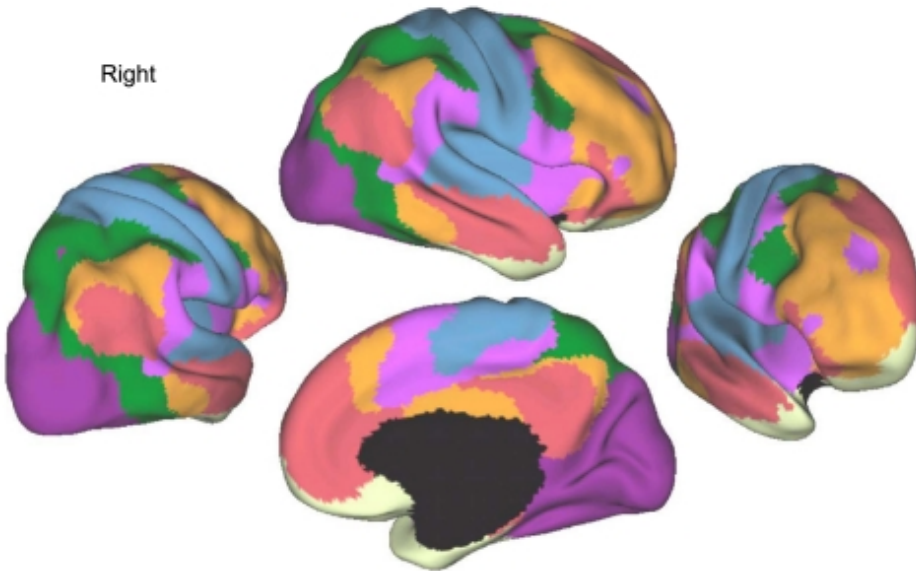
Zhang, D., & Raichle, M. E. (2010). Disease and the brain's dark energy. *Nature reviews. Neurology*, 6(1), 15–28.








7-Network Parcellation (N=1000)

Left



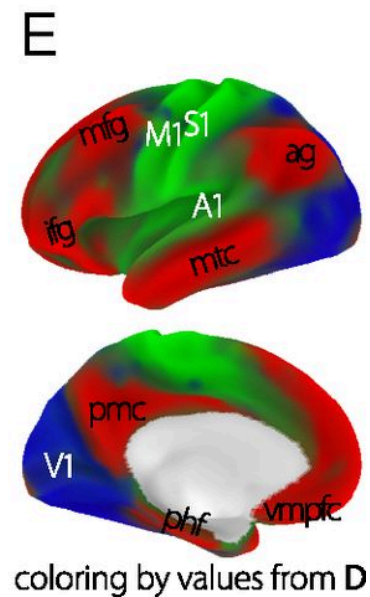
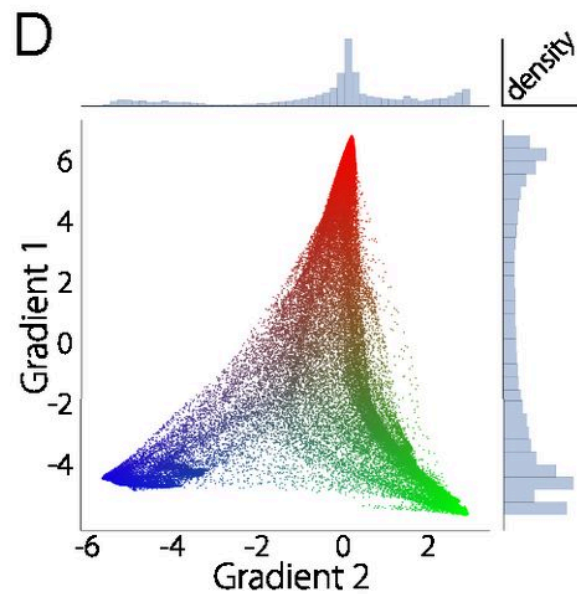
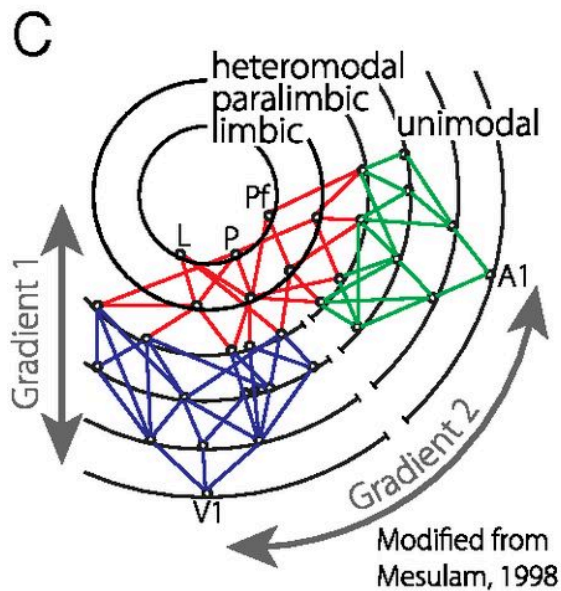
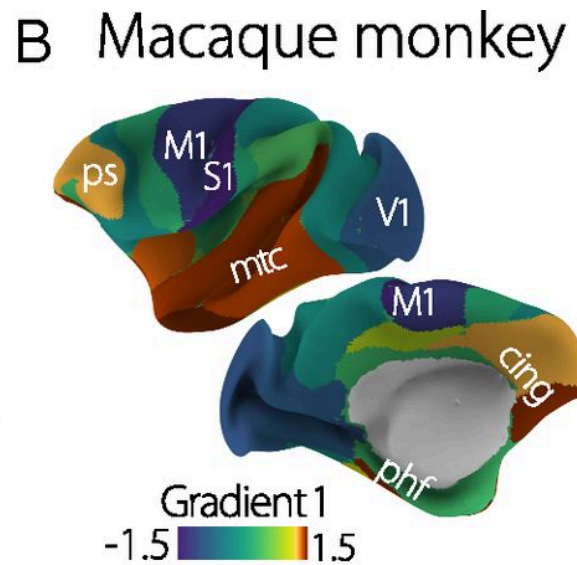
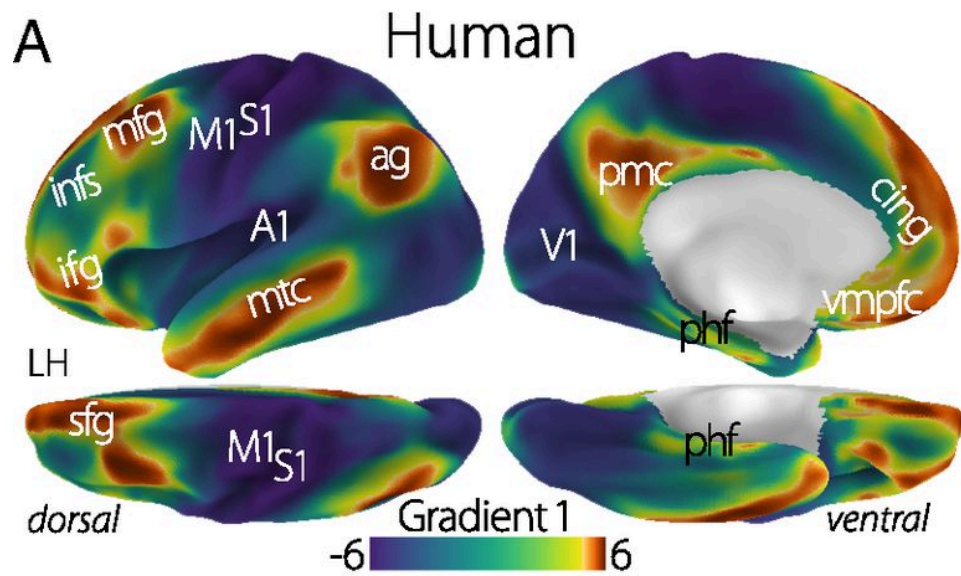
Right



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

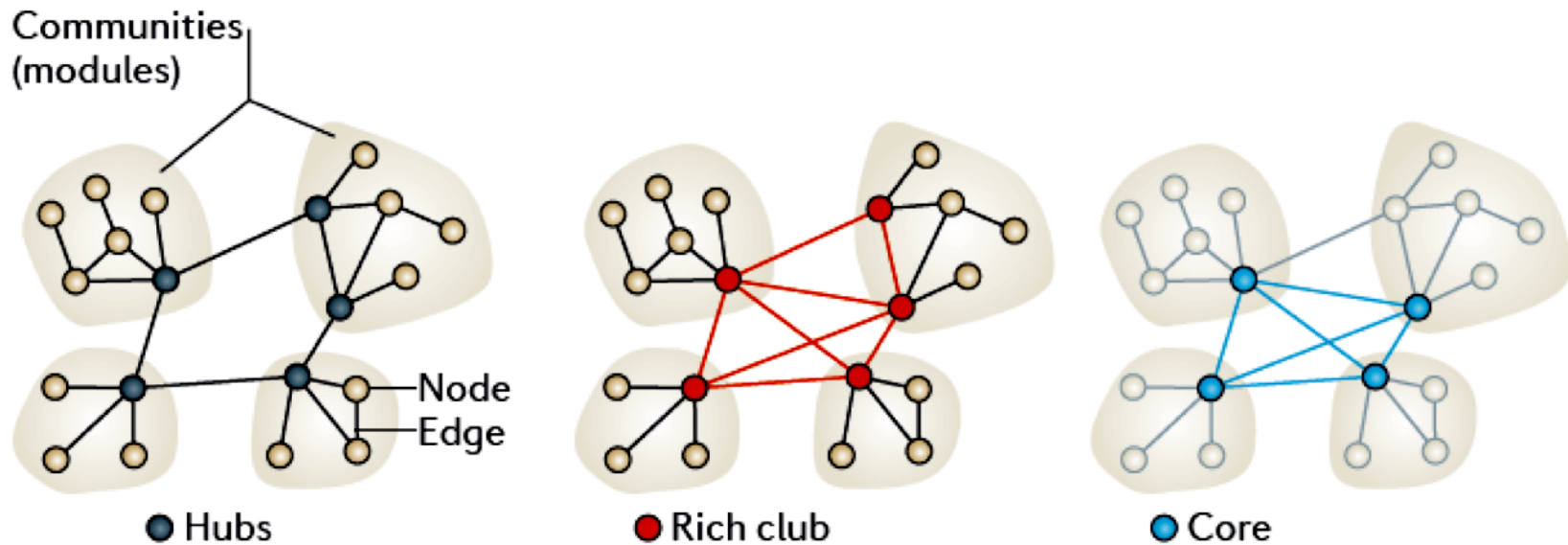
Yeo et al. (2011)

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



Margulies et al (2016) **Situating the default-mode network along a principal gradient of macroscale cortical organization.** PNAS

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



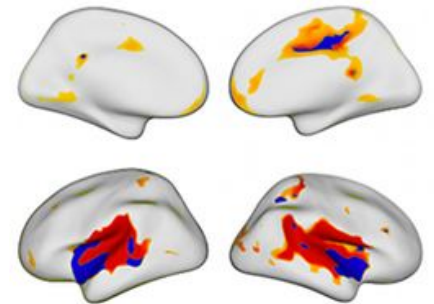
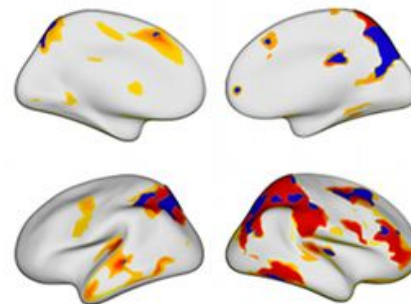
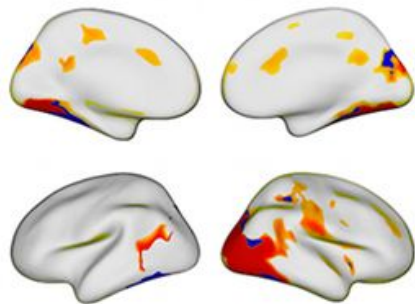
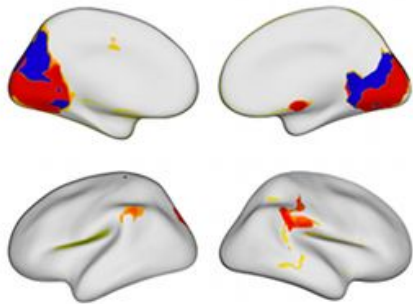
Bullmore, E., & Sporns, O. (2012). The economy of brain network organization. *Nature reviews. Neuroscience*, 13(5), 336–49

1. primary visual

2. extra striate visual

3. bilateral parietal

10. auditory

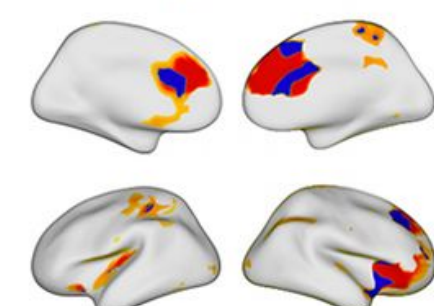
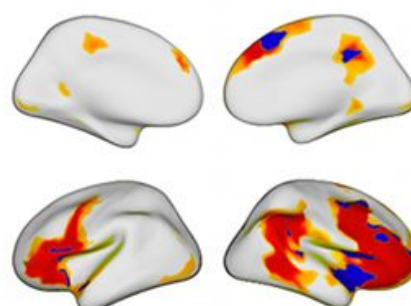
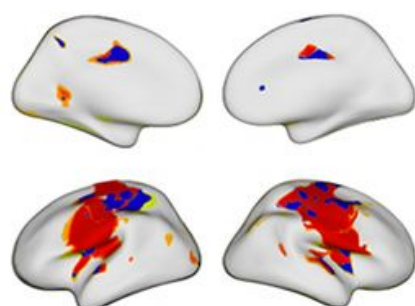
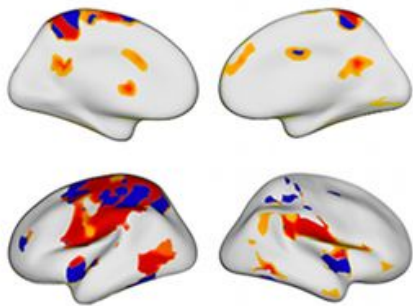


4. sensory

5. motor

6. right parietal frontal

11. frontal

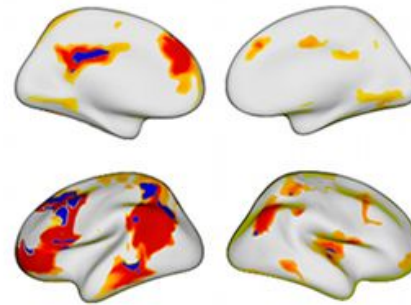
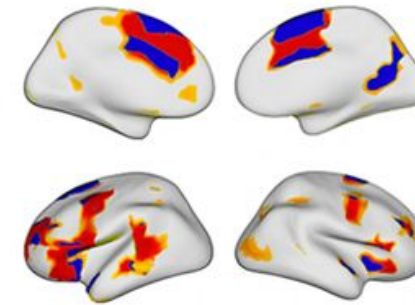
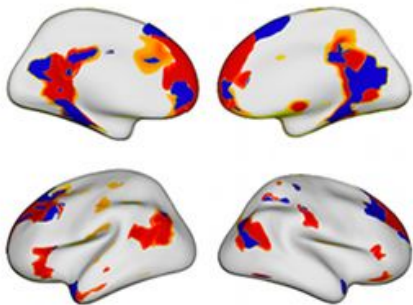


7. default mode

8. salience

9. left parietal frontal

Rich-club hubs (blue)



Modules (red)

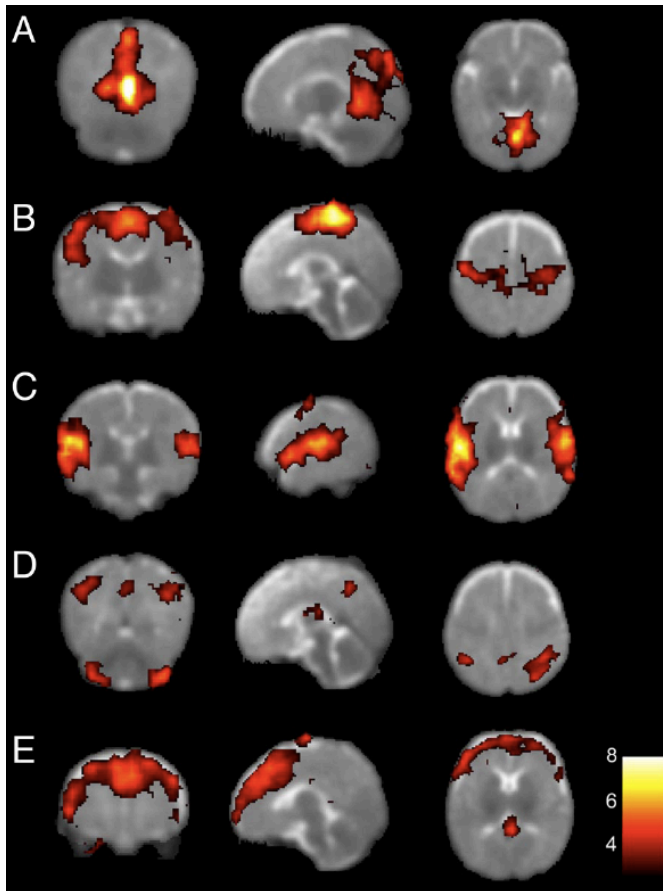
How?

HOW DOES CONNECTIVITY CHANGE IN
TIME?

Temporal scales of connectivity

- Changes across **(milli)seconds**
 - **Fast** functional changes due to **extrinsic** or **intrinsic** processes
- Changes across **years**
 - **Slow** structural changes due to **genetics, environment** and **noise**

Sub-network modules in the infant brain at rest with fMRI



Five consistent modules

A) primary visual

B) somatosensory/motor

C) primary auditory

D) Posterior lateral and midline of parietal cortex

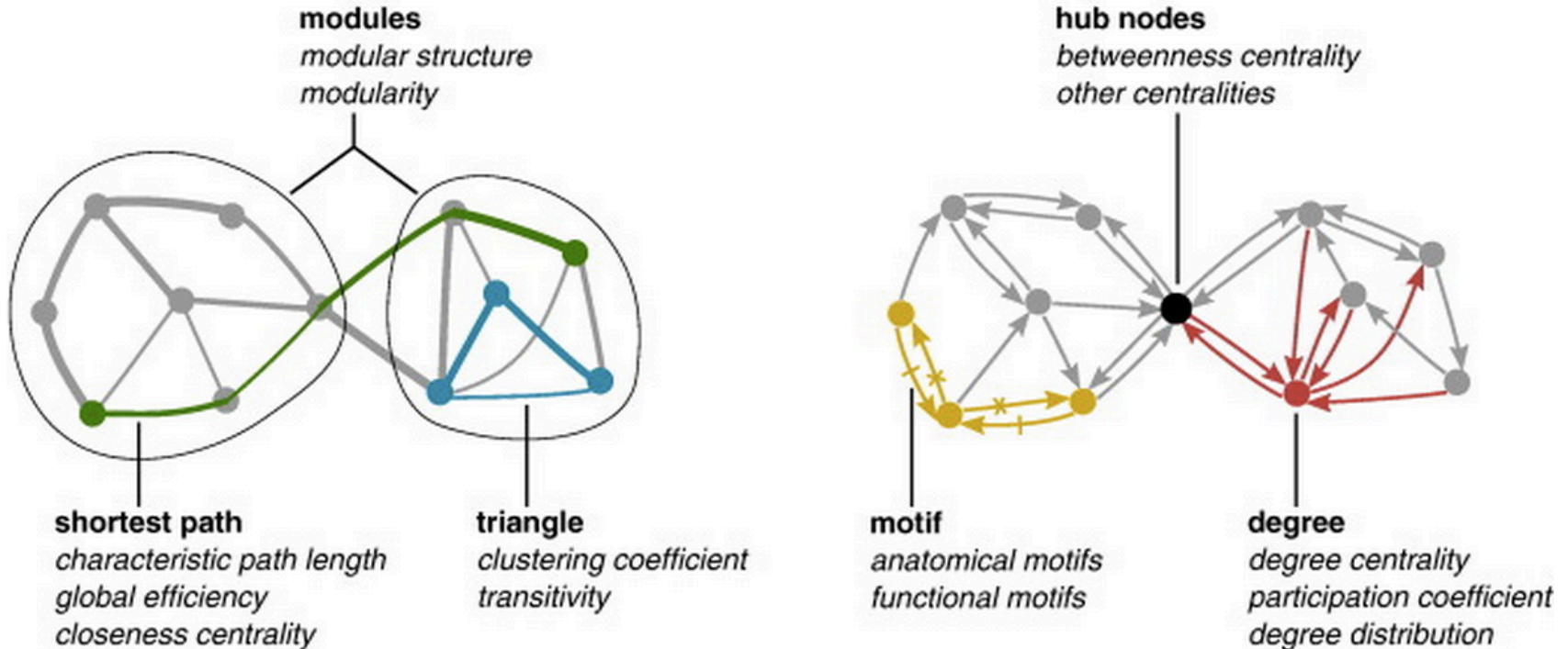
E) medial and lateral anterior frontal cortex

Fransson et al (2007) PNAS

How to estimate and compare network properties

How to calculate these network features?

- See Brain Connectivity Toolbox and its related papers



Rubinov & Sporns 2010, Neuroimage <http://www.neuroscience.cam.ac.uk/publications/download.php?id=17703>

Bullmore & Sporns 2009 Nature Review Neuroscience <http://www.nature.com/nrn/journal/v10/n3/full/nrn2575.html>

How to compare network properties?

- It's tricky because **network properties do not follow a gaussian distribution**
- Best is to NOT assume anything and use **permutation testing**: e.g. for a node, shuffle labels and compute surrogate group difference. Repeat x 5000 and get null distribution.
- **Remember to correct for multiple comparisons**

Part 3

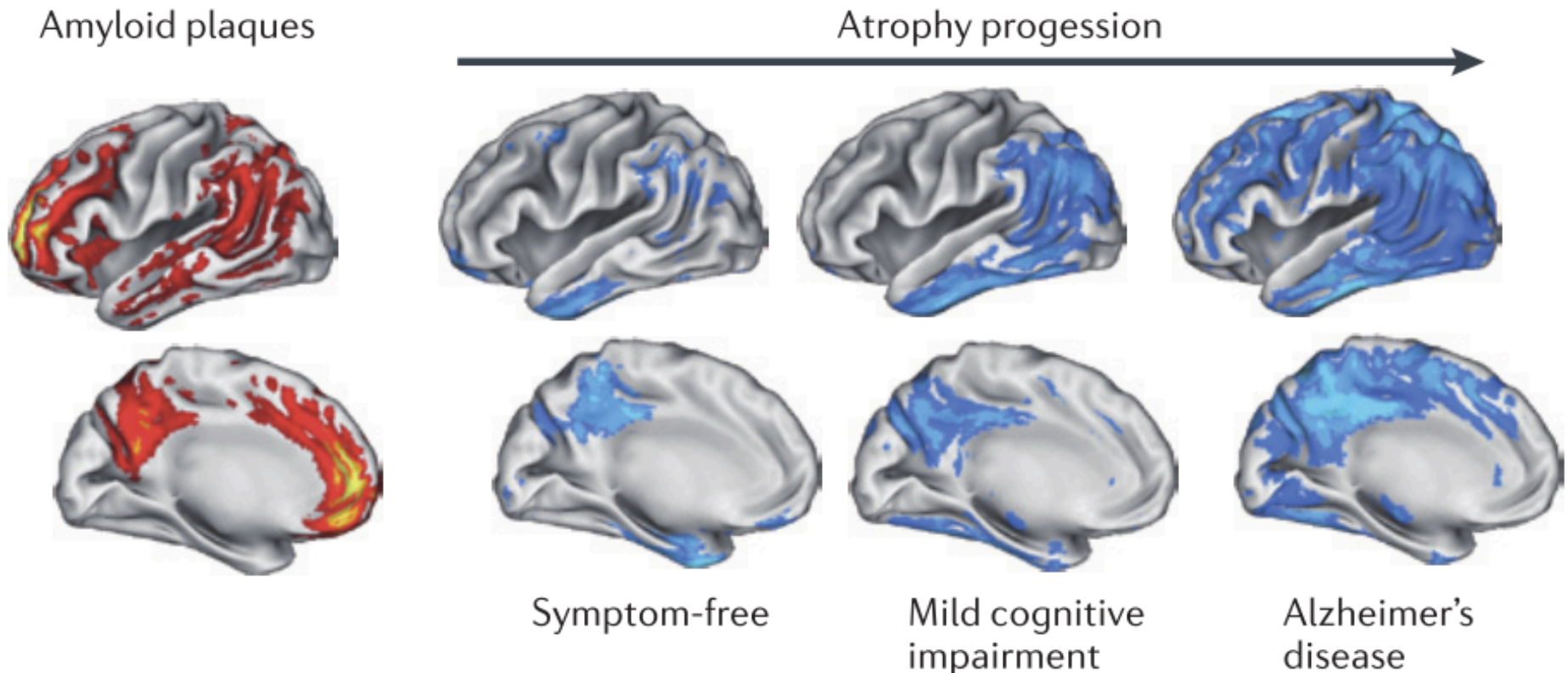
**Connectivity and
its impact**

Mapping the connectome and clinical applications

- The **connectome** will provide **novel insights on the functioning of the brain**
- There are multiple **mental diseases that are caused by dysfunctions of brain networks**, for example:
 - Alzheimer's disease
 - Schizophrenia
 - Autism

Alzheimer's disease

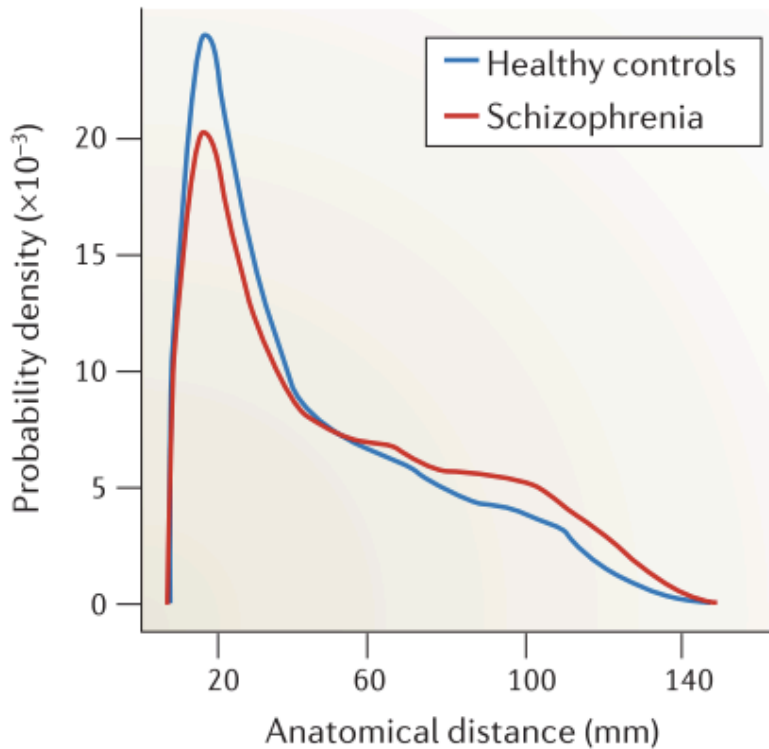
- The most expensive hubs are attacked by the disease



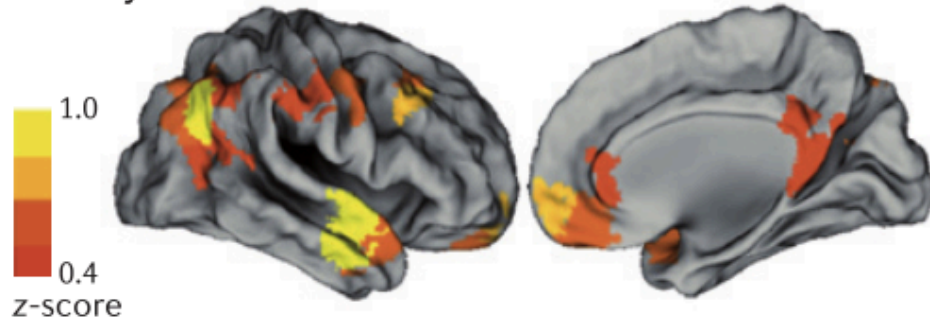
Schizophrenia

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

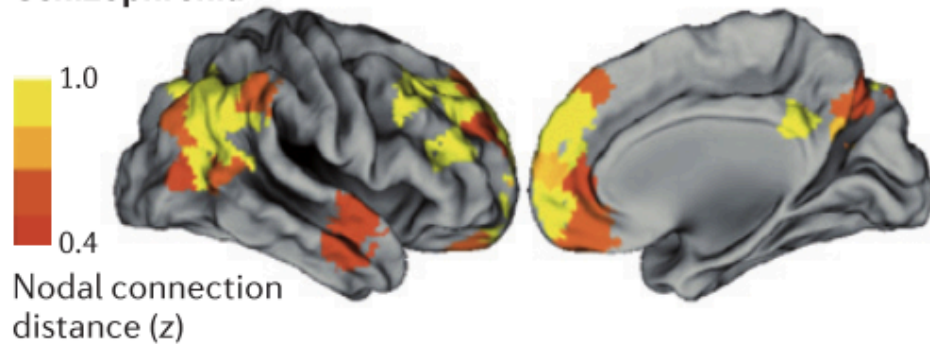
- **Unbalanced small-worldness**



Healthy volunteers



Schizophrenia

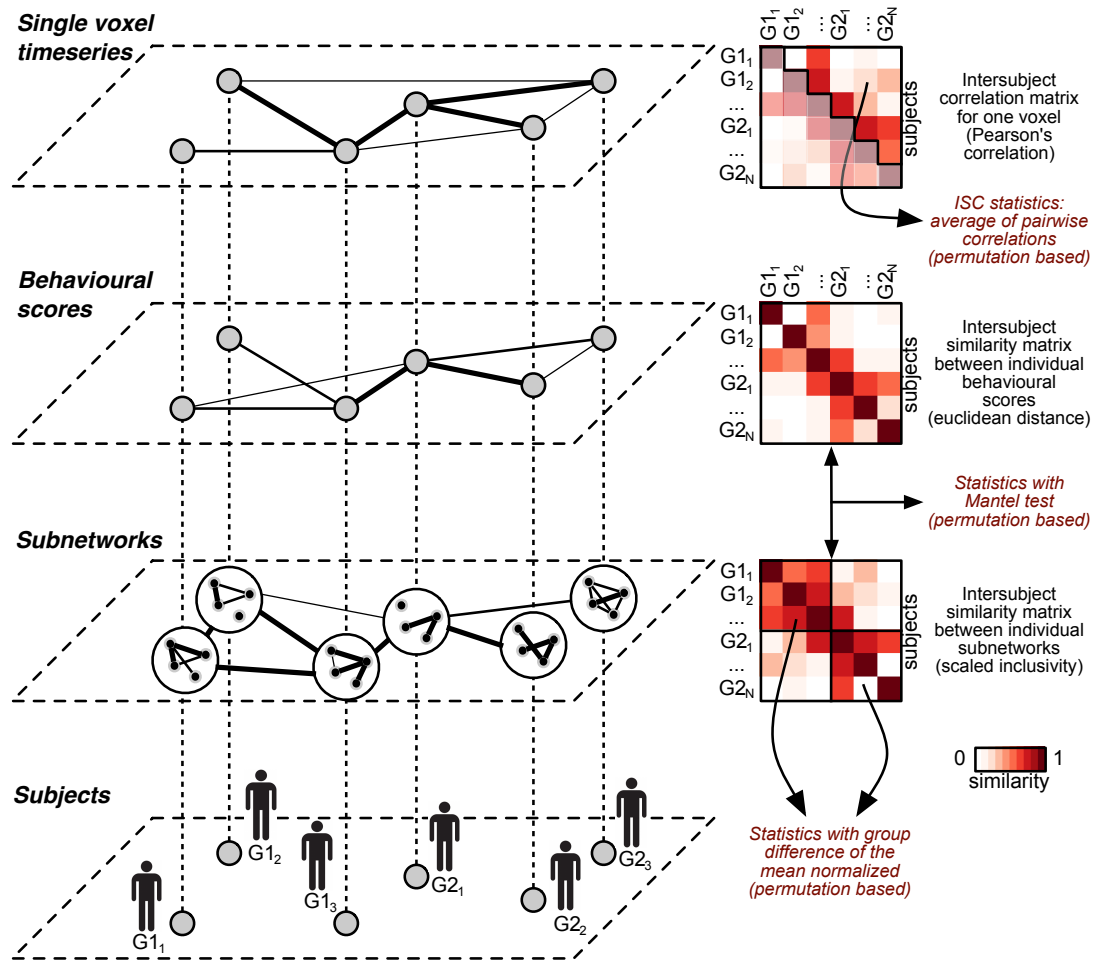


Reorganization of functionally connected brain subnetworks in high-functioning autism (Glerean et al 2016)

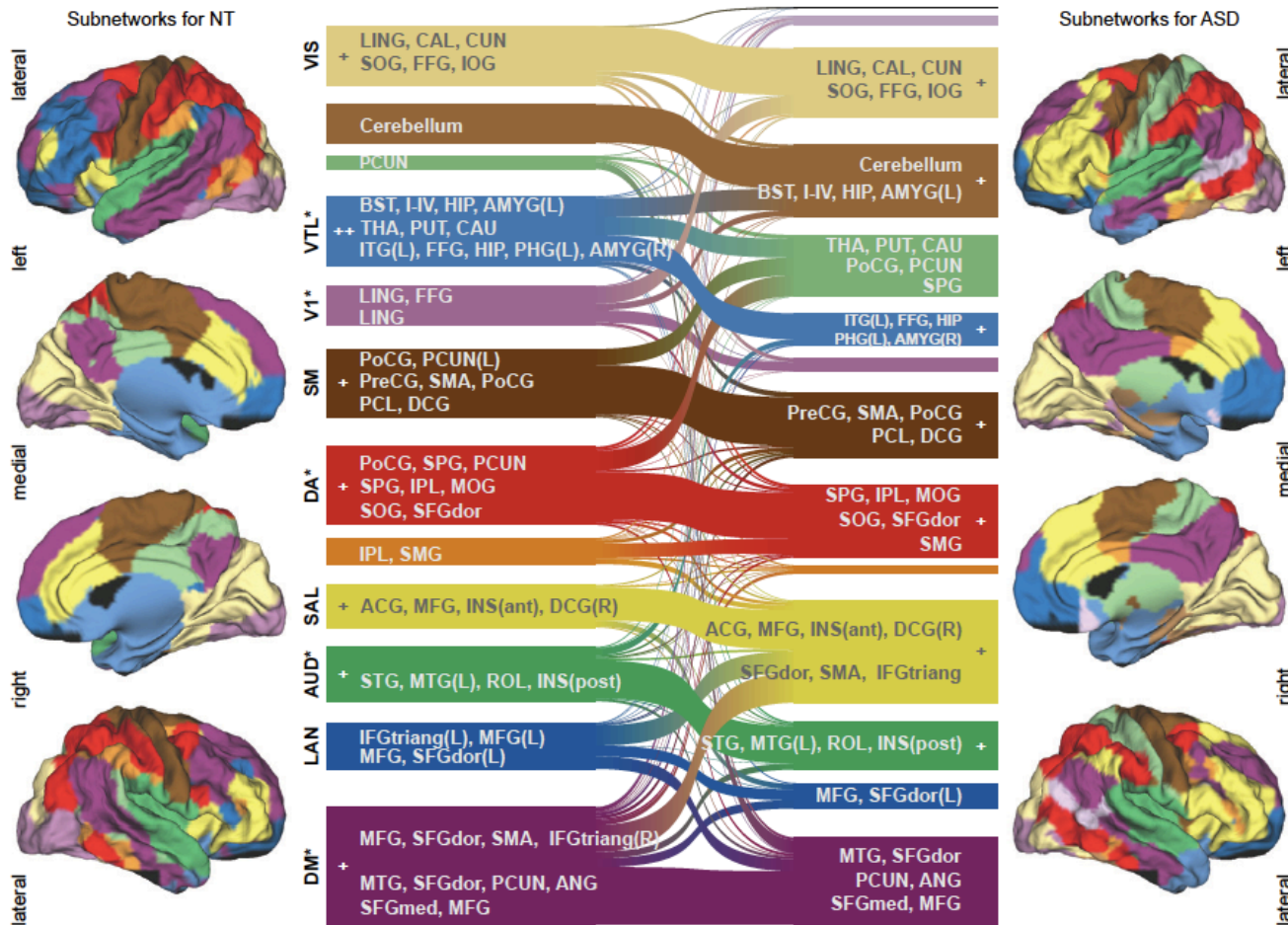
- Neuroimaging literature of ASD reports a **mixture of decreased and increased functional connectivity**.
AIM1) intersubject analysis framework to take into account the **heterogeneity of the disorder**.
AIM2) analyze connectivity at the subnetwork level to possibly resolve the mixture of findings at single node/link level.
- **Data:** 26 participants (13 with ASD), watching the movie *Tulitikkutehtaan tyttö* while undergoing fMRI. A replication resting-state dataset was included (data from the ABIDE initiative).

Intersubject analysis framework

- Assessing significance of ISC matrix
- Mantel test (comparison between similarity matrices)
- Comparing within groups/conditions similarities



Autism subnetworks (Glerean et al 2016)



Significant differences in:
Default Mode

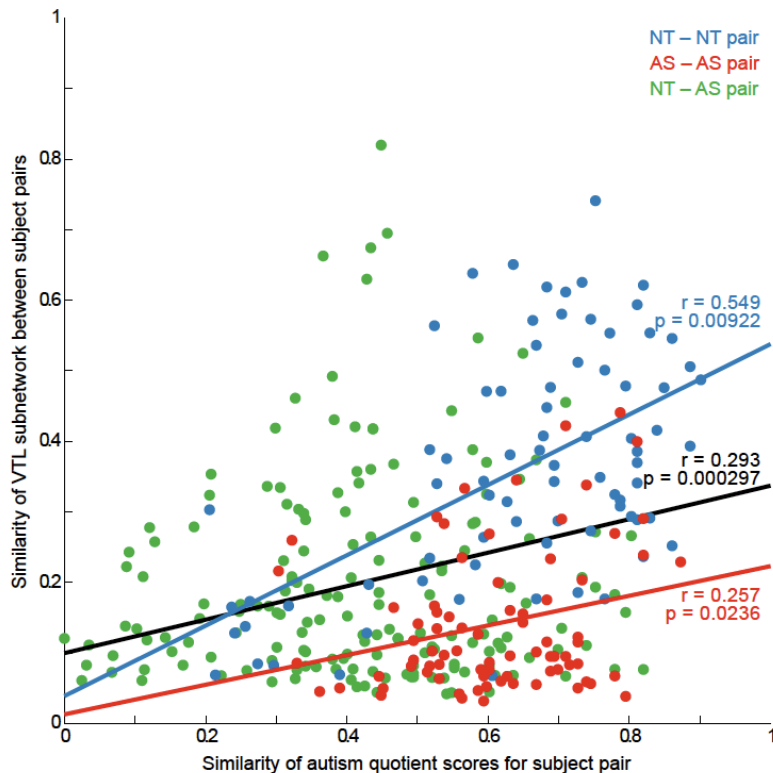
Auditory

Dorsal attention

Visual primary

**Ventro-temporo-
limbic (VTL)**

Results: correlation between AQ similarity and VTL similarity



The more two subjects have a similar VTL subnetwork, the more they have similar symptoms (amygdala, nucleus accumbens, putamen, caudate, thalamus, ventral visual pathway, ventro-medial prefrontal cortex)

The relationship is **also significant for controls**

Clinical uses?

CAN WE USE THESE TOOLS FOR
DIAGNOSTIC/NEUROSURGICAL
PURPOSES?

Clinical applications of resting state fMRI and network analysis

- Idea of putting a patient in the MRI scanner resting for ~5 minutes and get a diagnosis is intriguing, but **does it work?**
- **Open discussion in the field:**
 - Lee et al. 2013, **Resting-State fMRI: A Review of Methods and Clinical Applications**, AJNR doi: 10.3174/ajnr.A3263
 - Lang et al. 2014, **Resting-State Functional Magnetic Resonance Imaging: Review of Neurosurgical Applications**, Neurosurgery doi: 10.1227/NEU.0000000000000307
 - Castellanos et al, 2013, **Clinical applications of the functional connectome**, Neuroimage, doi: 10.1016/j.neuroimage.2013.04.083

Clinical applications of resting state fMRI and network analysis

- **Examples:**
 - **Presurgical planning** in patients with **brain tumor** or **intractable epilepsy** (less demanding than an active task in the scanner) [e.g. tumor in sensorimotor cortex, medial temporal lobe epilepsy]
 - Diagnosis of **Alzheimer's disease** (classification based on network clustering coefficient of hippocampus), children with **ADHD** (although another paper has shown that classification based on behavioural score had the same or better performance than resting state)
 - Resting state fMRI and **deep brain stimulation** (please refer to previous references for more detailed examples and discussions)

Clinical applications of resting state fMRI and network analysis

- **My two cents**

- there are still **methodological issues** to consider (what is a node? Best way of computing a network? Global signal and other BOLD related artifacts: **head motion**, breathing rate, heart rate)
- Shifting from a “biomarker from a distribution” approach to **combination of biomarkers** and comparison between large pools of subjects using machine learning (**UK Biobank project**)

Future?

**FUTURE DIRECTIONS IN THE FIELDS OF
NETWORK SCIENCE AND BRAIN
CONNECTIVITY**

Future directions in the field

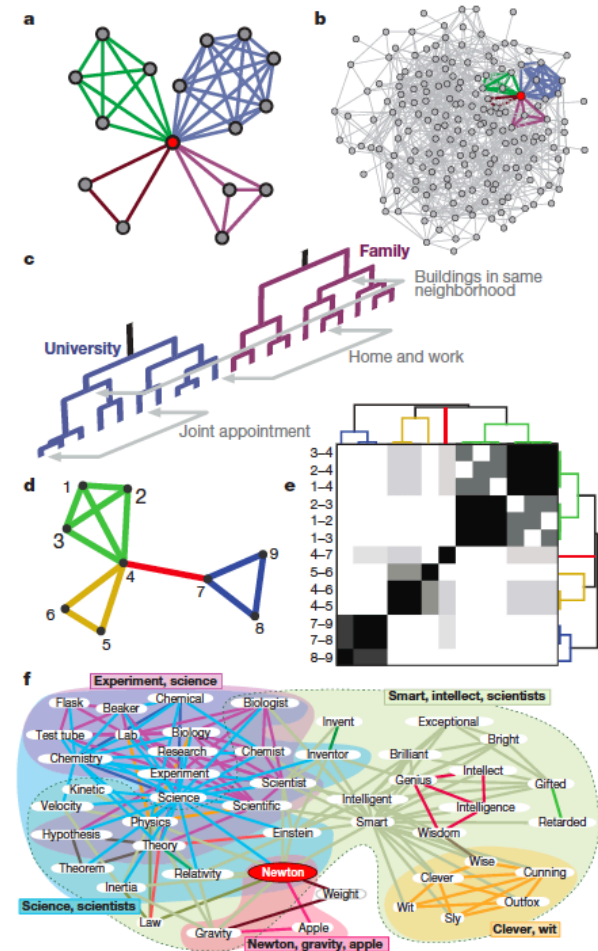
- **Line networks (link networks) and overlapping communities**
- **Multilayer and multiplex networks**
- **Networks of networks**

Overlapping communities

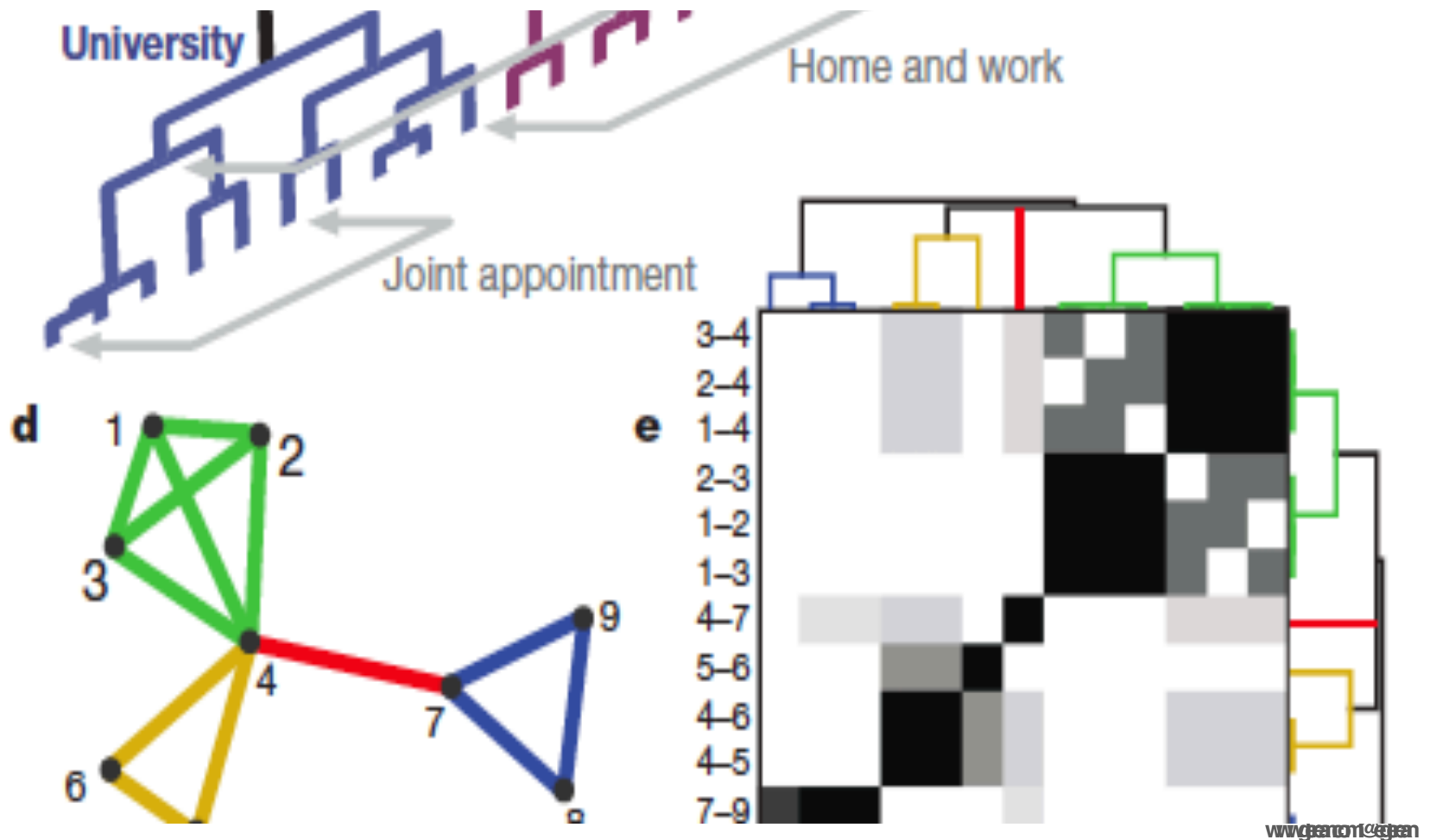
- Line networks (link networks) and overlapping communities

- See paper:

<http://www.nature.com/nature/journal/v466/n7307/abs/nature09182.html>

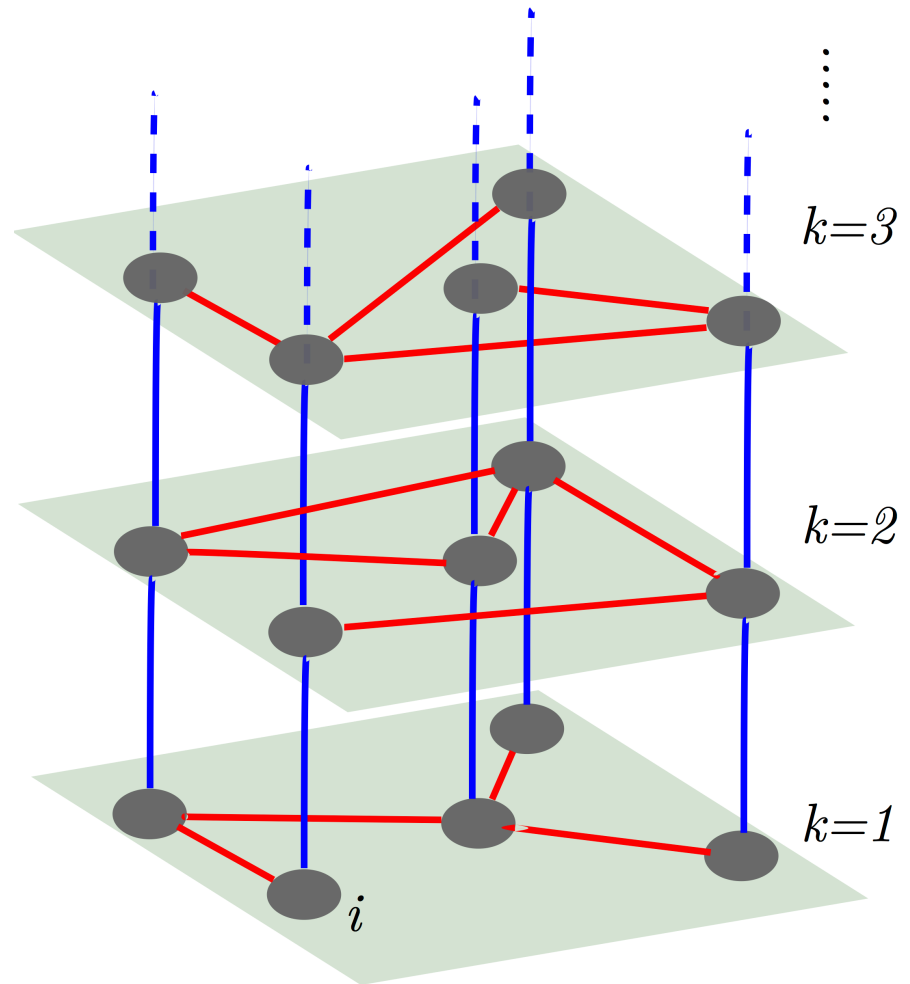


Overlapping communities



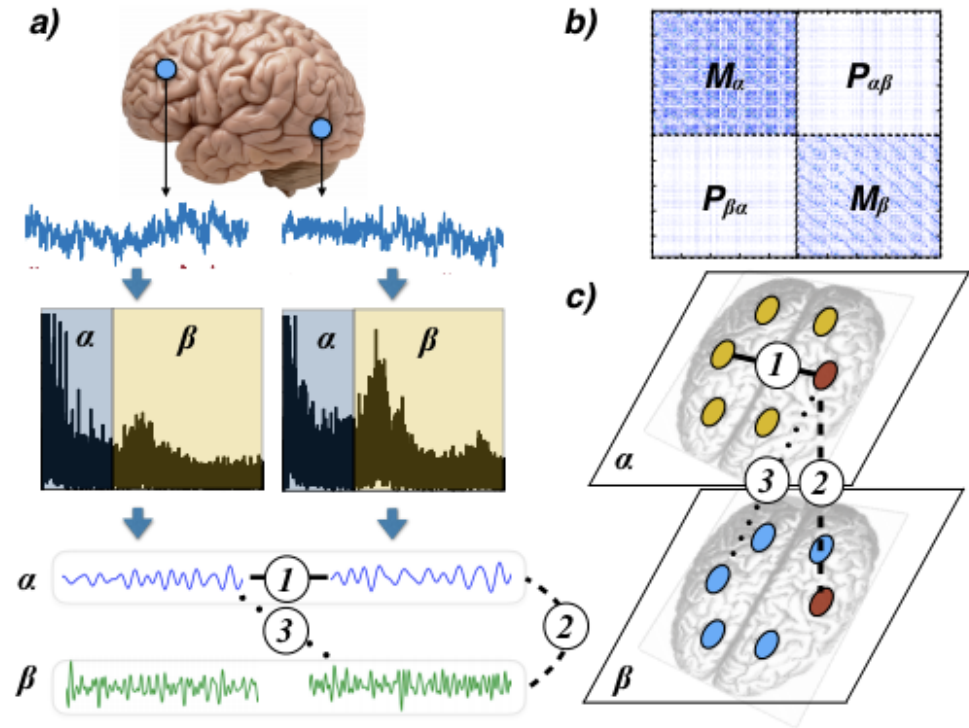
Multiplex networks

- Multiple networks where nodes are the same and connected with themselves through a 3rd dimension (e.g. subjects, time points, frequency bands)



Multilayer networks

- Multiple networks where nodes are connected with all other nodes in other layers

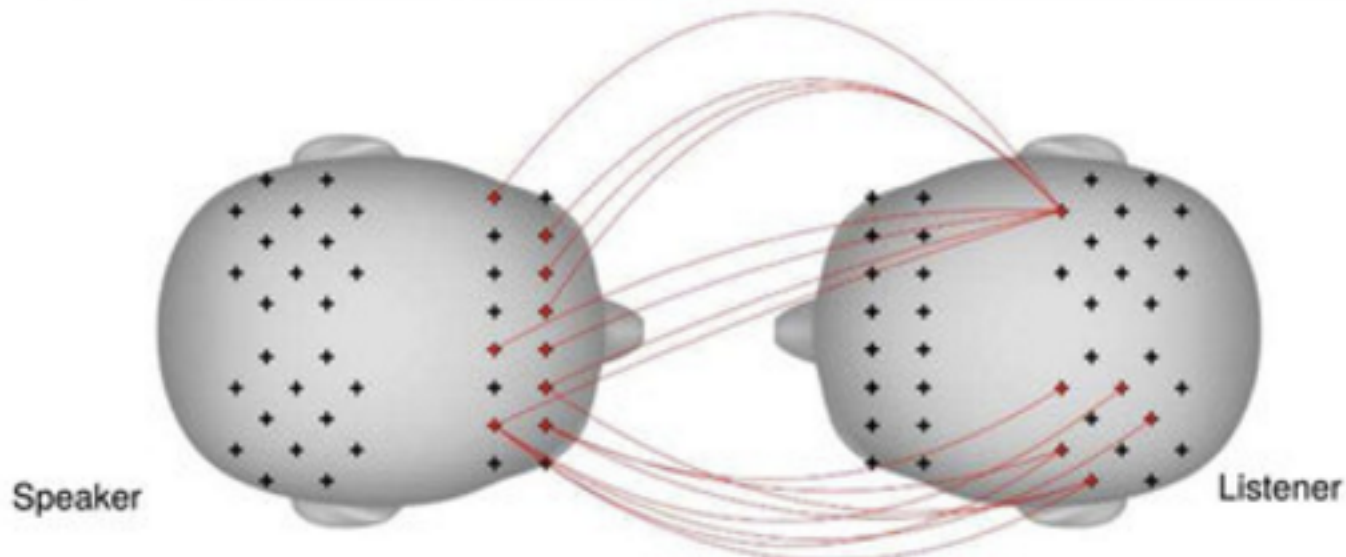


<https://arxiv.org/pdf/1703.06091.pdf>

Networks of networks

- Functional networks between subjects

C Speaker-listener significant couplings at 5 sec shift (FDR $q < 0.01$)



<https://www.nature.com/articles/srep43293>

Take home messages

Human brain networks

Take home messages

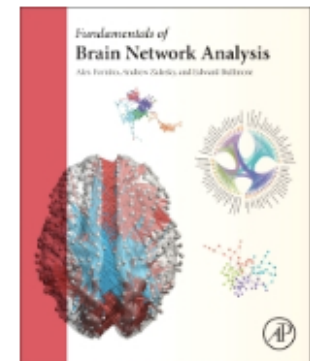
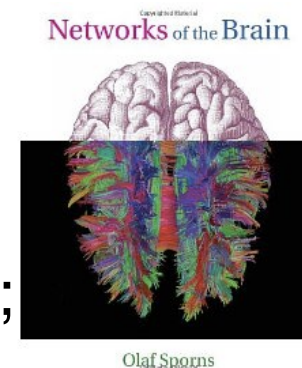
- **Brain network science** is a relatively recent field that is still evolving as new graph-theory methods are coming out. I personally think it is the way to go, and recent top papers in the field have been using brain connectivity methods.
- There are **multiple ways of modelling the brain as a network** and you just saw a glimpse. Do not be scared by the vast amount of options, start by replicating a paper you like.
- **Tools are still a bit scattered and choice of many parameters are left to the end user.** More rigorous automatic approaches should be devised

Some fundamental references

- **Bullmore, E., & Sporns, O. (2012). The economy of brain network organization.** Nature reviews. Neuroscience, 13(5), 336–49.
- **Craddock, et al. (2013). Imaging human connectomes at the macroscale.** Nature Methods, 10(6), 524–539.

- **Networks of the Brain**
Sporns, O; 2010, MIT Press.
- **Fundamentals of Brain Network Analysis.** Fornito, Zalesky, Bullmore; 2017, Elsevier

(They can be taken as book exams <http://www.brain-mind.fi/courses.html>)



...and something in Finnish about network science

<https://www.researchgate.net/publication/242719764> *Kompleksisten verkostojen fysiikkaa*