

Functional connectivity models: from blobs to (dynamic) networks

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 @dvdevill

Juan-les-Pins / November 22, 2017

Overview

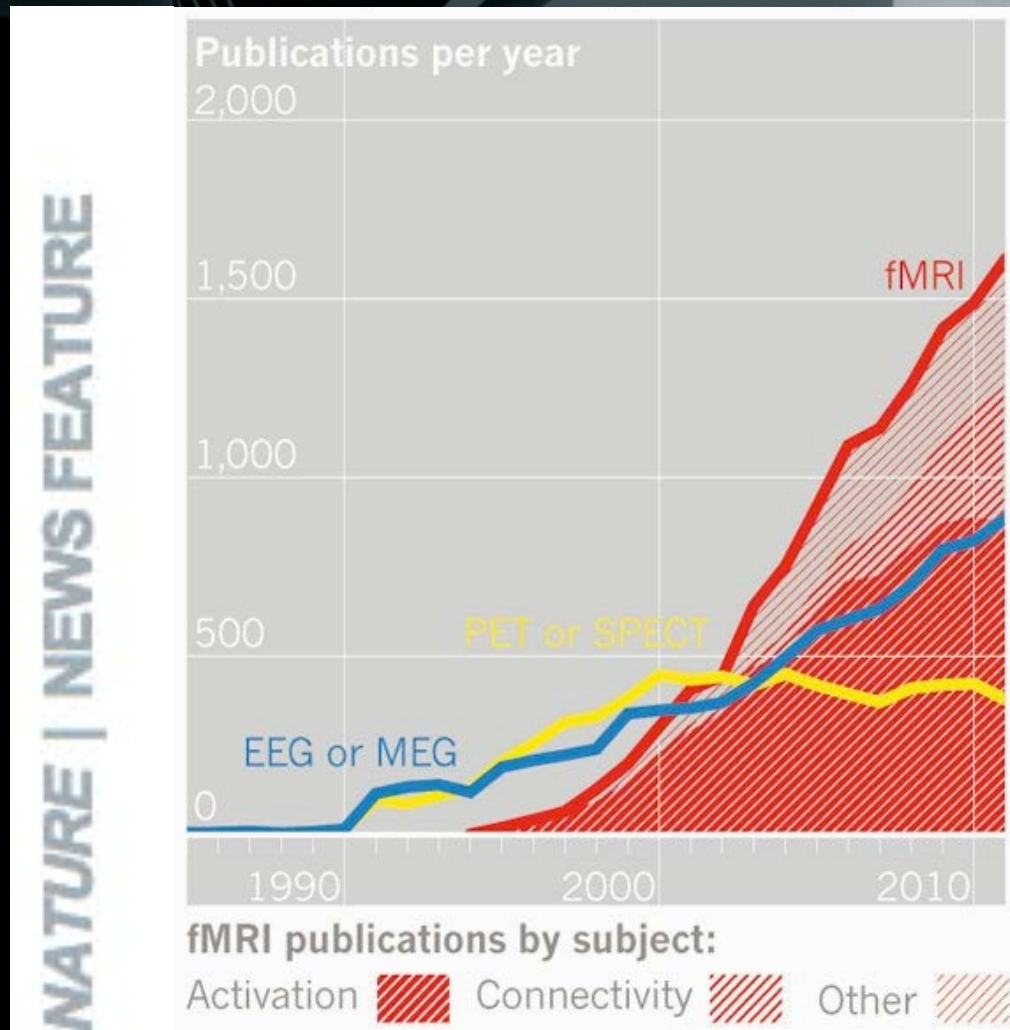
- The rise of (functional) MRI
- Activation mapping
 - Task-evoked activity
 - Brain activation maps using confirmatory analysis
 - General Linear Model (GLM)
- Connectivity mapping
 - Resting-state condition reveals intrinsic functional networks
 - Blind source separation (independent component analysis)
 - Statistical interdependencies and graph analysis
- Dynamic connectivity mapping
 - Moment to moment fluctuations of connectivity
 - Windowed and event-based approaches
- Open challenges

“the quest for an understanding of the functional organization of the [...] human brain, using techniques to assess changes in brain circulation, [a search that has occupied] mankind for more than a century”

Marcus Raichle, 1998

Magnetic resonance imaging (MRI)

- Widely deployed in hospitals and research centers
- Endogenous contrast mechanism
- Non-invasive imaging tool to study human brain anatomy and function



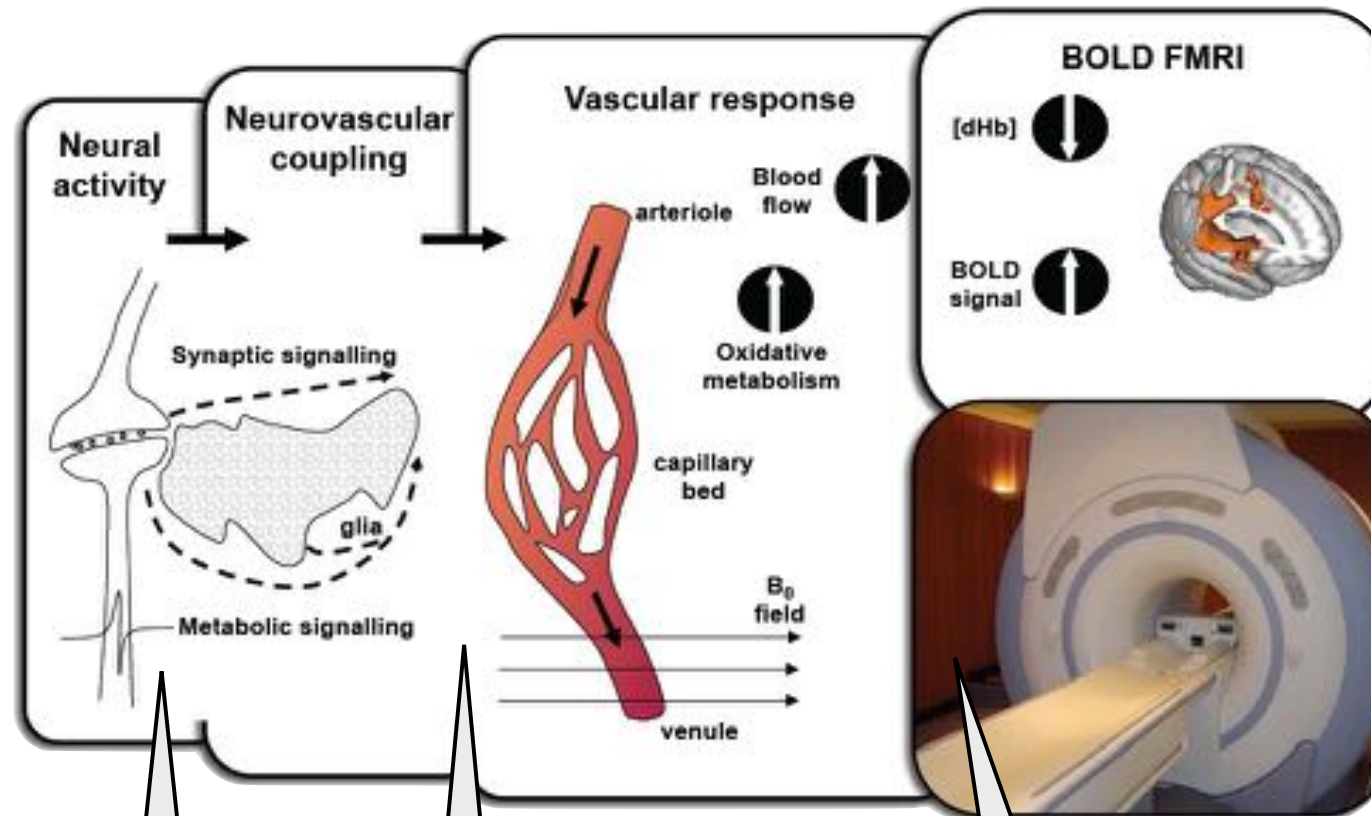
Functional MRI



- Series of 3D volumes
 - 3x3x3 mm³
 - 20-30 slices
 - every 2-4 sec
 - during 5-10 minutes



FMRI blood-oxygenation-level-dependent (BOLD) signals are slow proxy for neuronal activity



Neurons in the visual cortex are active (metabolic demand at synapses)

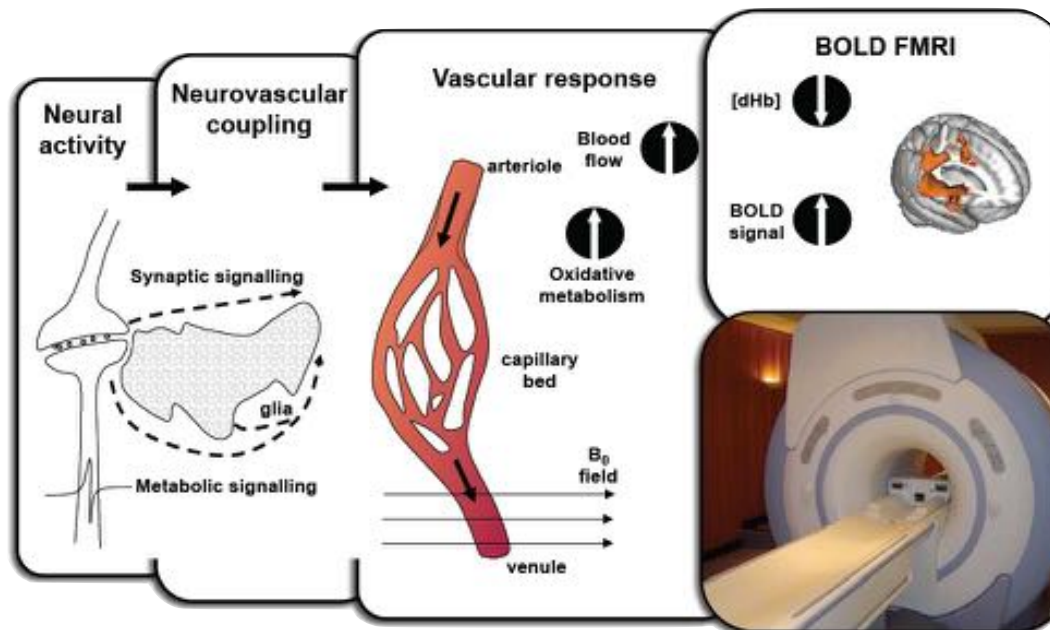
Neurovascular coupling: blood flow increases locally to bring in glucose and oxygen

Oxygen is carried by hemoglobin in RBCs

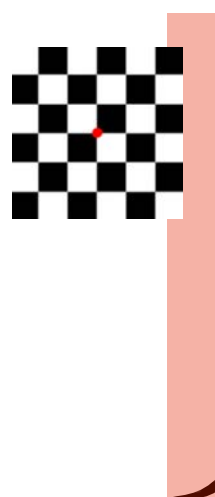
Increase of ratio in oxygenated/deoxygenated hemoglobin

Deoxygenated Hb is paramagnetic and drives BOLD signal

FMRI blood-oxygenation-level-dependent (BOLD) signals are slow proxy for neuronal activity



20 sec

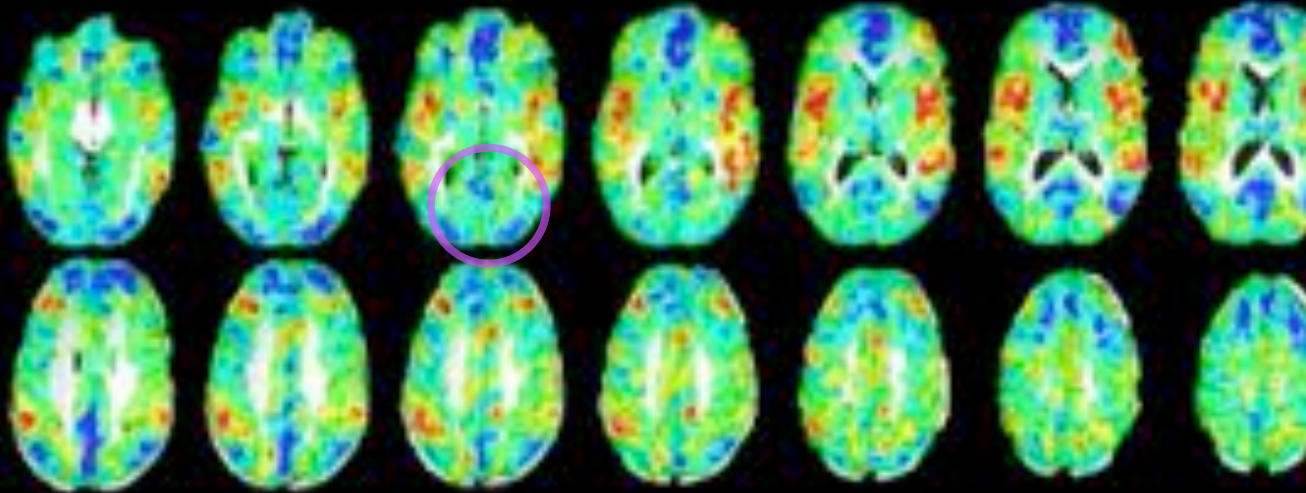


hemodynamic response function

- *Impulse response to short period of neural activation*
- *Notice the response timing: ~2 sec delay, 4-6 sec to peak, up to 20 sec back to baseline*

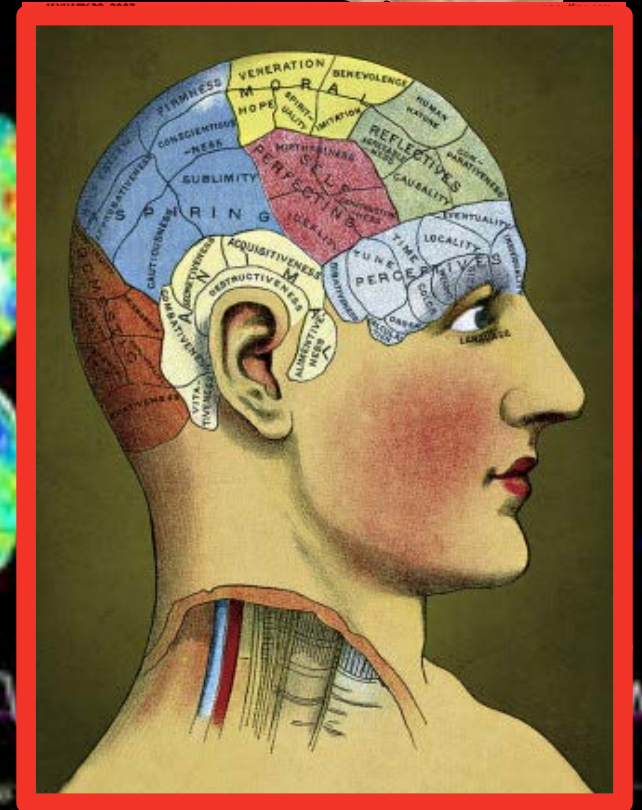
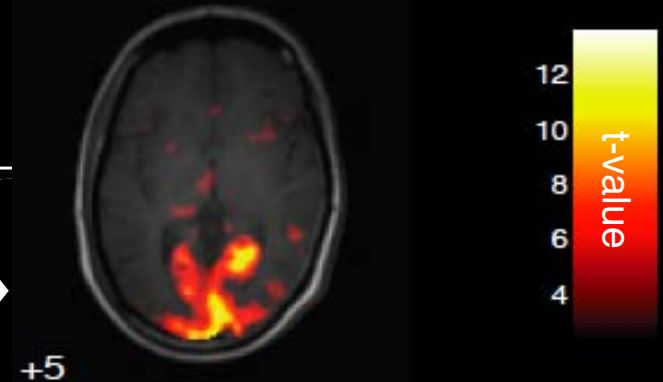
fMRI of evoked activity

min 0 max

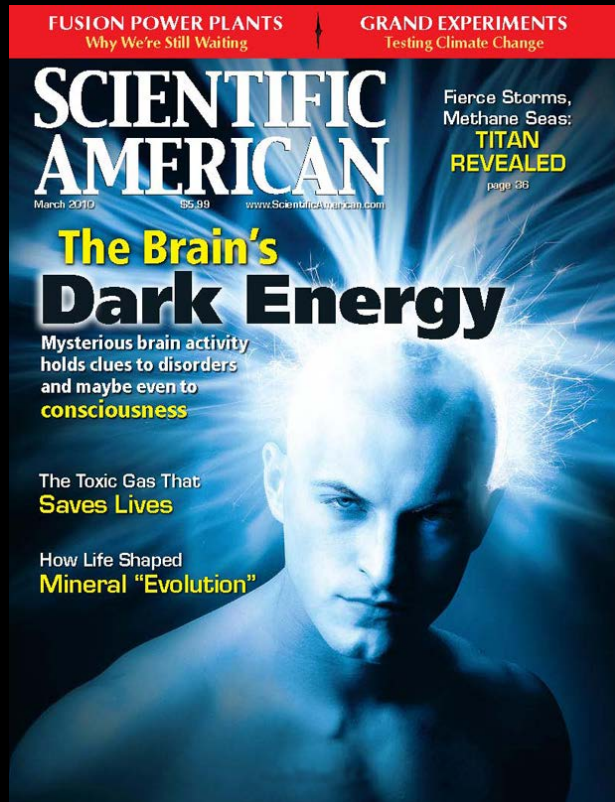


movie accelerated 4 times

GLM / statistical testing

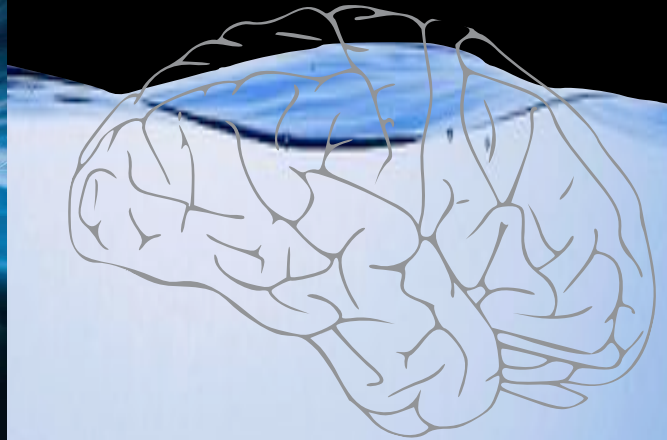


fMRI of evoked and intrinsic activity

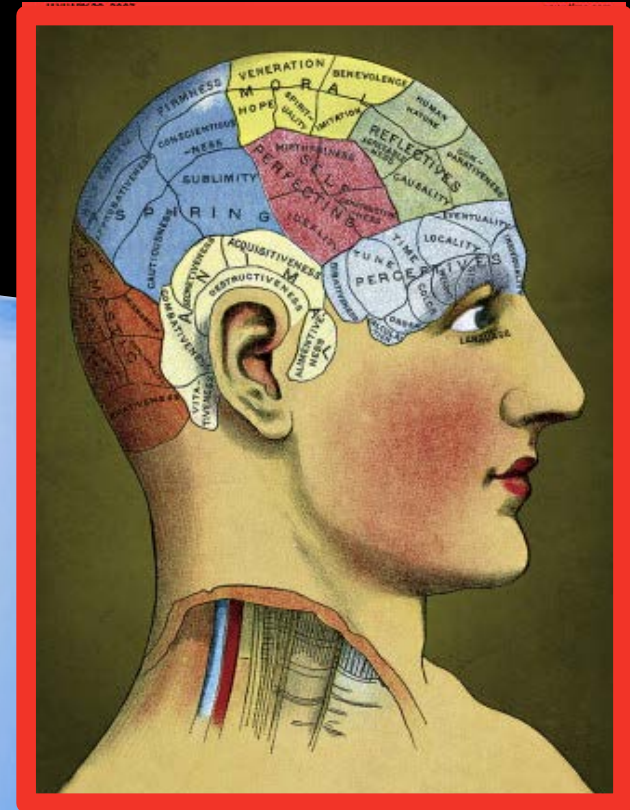


energy budget

5% evoked activity



95% intrinsic activity



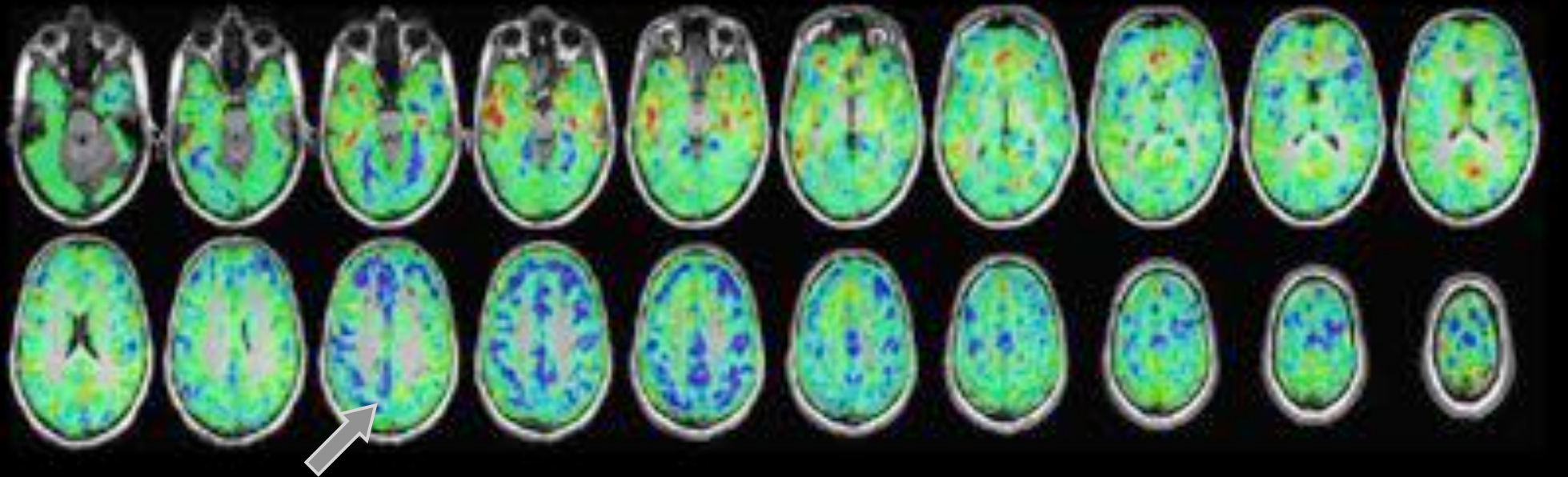
*Condition of resting state:
Conscious and unconscious brain activity
happen without premeditation or external
stimulus*

*More profound change of viewpoint:
From brain processing stimuli/performing tasks
to internal dynamics being modulated*

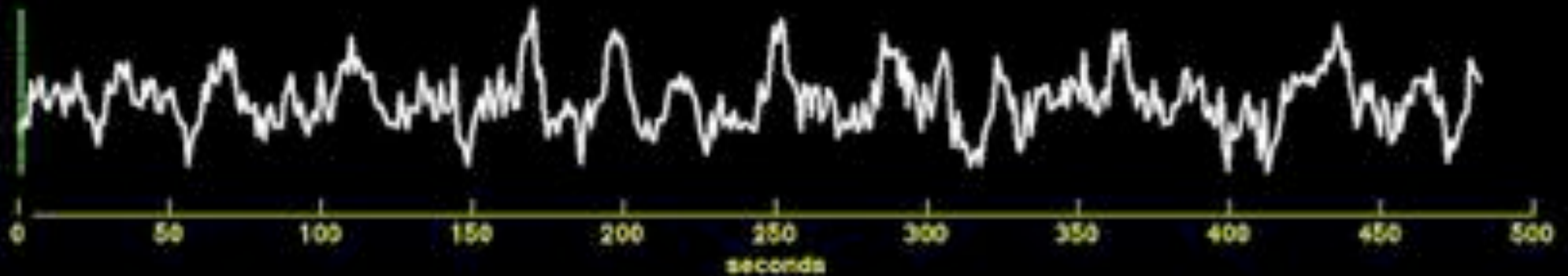
fMRI of spontaneous activity

resting-state scan (minimally preprocessed)

changes w.r.t. baseline
min 0 max



— BOLD signal (PCC)



movie accelerated 4 times

minimal compliance for patient studies!

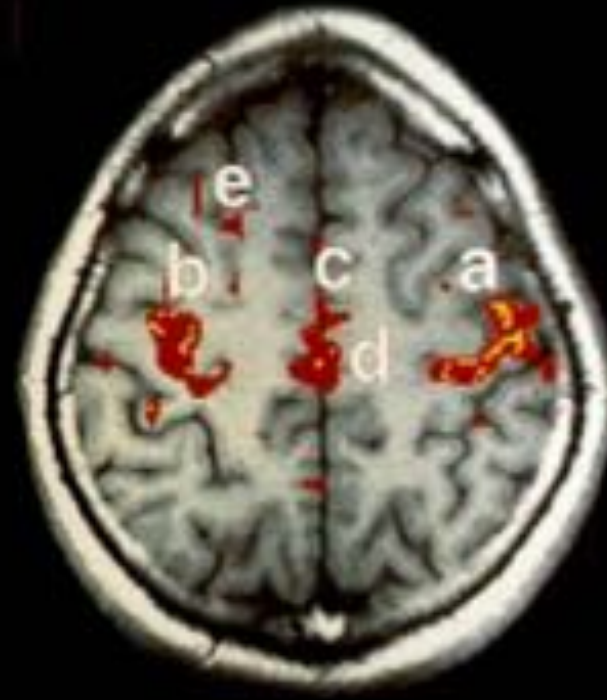
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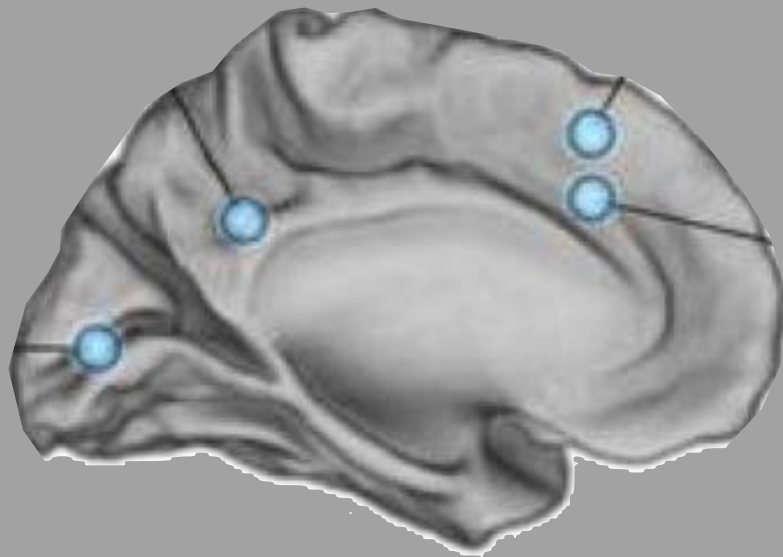
**Modeled activity
during finger tapping**



**High correlations
in spontaneous signal**



Seed-based connectivity maps



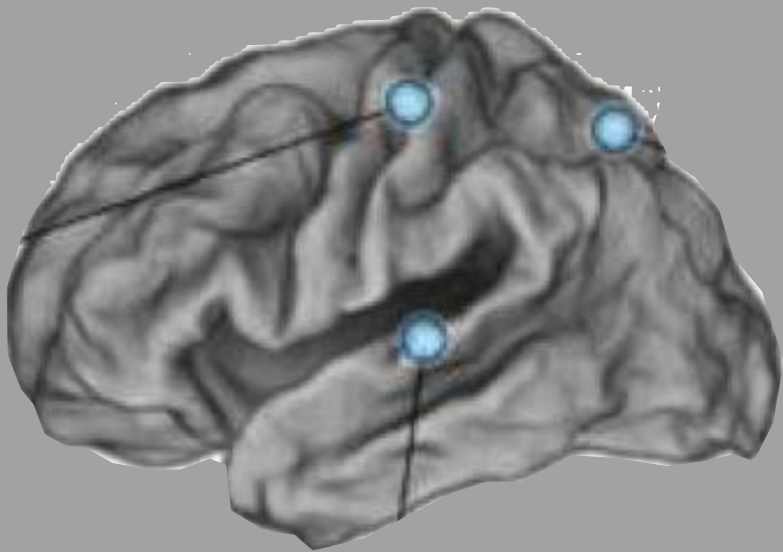
Default-mode network



Auditory



Visual



Somato-motor

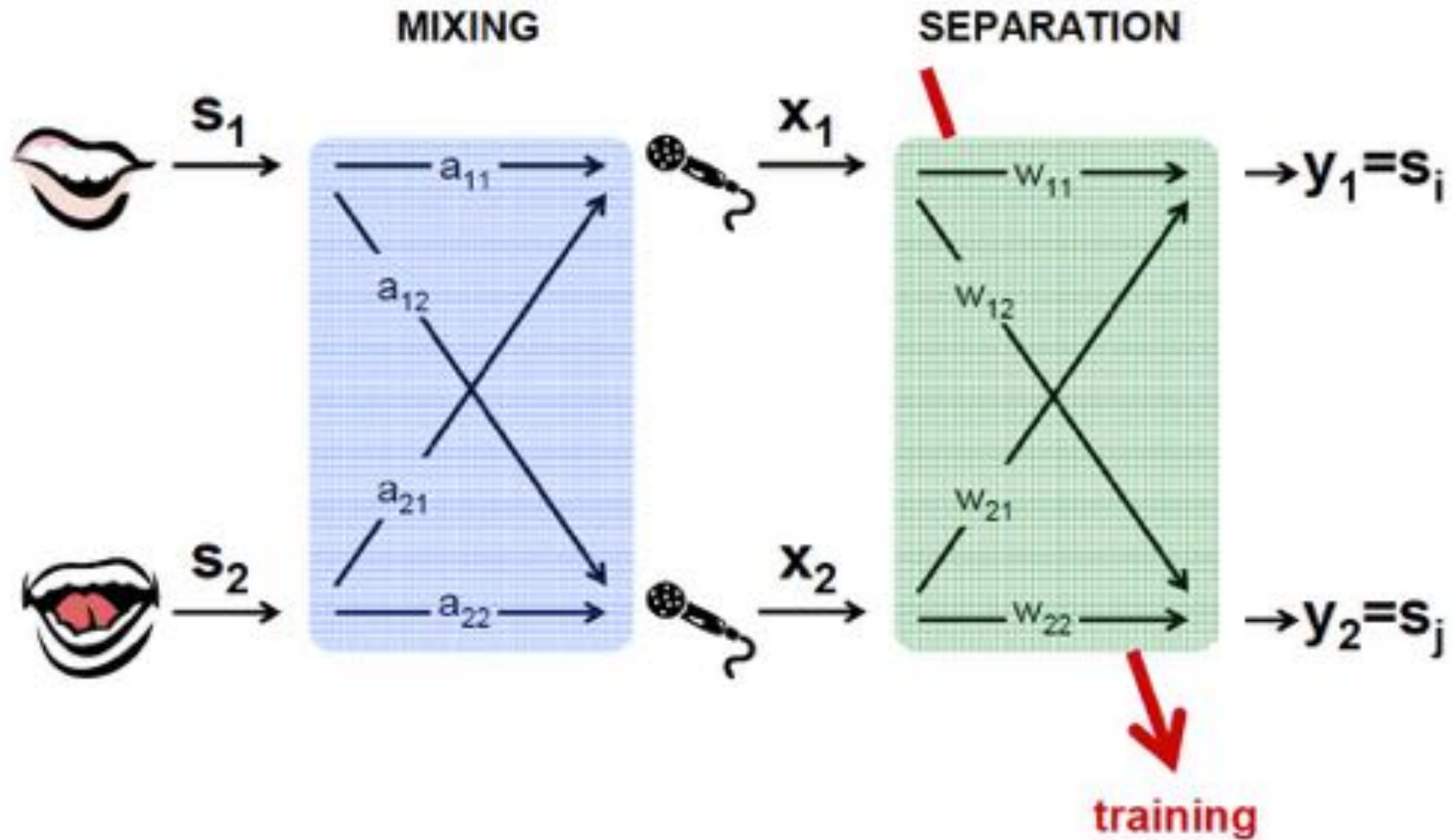


Dorsal attention

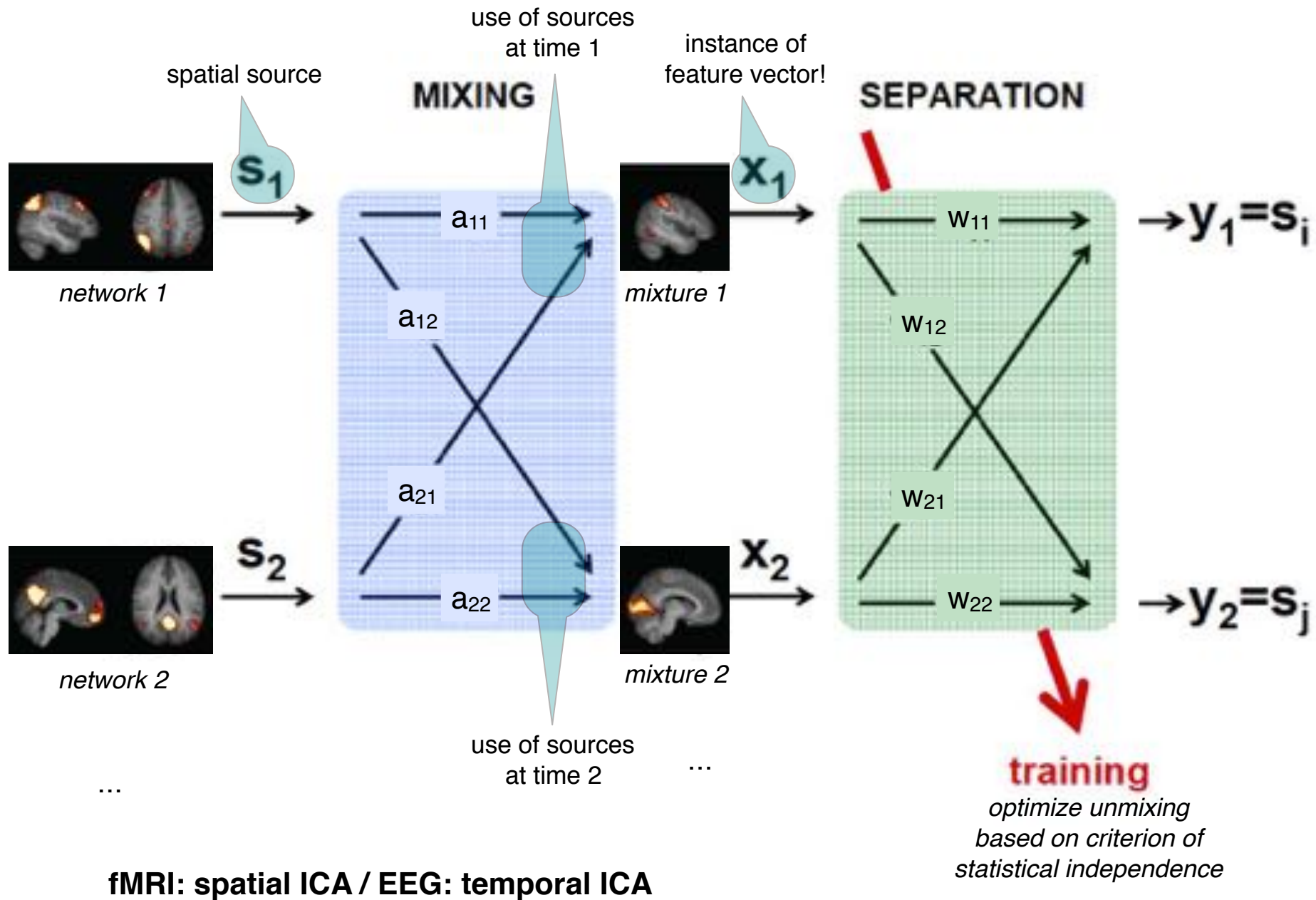


Executive control

Cocktail problem: blind source separation

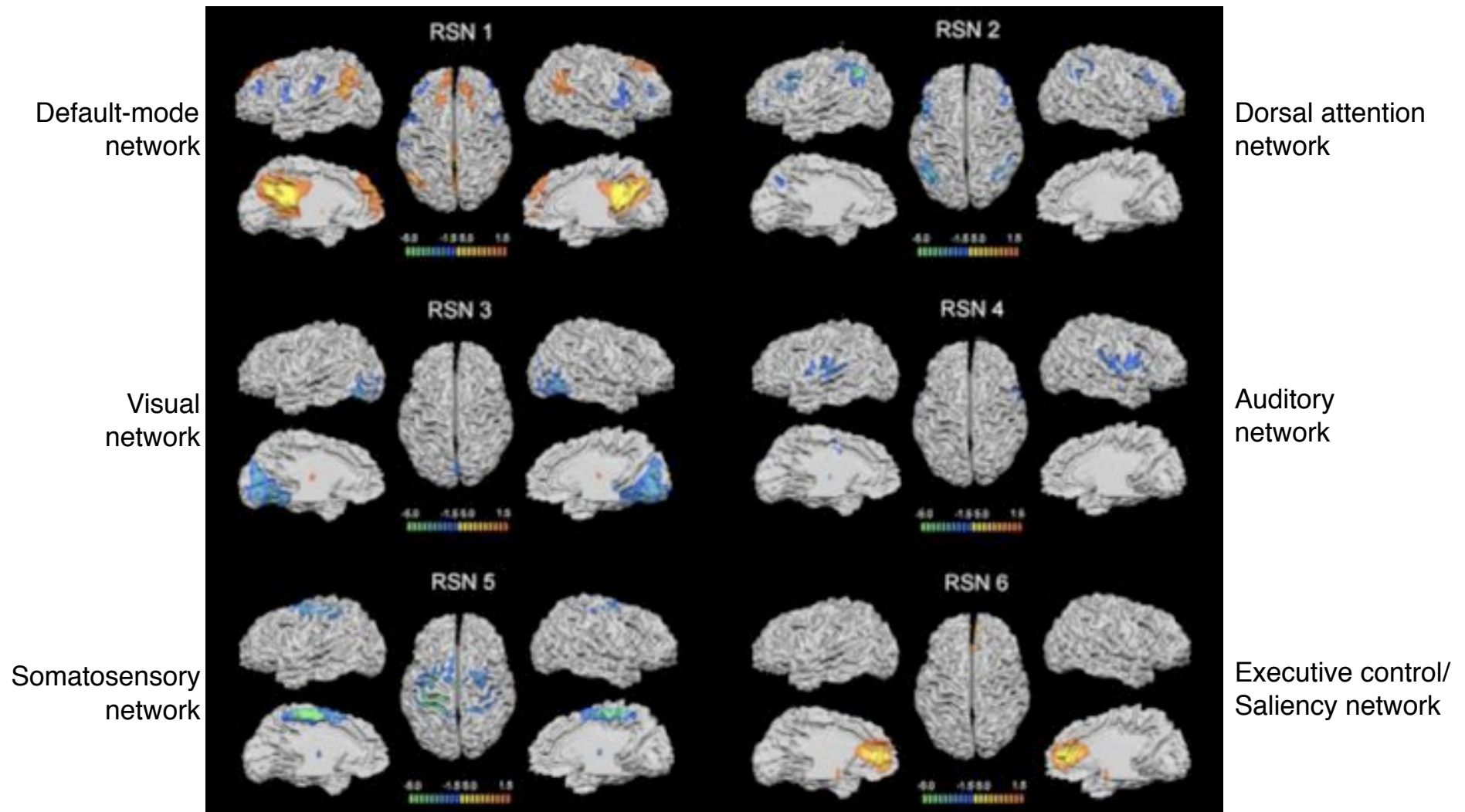


Mixing of spatial brain sources



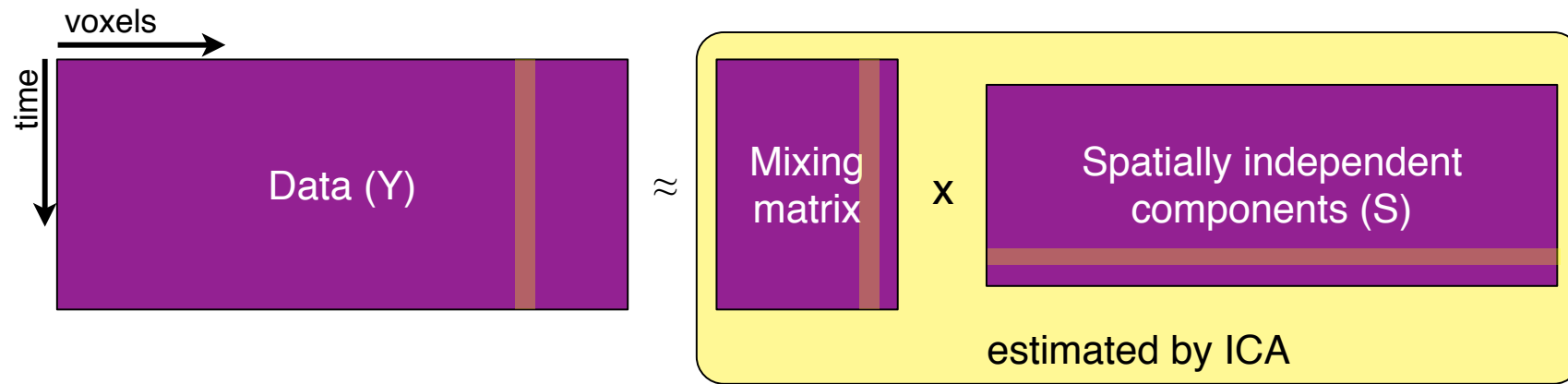
Unmixed spatial brain sources

- ICA reveals several large-scale brain networks (similar to task-evoked networks!)

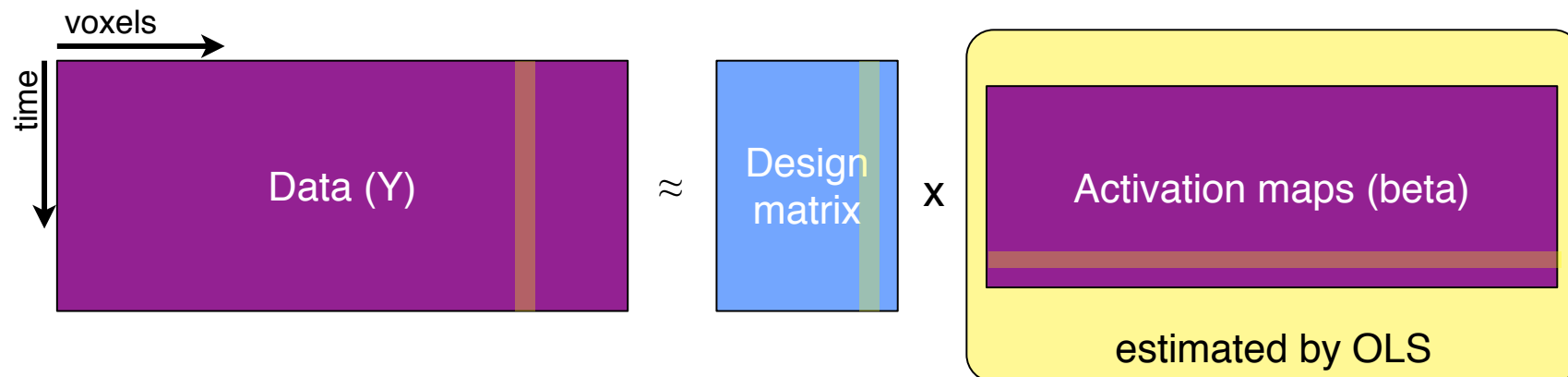


Spatial ICA

- Unmixing of brain regions from fMRI data

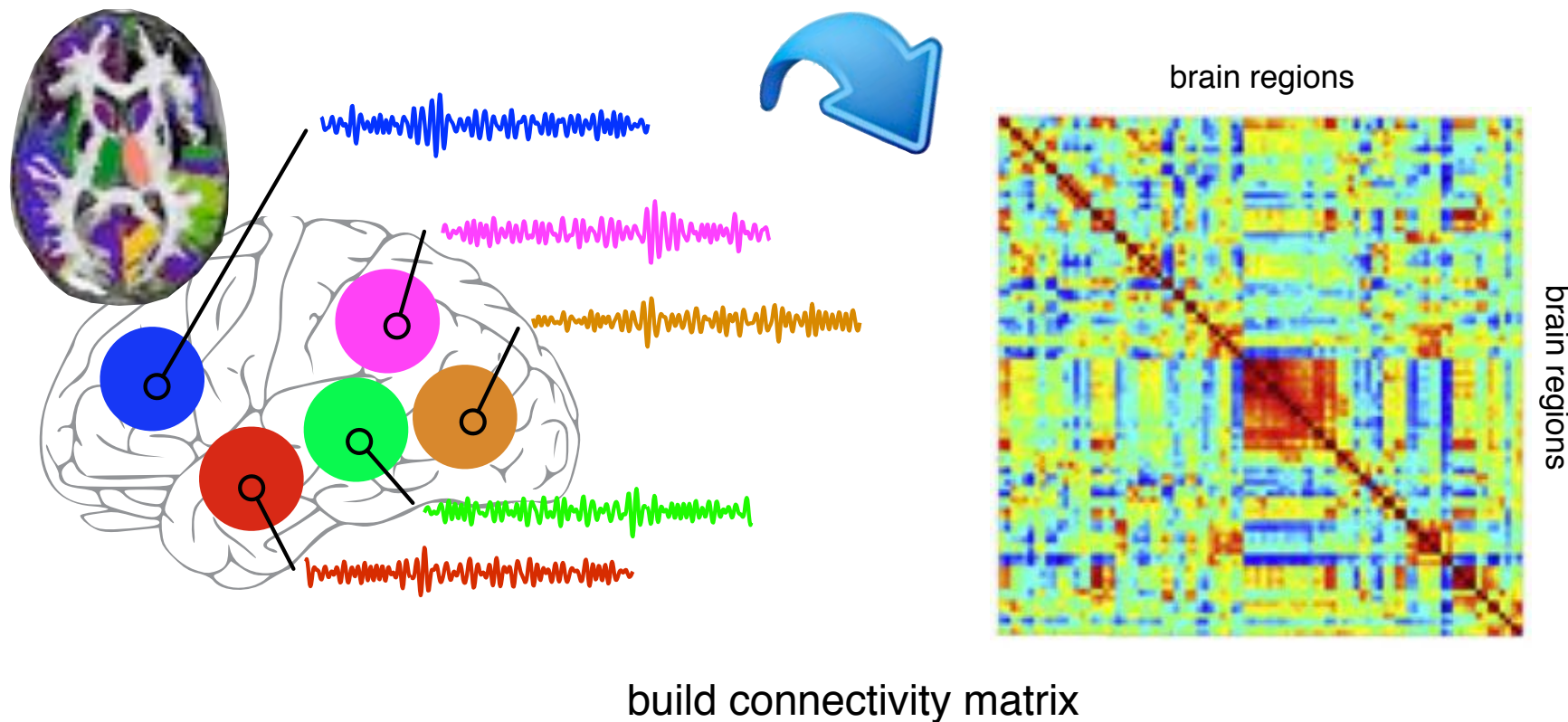


- Compare against the classical GLM



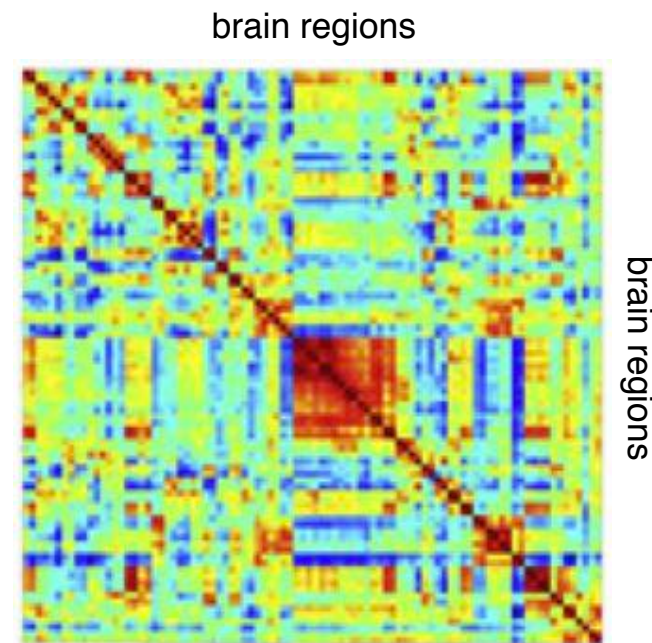
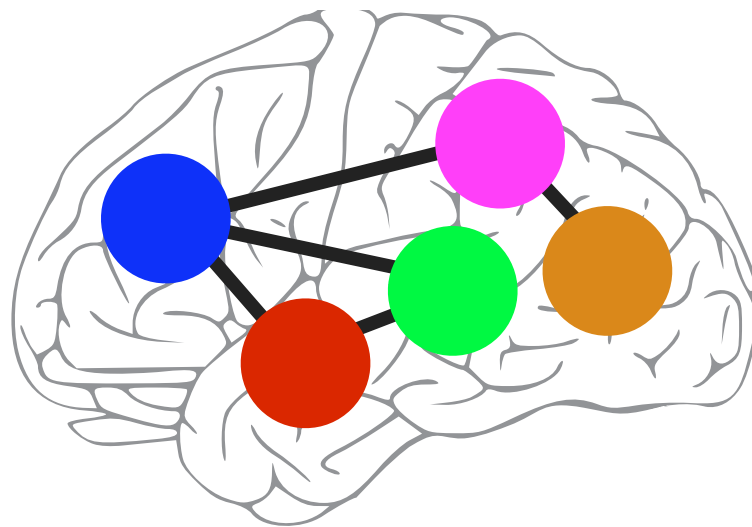
Graph analysis

- Parcellation into brain regions based on atlas
- Pairwise correlations between regionally-averaged timecourses of all brain regions

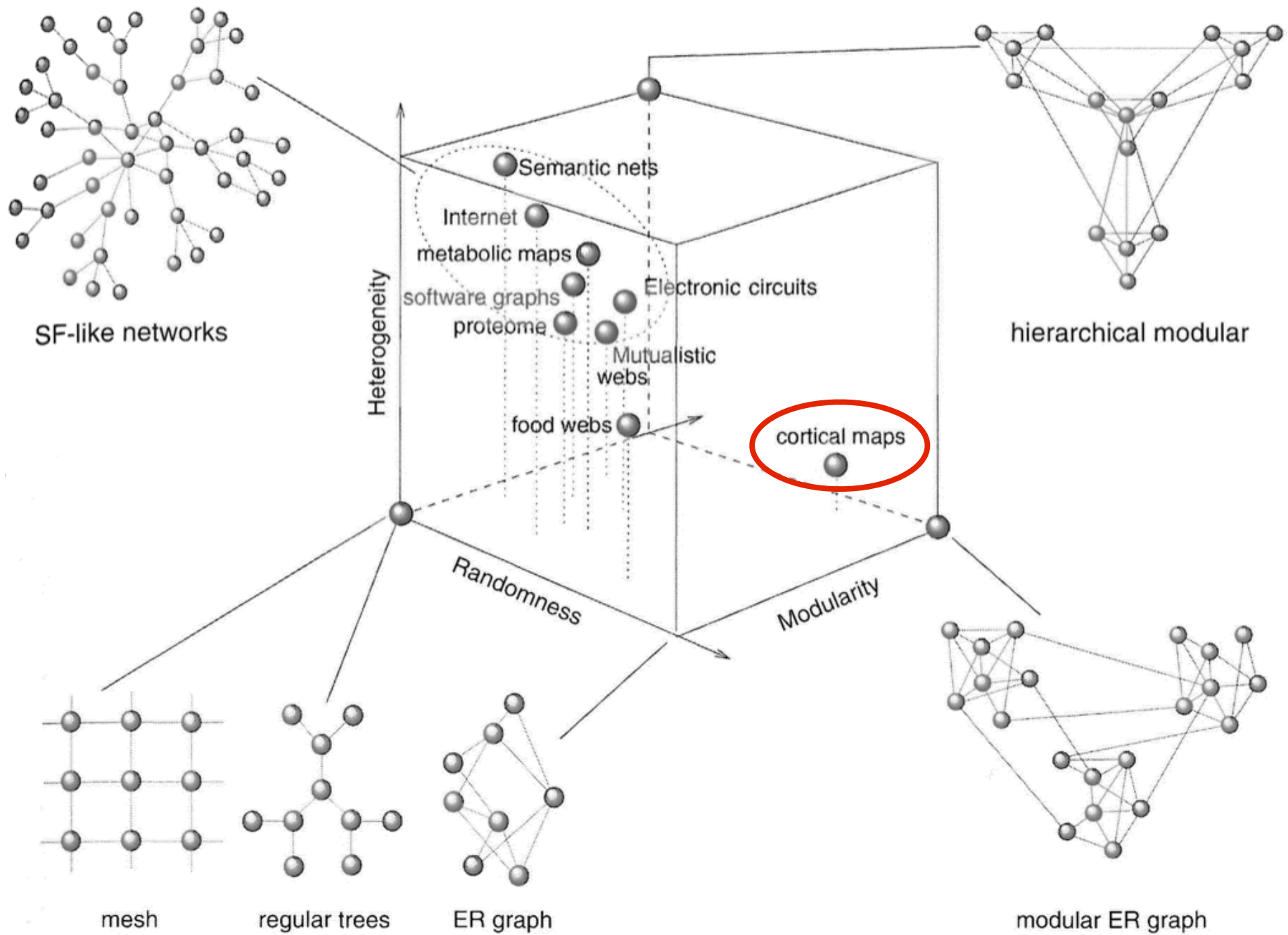


Graph analysis

- Parcellation into brain regions based on atlas
- Pairwise correlations between regionally-averaged timecourses of all brain regions



apply graph/network analysis



[Sole and Valverde, 2004]

■ Small-world

- high clustering
~functional segregation
- high efficiency
~functional integration

■ Cost-efficient

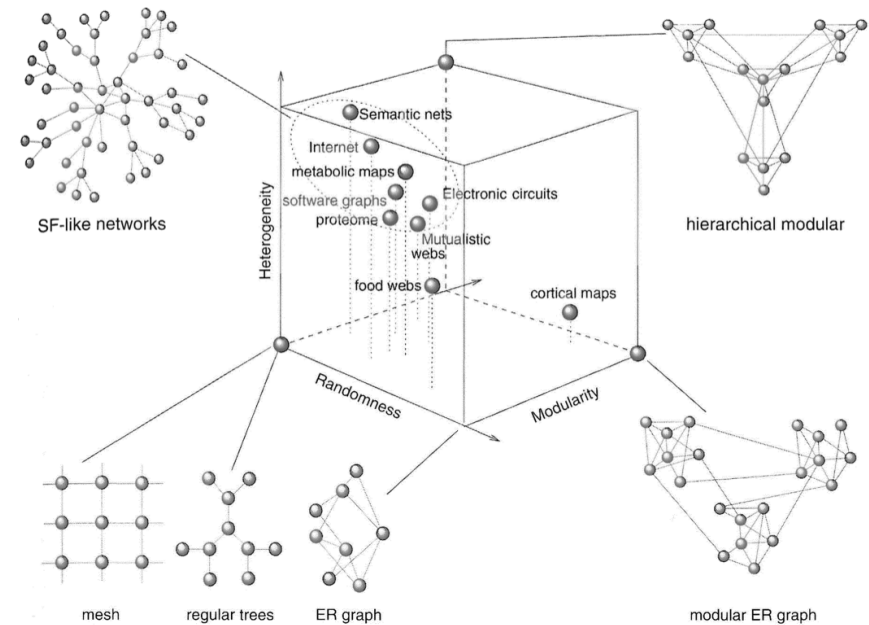
- high efficiency for
low connection cost

■ Hubby

- fat-tailed degree distributions
- hierarchical,
but still resilient to attacks and errors

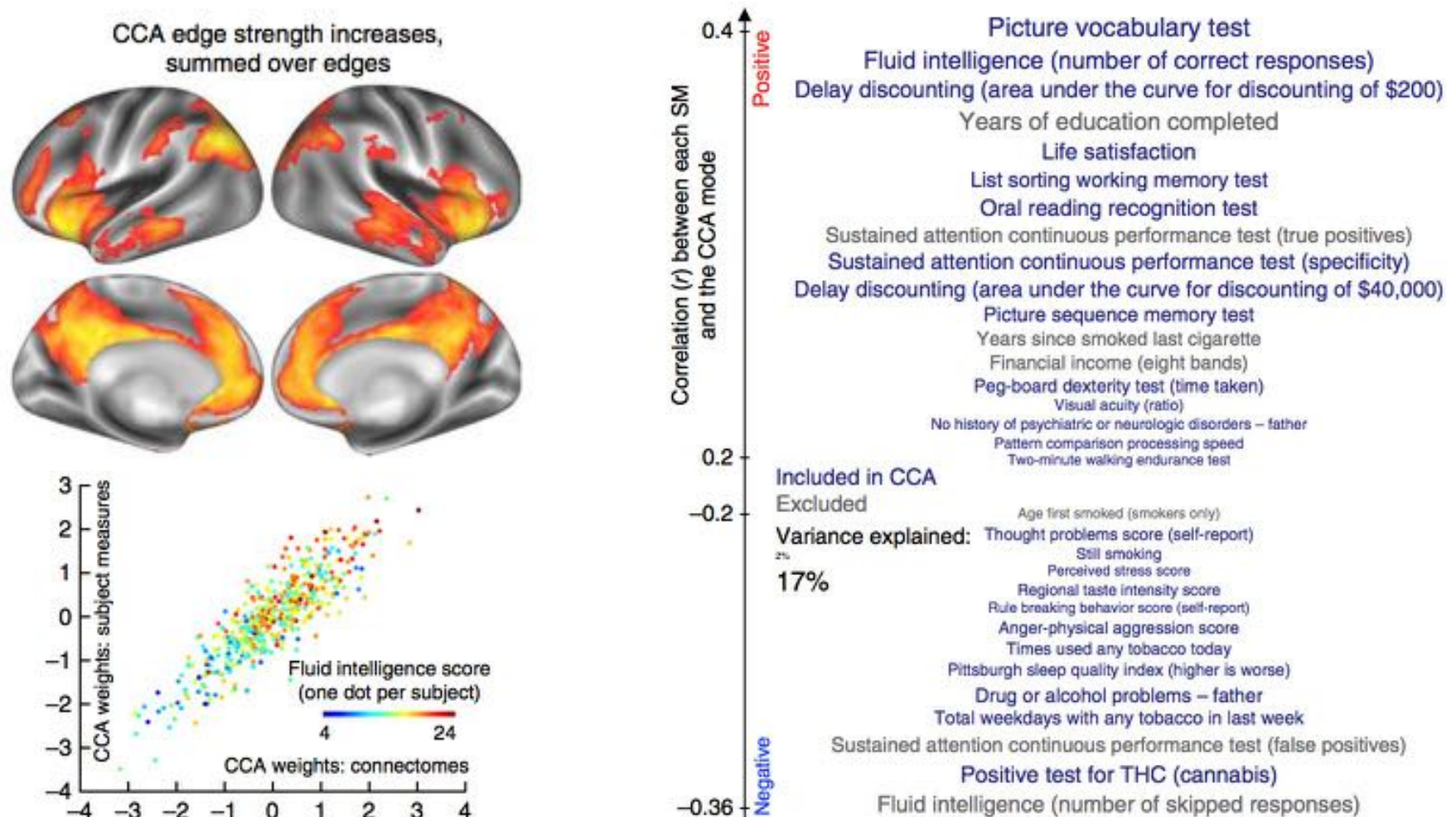
■ Modular

- more dense connections to nodes in
module than to nodes in
other modules



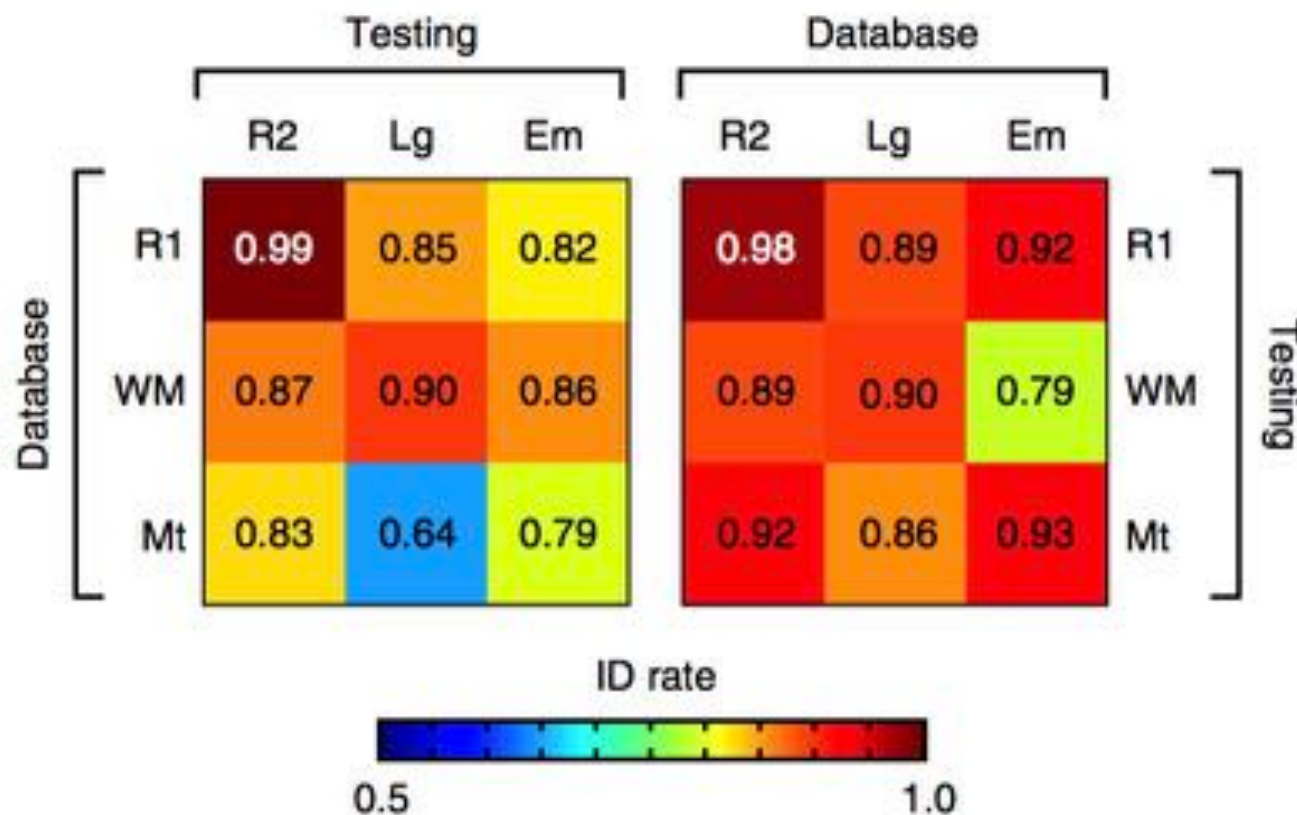
Show me your rest, and I tell your success

- Linking resting-state functional connectomes (280!) to life style/demographics/psychometric measures



Show me your rest, and I tell who you are

- Functional connectome serves as fingerprint to identify individuals
 - 126 individuals, across resting state & task sessions



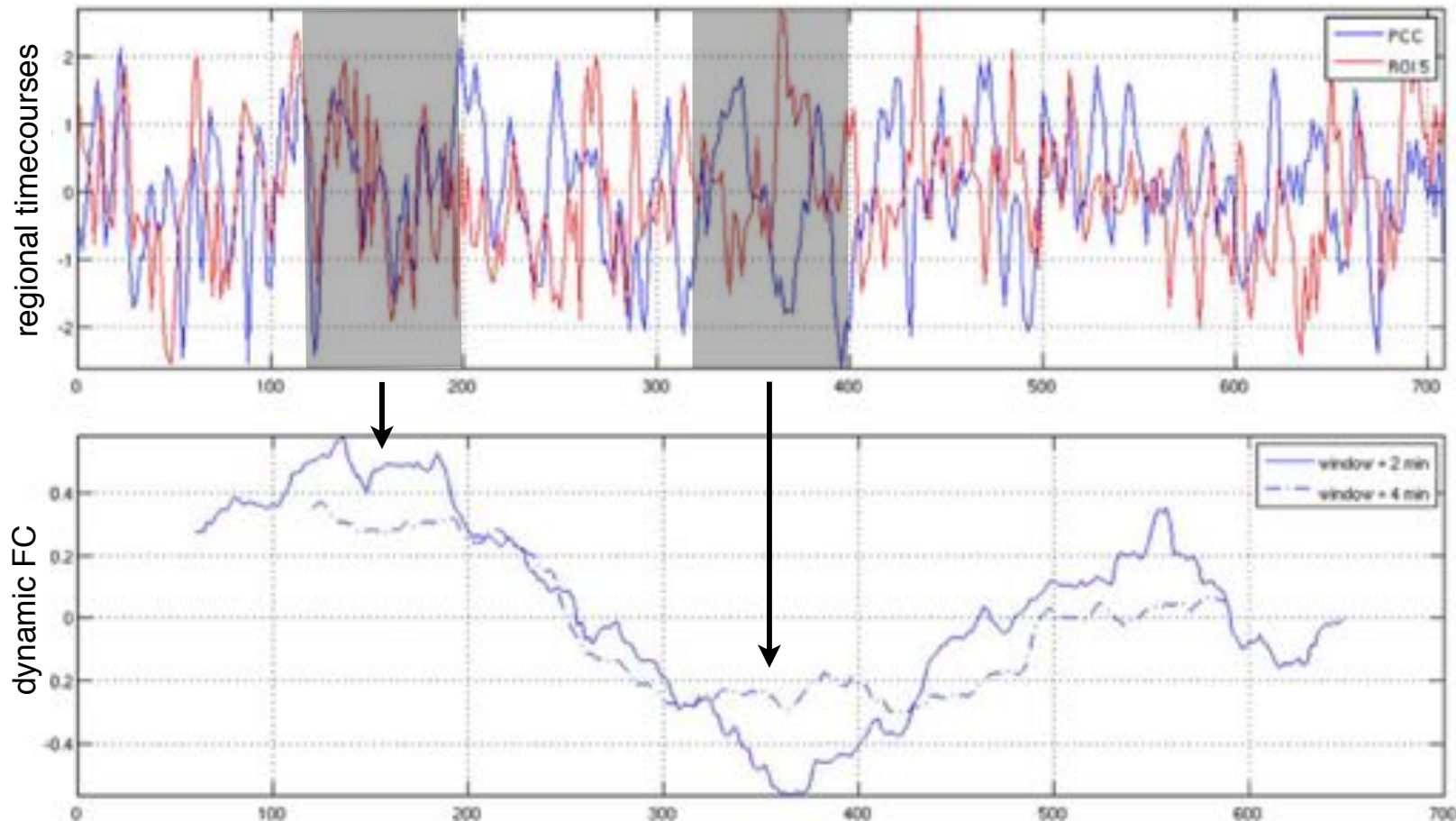
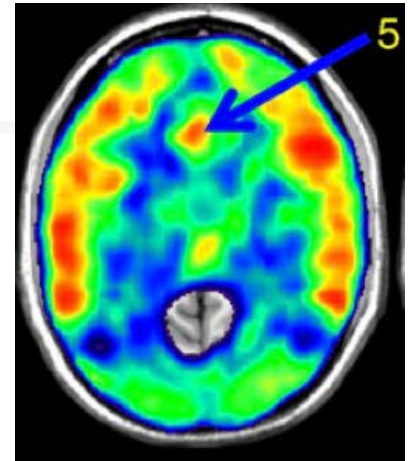
R1: rest 1, R2: rest 2, WM: working memory, Mt: motor task, Lg: language, Em: emotion

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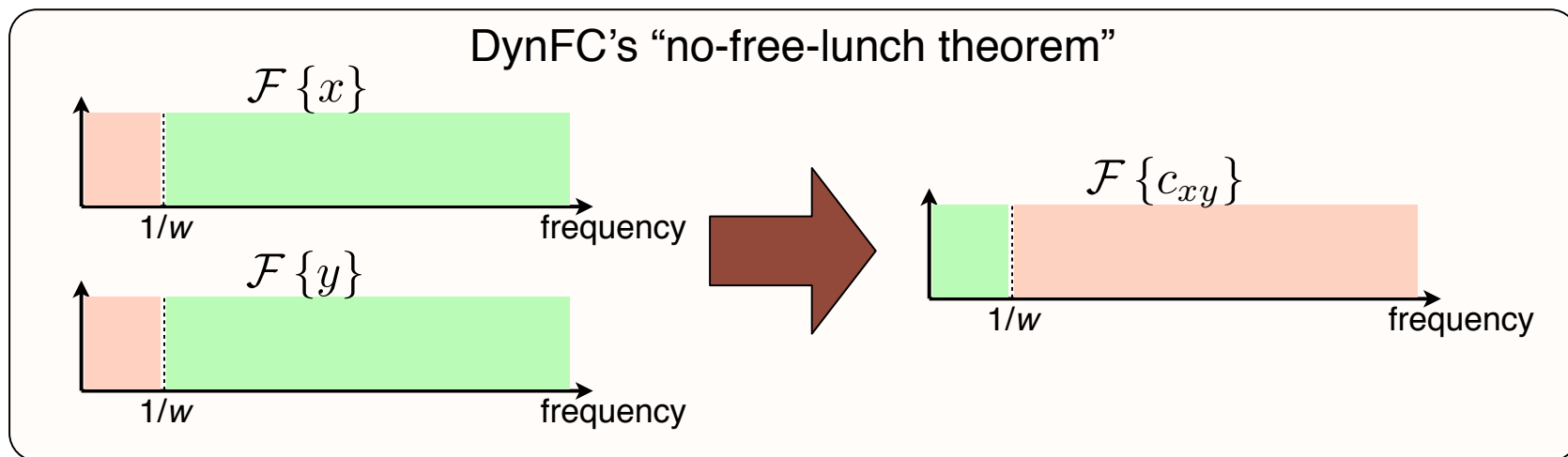
Dynamic functional connectivity

- Extract network dynamics by sliding-window pairwise correlations



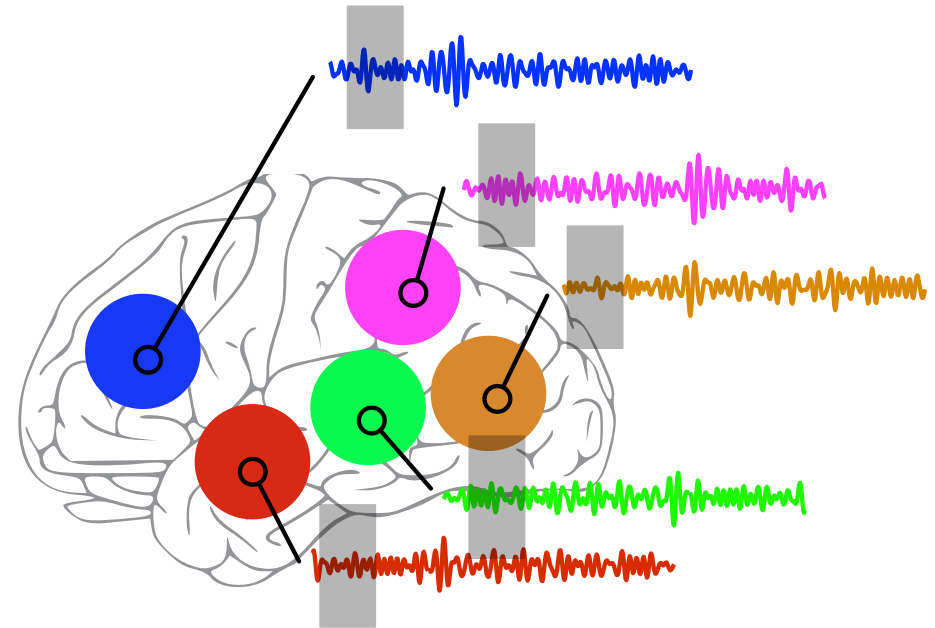
Dynamic functional connectivity

- Proper preprocessing of timecourses is required
 - To avoid aliasing artefact: high-pass filtering of input timecourses with reciprocal of window length w

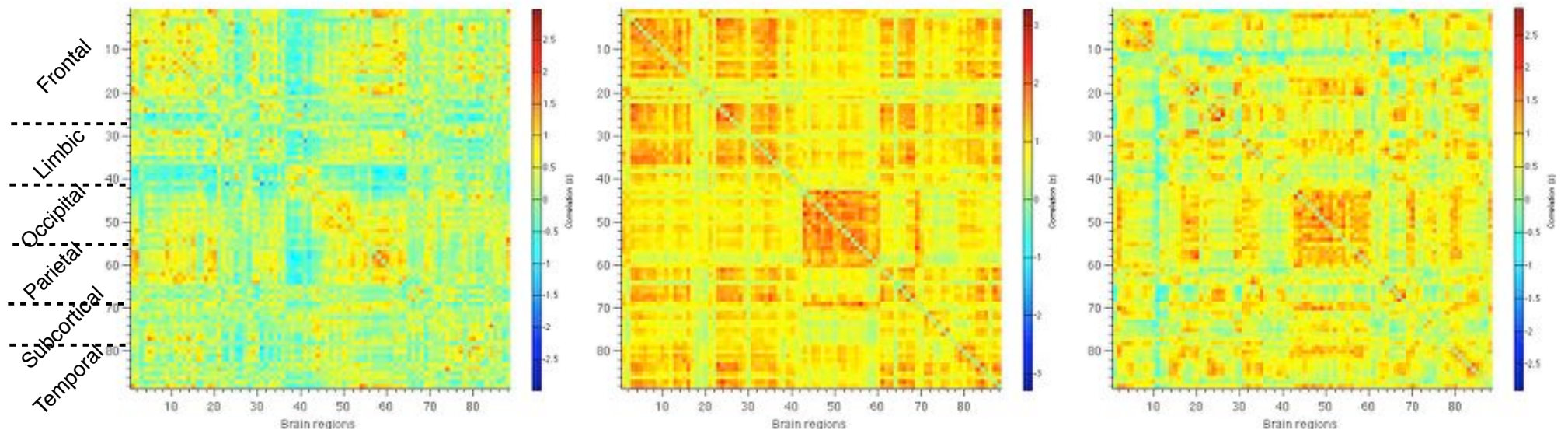


Dynamic functional connectivity

- Sliding window correlation*
 - Window length is 30 TRs, step of 2 TRs, TR=1.1sec

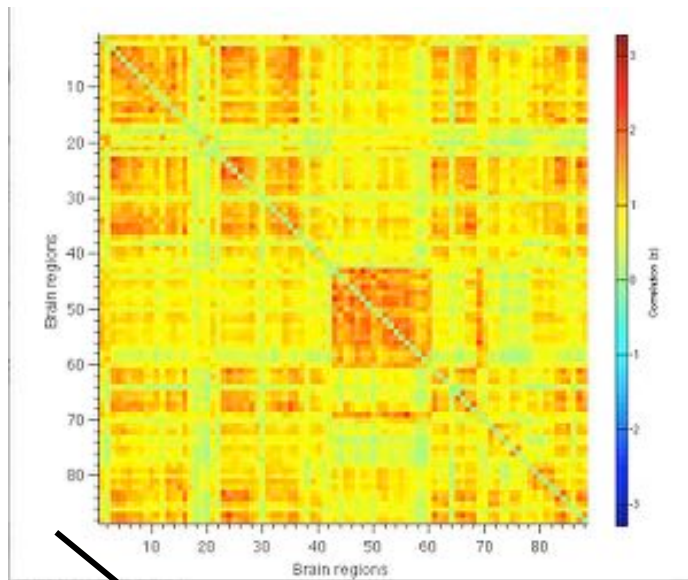


- Three healthy subjects:

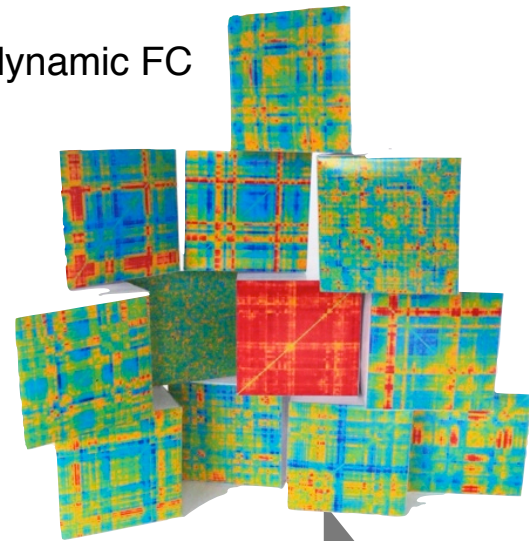


Eigenconnectivities

“Lego” of dynamic FC

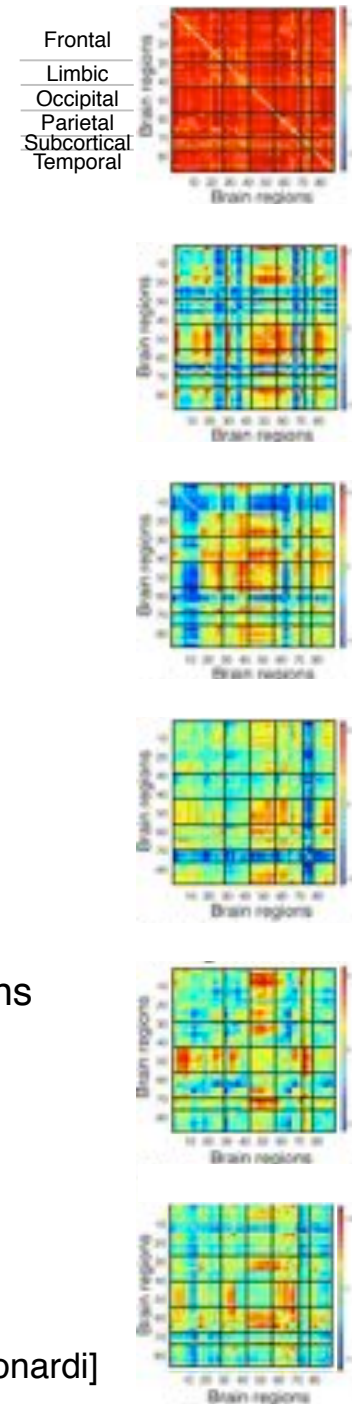


time x subjects



PCA

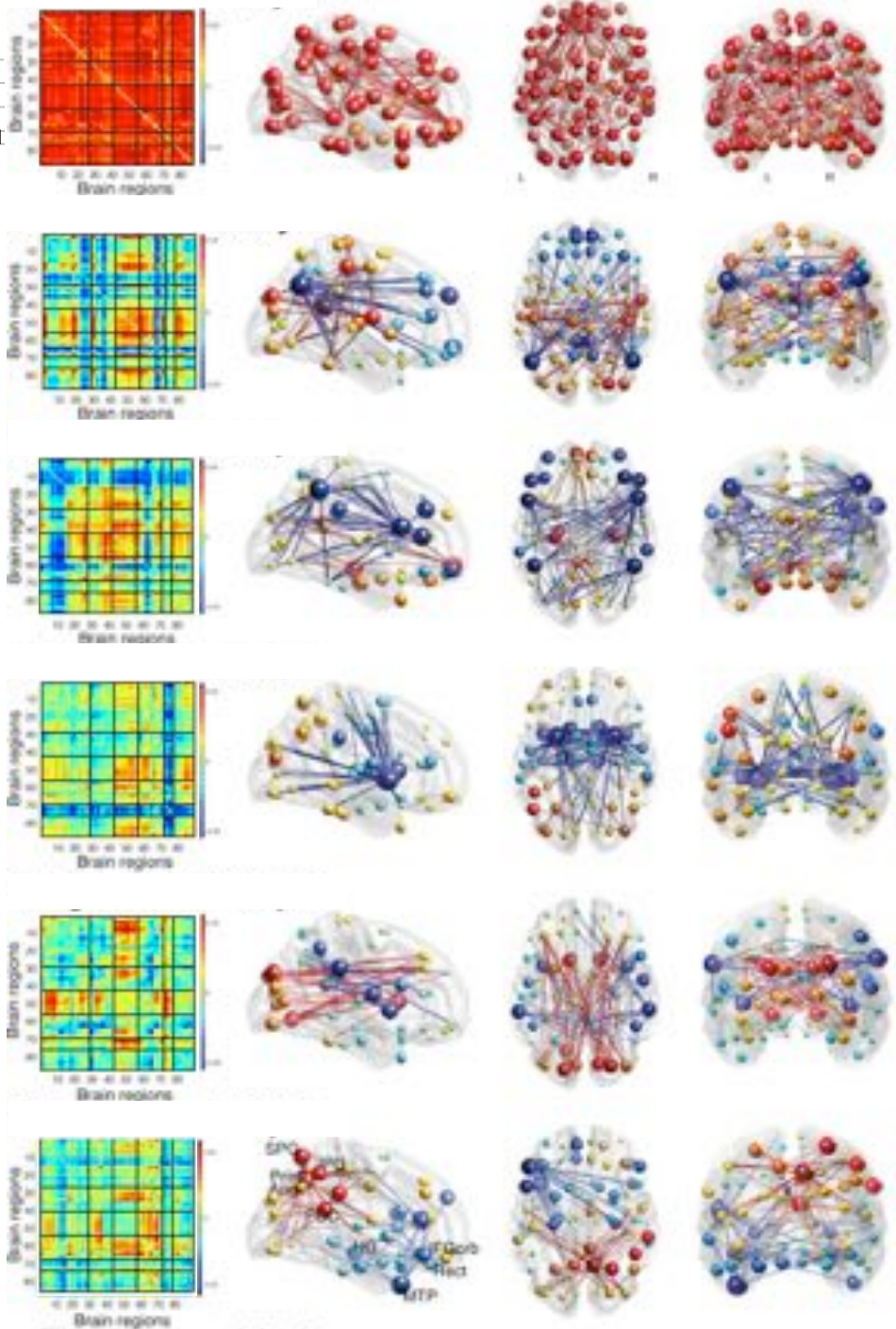
- average dynamic FC is subtracted out (i.e., driven by fluctuations of FC only!)
- optimized for explained variance
- orthogonality constraint



positive
0
negative

Eigenconnectivities

Frontal
Limbic
Occipital
Parietal
Subcortical
Temporal



global fluctuations in FC

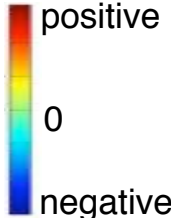
cingulate gyri, medial frontal gyri,
precuneus (default-mode network)
primary sensory in red

inferior and middle frontal gyri, inferior
parietal
(fronto-parietal)

subcortical

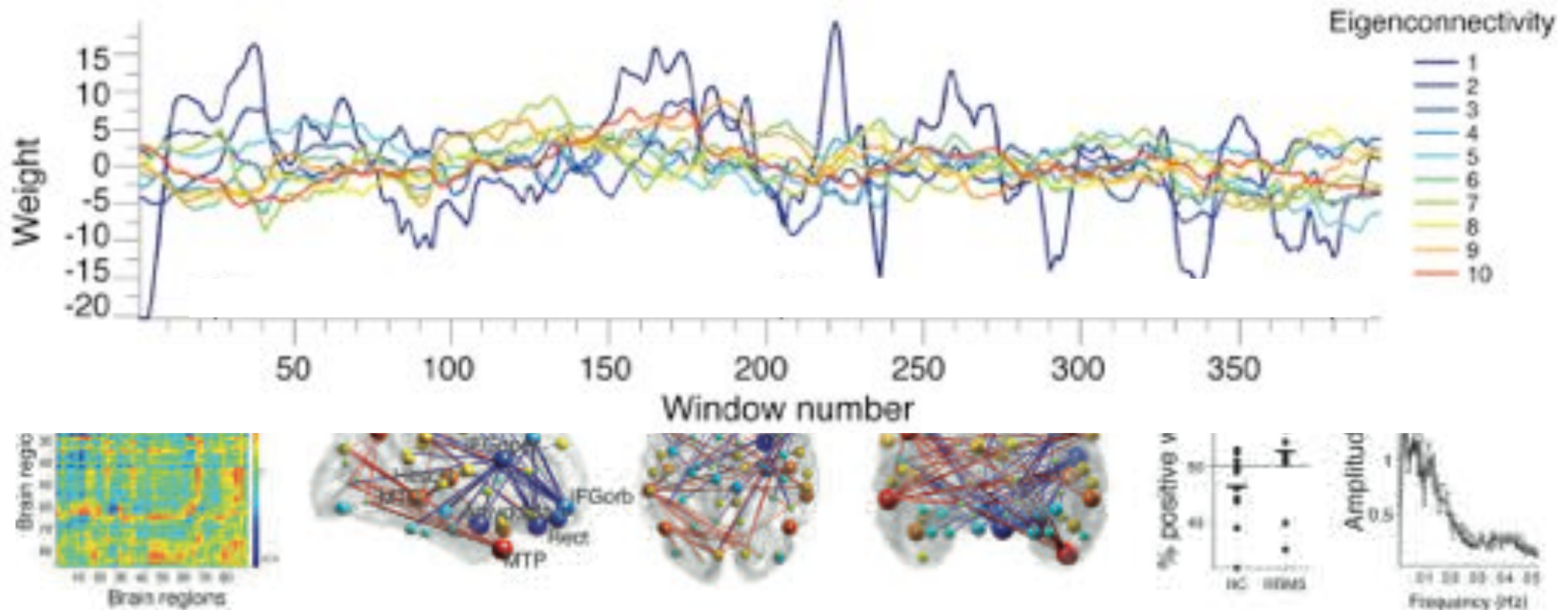
visual

posterior DMN
temporal and inferior frontal



Altered dynamics in multiple sclerosis

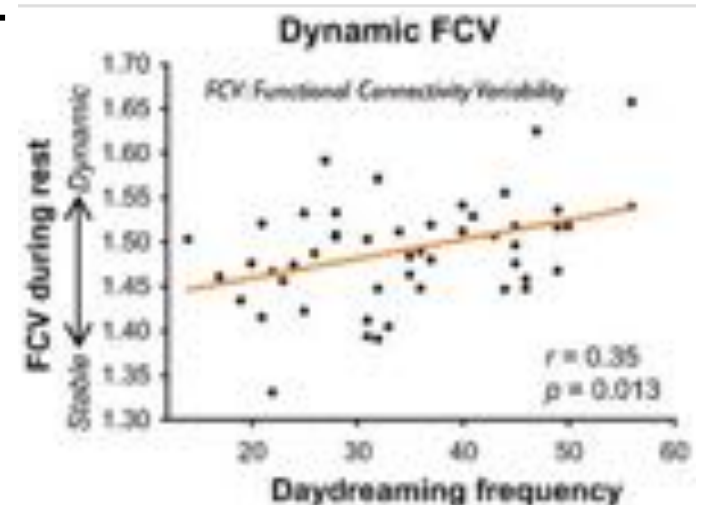
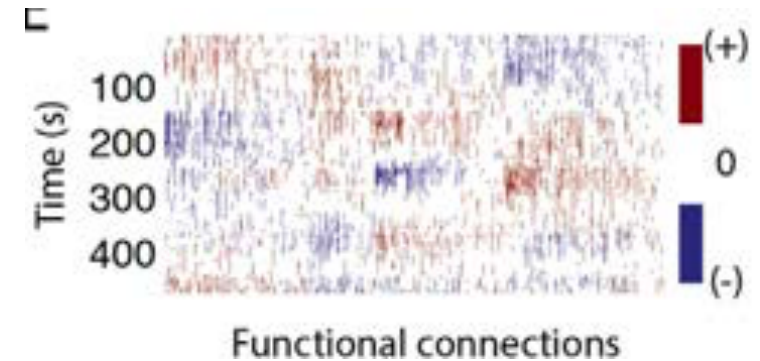
- Temporal contributions of eigenconnectivities is altered in minimally-disabled relapse-remitting multiple sclerosis patients ($F_{(10,19)}=2.6$, $p=0.005$)



- Schizophrenia, autism, temporal lobe epilepsy, ...

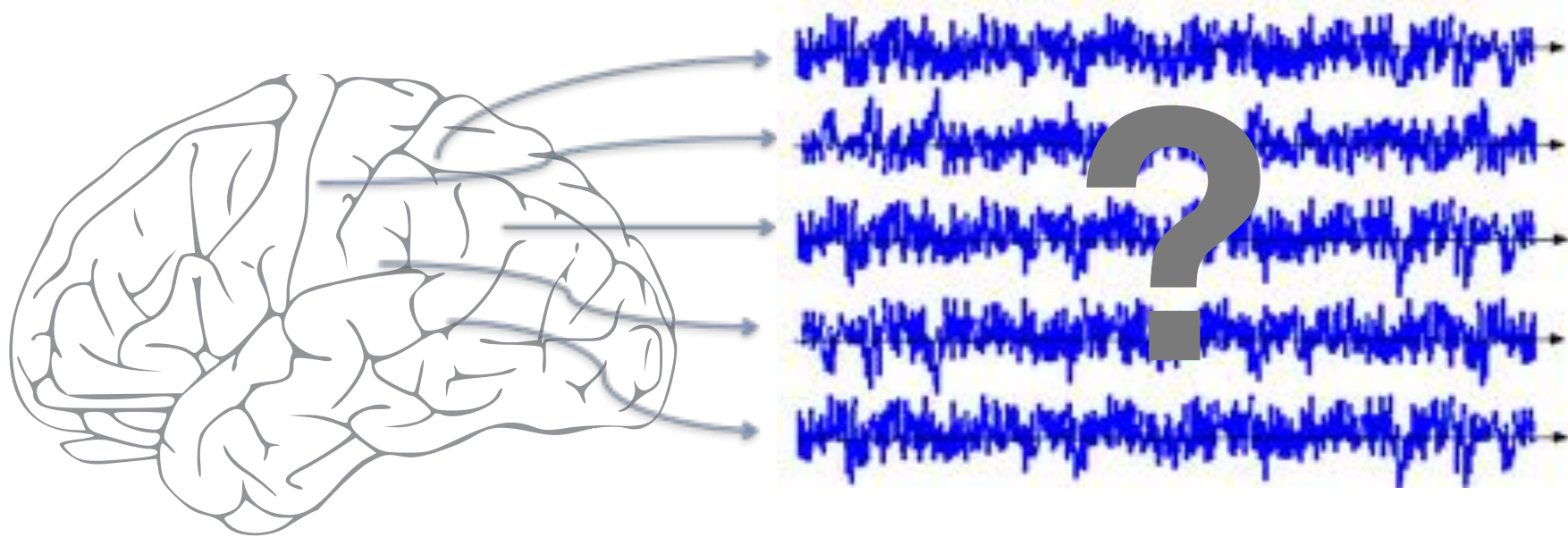
Origin and relevance of dynamic FC

- Parts of dynFC **survive** rigorous statistical testing: non-parametric randomization
 - Betzel et al, 2016; Zalesky, 2014; Keilholz et al, 2013; Huang, et al, ArXiv
- DynFC correlates with **electrical** activity
 - Thompson et al, 2013; Tagliazucchi et al, 2013; Chang et al, 2013
- DynFC varies along **demographic** variables: age, gender
 - Hutchison, Morton, 2015; Yaesoubi et al, 2015
- DynFC is modulated by changes in **arousal**: anesthesia, caffeine
 - Rack-Gomer and Liu, 2012; Barttfeld et al, 2015; Tagliazucchi et al, 2014
- DynFC correlates with **cognition**: daydreaming, cognitive flexibility
 - Kucyi, Davis, 2013; Yang et al, 2014; Cheng et al, 2016

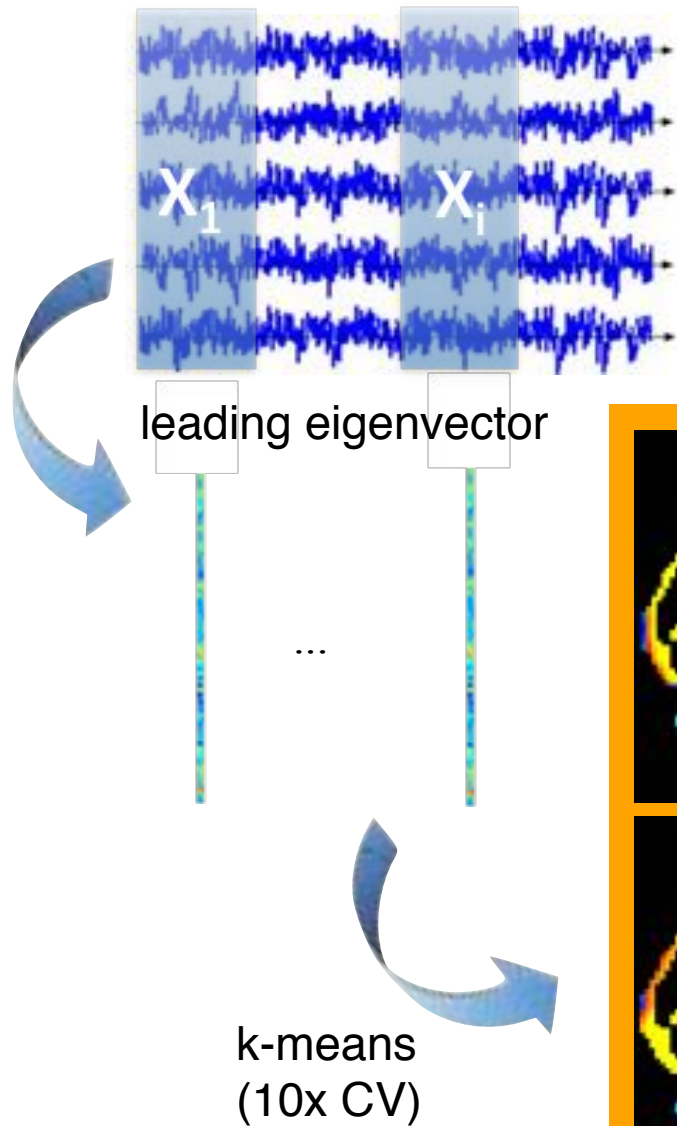


Spatially resolved dynamic FC

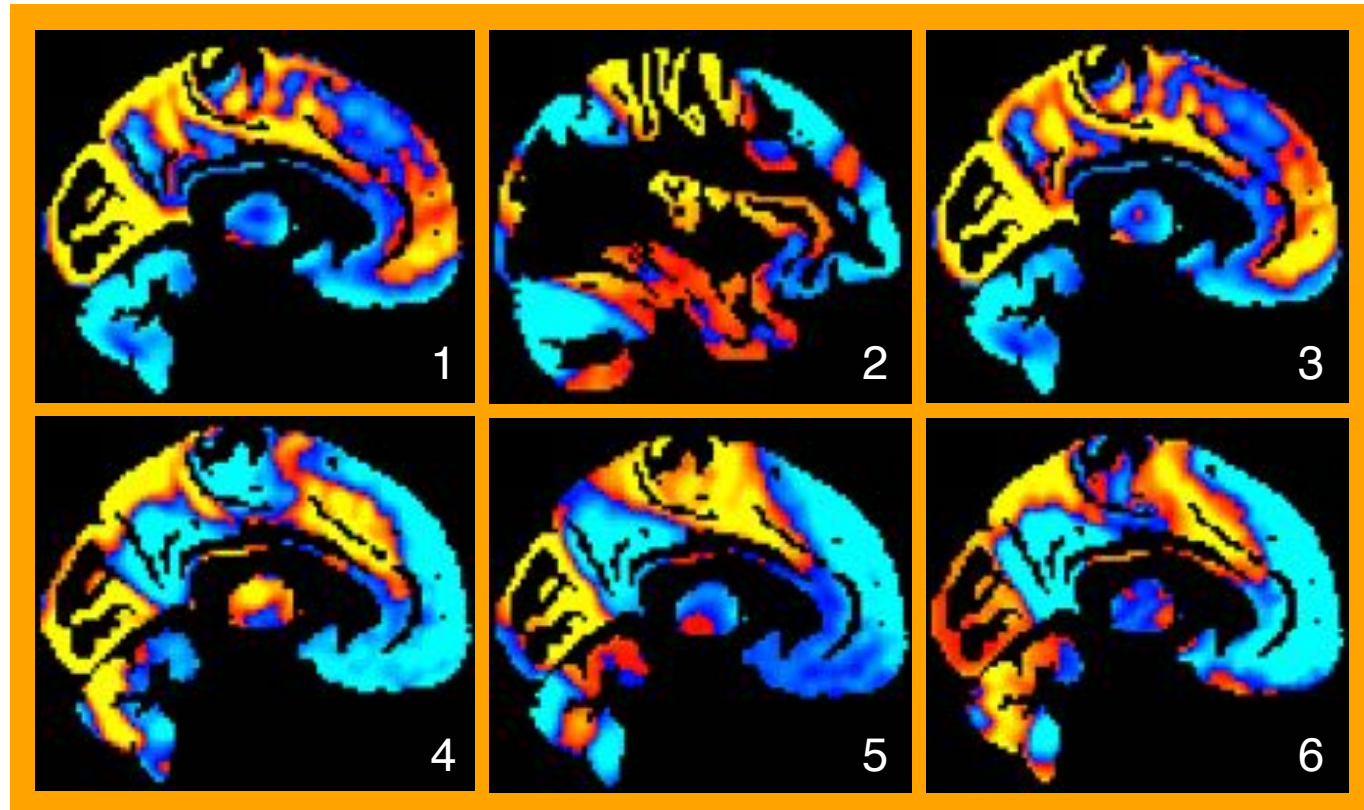
- Voxel-wise (dynamic) connectome
 - 10^5 timecourses
 - Connectivity matrix is huge!
 - Sliding-window approach... %@U\$)!
- But matrix is low-rank!
 - Rank is at most #timepoints $\ll 10^5$



Representative dominant patterns

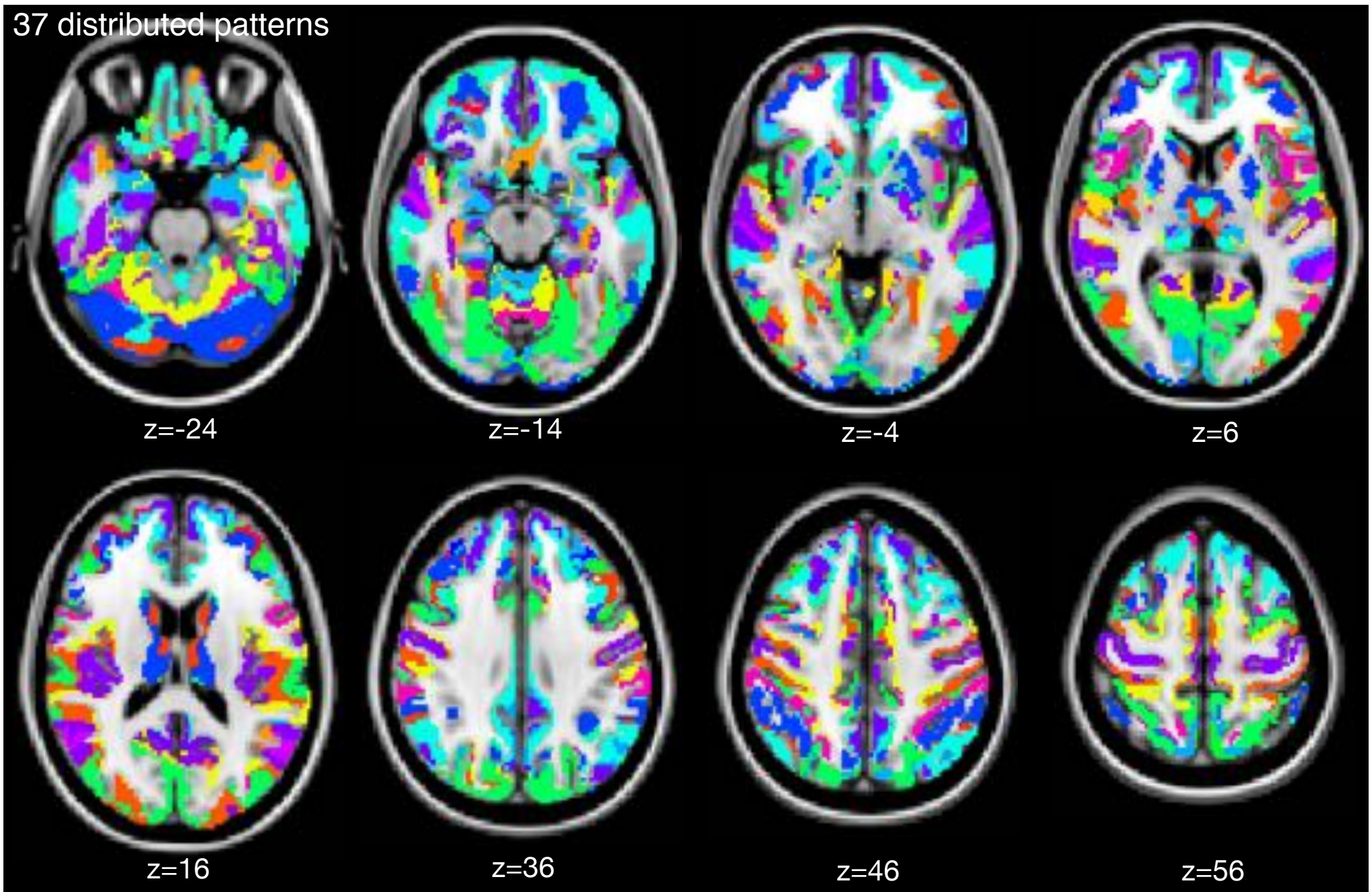


- Interplay between major networks
 - RDP5: full DMN vs sensorimotor
 - RDP4,6: segregation of DMN in dorsal/ventral parts



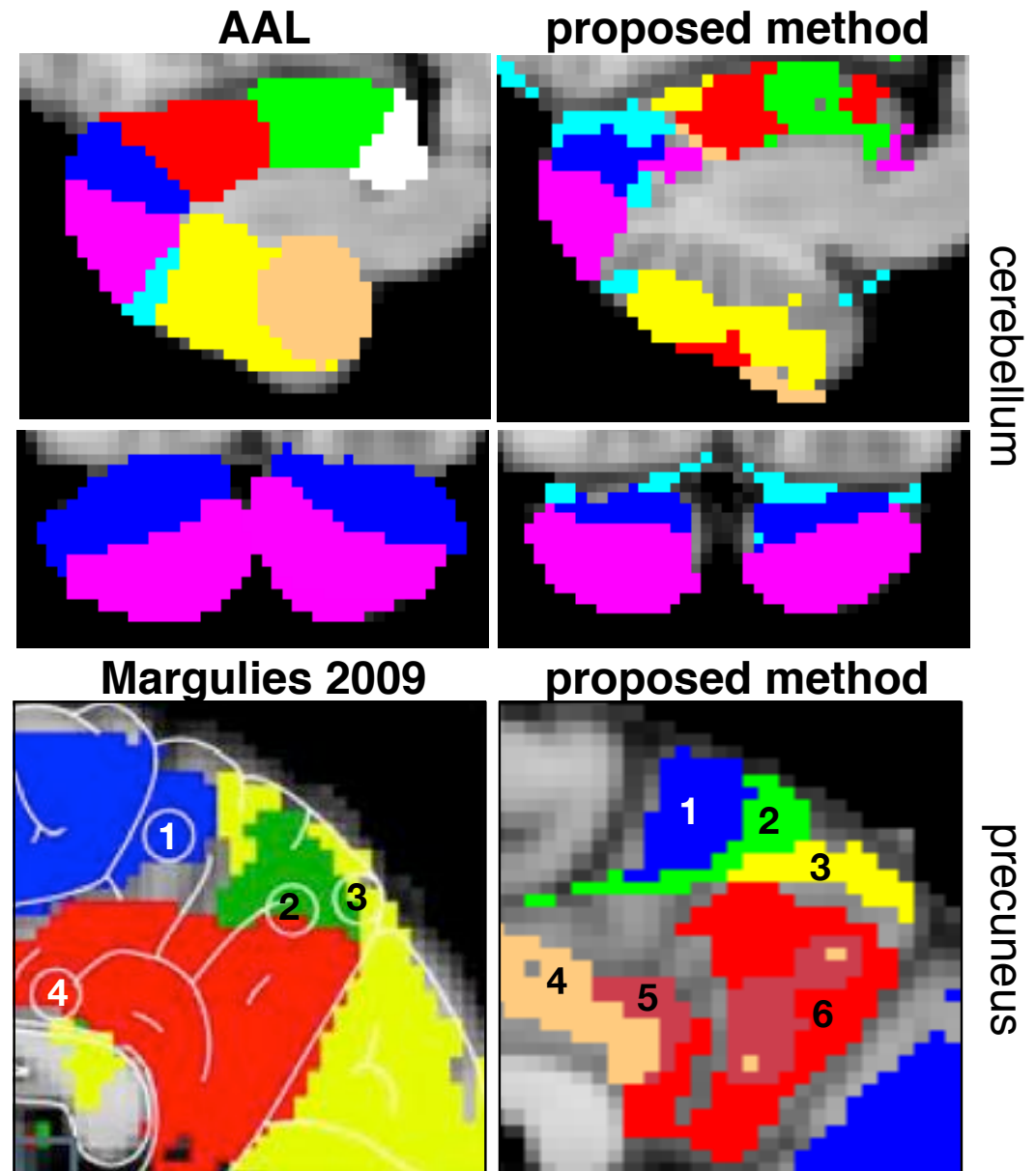
Dynamic FC voxel-wise parcellation

37 distributed patterns



Dynamic FC voxel-wise parcellation

- Split into contiguous regions
 - Leads to 449 parcels
- Fluctuations of dynamic FC are meaningful in terms of long-range and short-range interactions
- Extensions
 - Higher rank
 - Clustering
 - ...

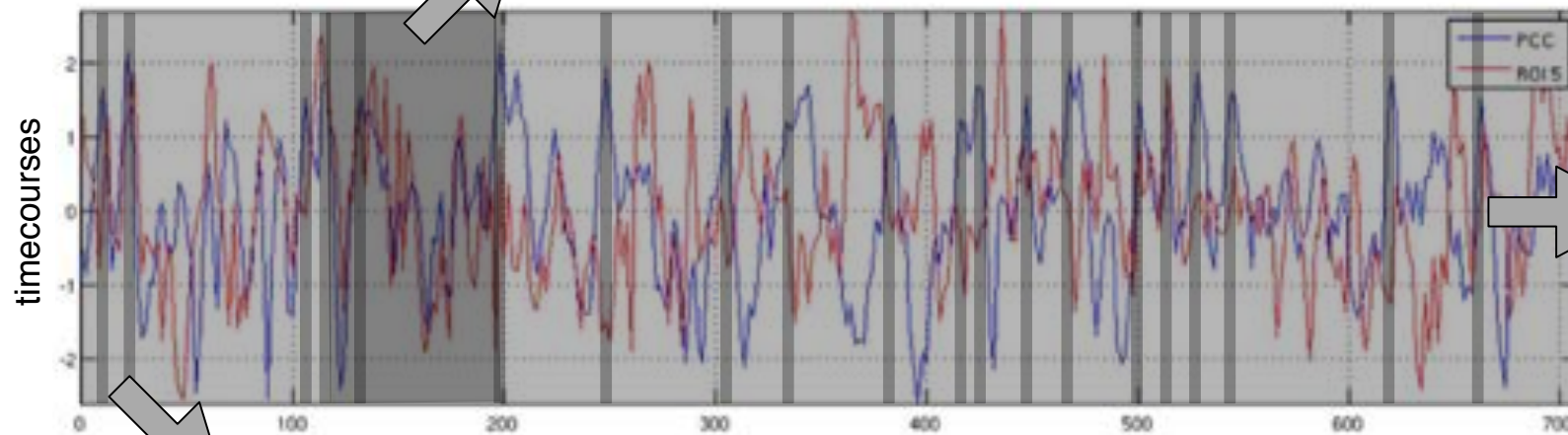


Better capturing of dynamics



windowed correlation is temporally restricted, but suffers from low SNR and potential aliasing

[Chang and Glover, 2010; Leonardi, VDV, 2013, 2014]



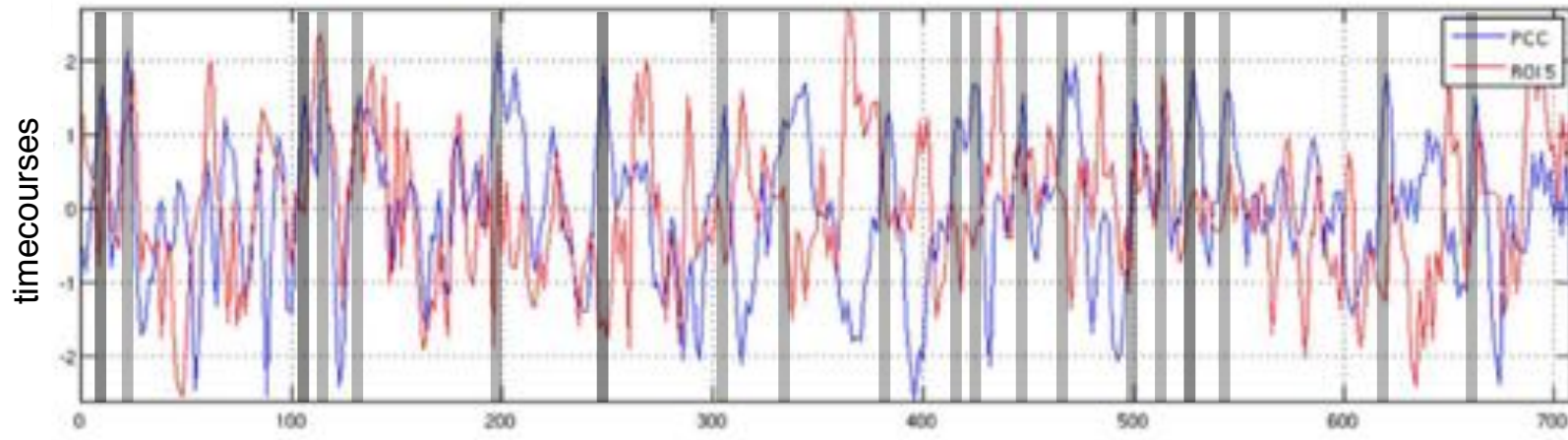
statistical dependencies characterize “average” behavior during resting state



can we extract ‘key points’ that capture the important events?

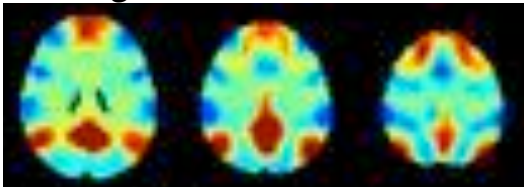
Co-activation patterns (CAPs)

- Selection of timepoints with extreme values of seed region (point process model)

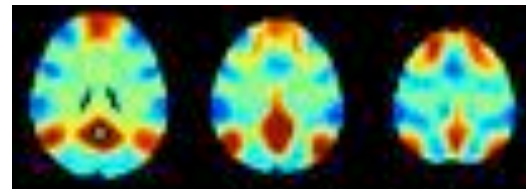


- Averaging is proxy of seed connectivity:

average over all selected frames:

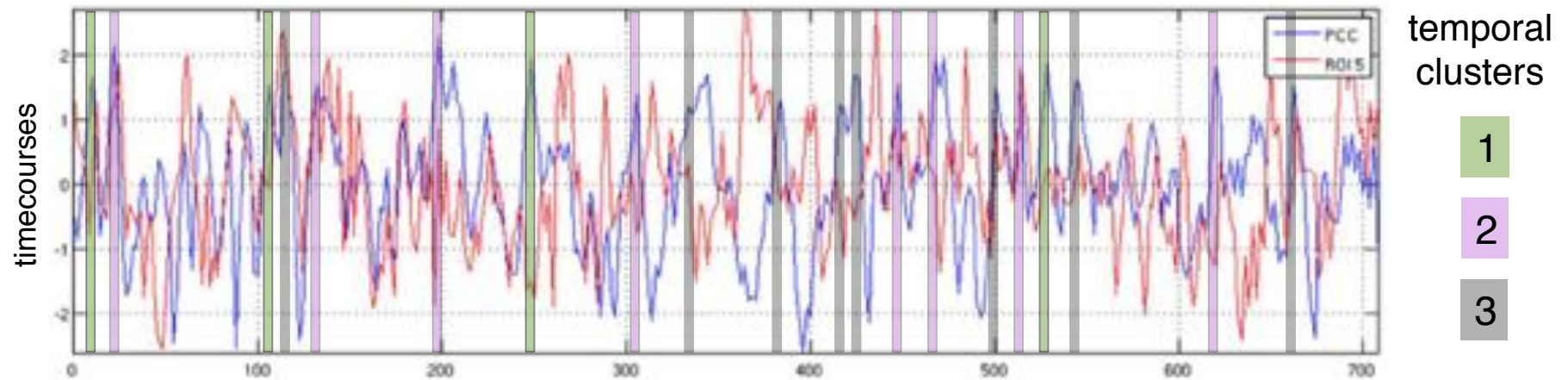


seed-based correlation:

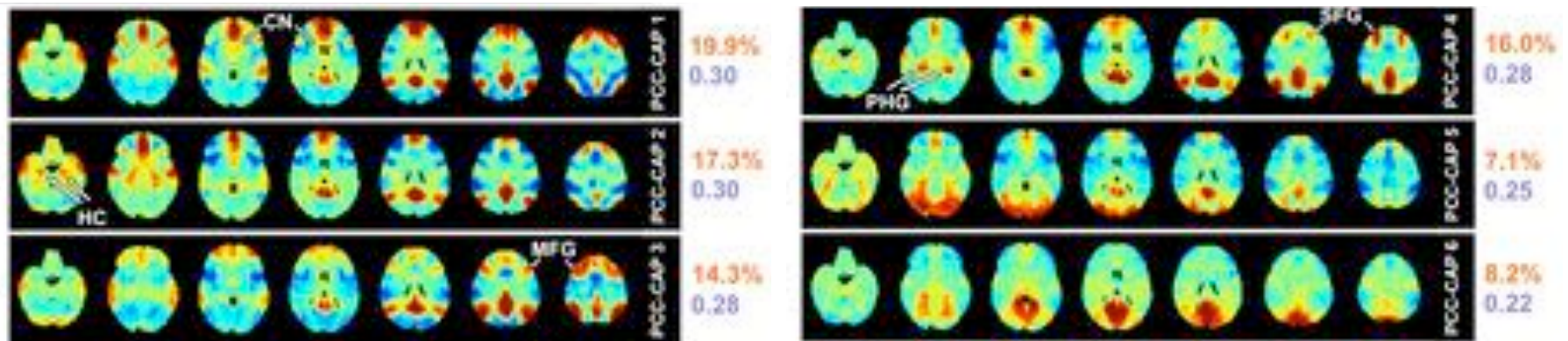


Co-activation patterns (CAPs)

- Temporal clustering of selected frames

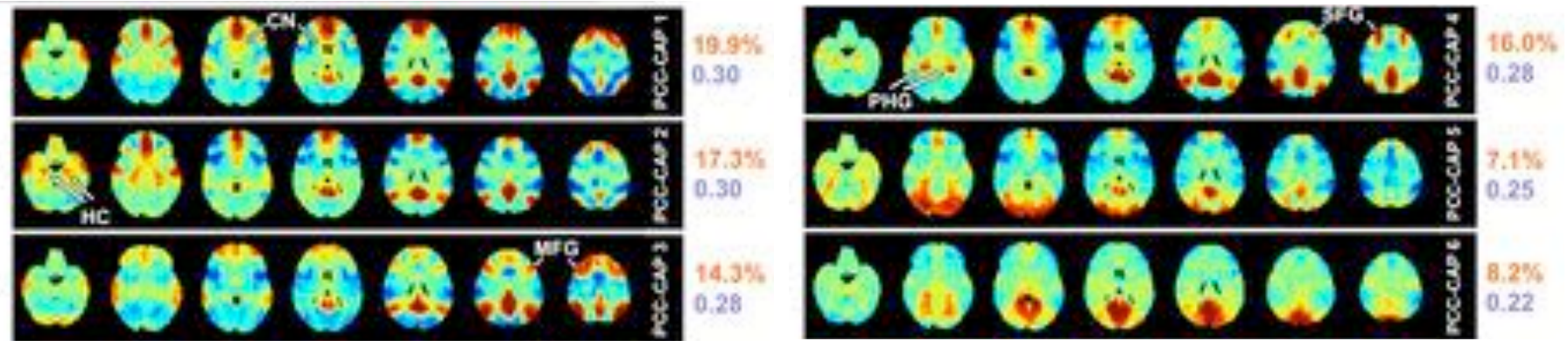


- Averaging of spatial activity patterns for each temporal cluster leads to “co-activation patterns” (CAPs)



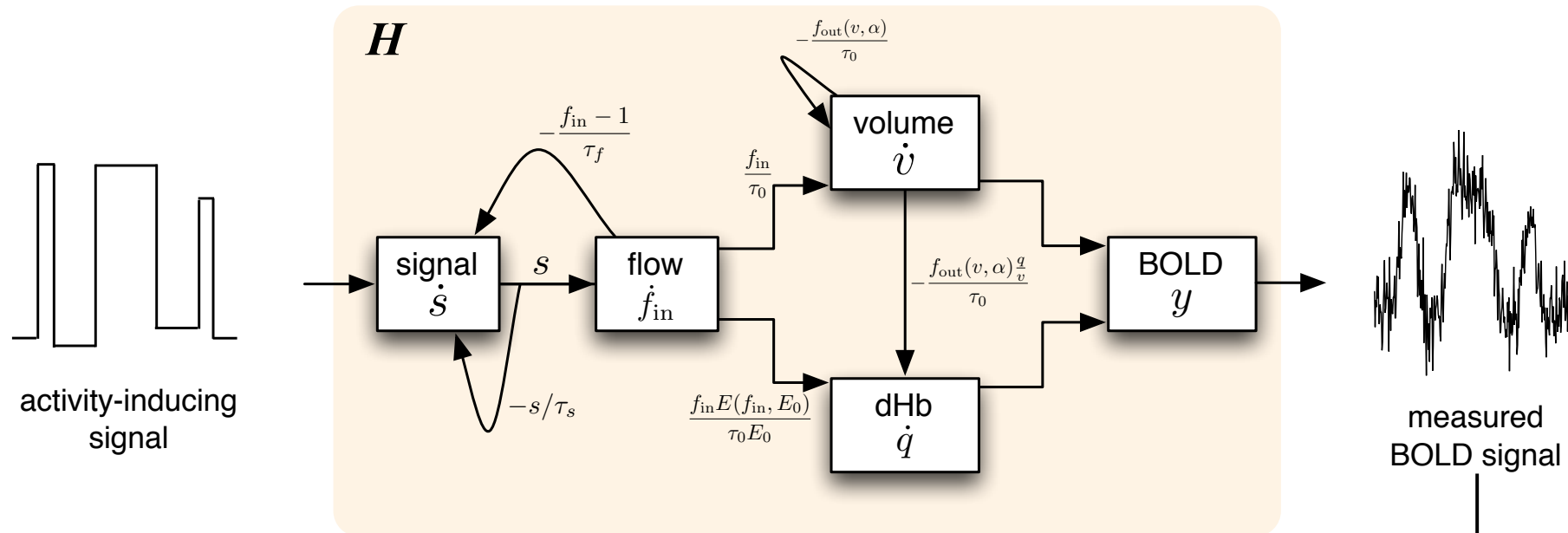
Co-activation patterns (CAPs)

- Averaging of spatial activity patterns for each temporal cluster leads to “co-activation patterns” (CAPs)



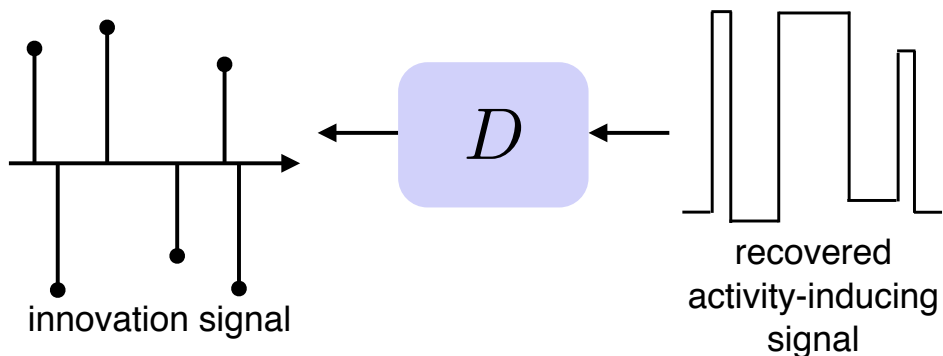
- Temporal clustering allows CAPs to have spatial overlap, but it does not disentangle temporal overlap:
 - #CAPs suffers from combinatorial explosion:
 - N “fundamental” networks that can temporally overlap K at a time would lead to $\sim N^K$ CAPs
 - CAPs can be contaminated by non-seed related activity

Total activation regularized deconvolution of fMRI BOLD timeseries



■ Total activation regularization:

$$\tilde{\mathbf{x}} = \arg \min_{\mathbf{x}} \underbrace{\frac{1}{2} \|\mathbf{y} - \mathbf{x}\|_2^2}_{\text{data fitness}} + \lambda \underbrace{\|DH^{-1}\{\mathbf{x}\}\|_1}_{\text{regularization}}$$



H^{-1} First-order Volterra kernel

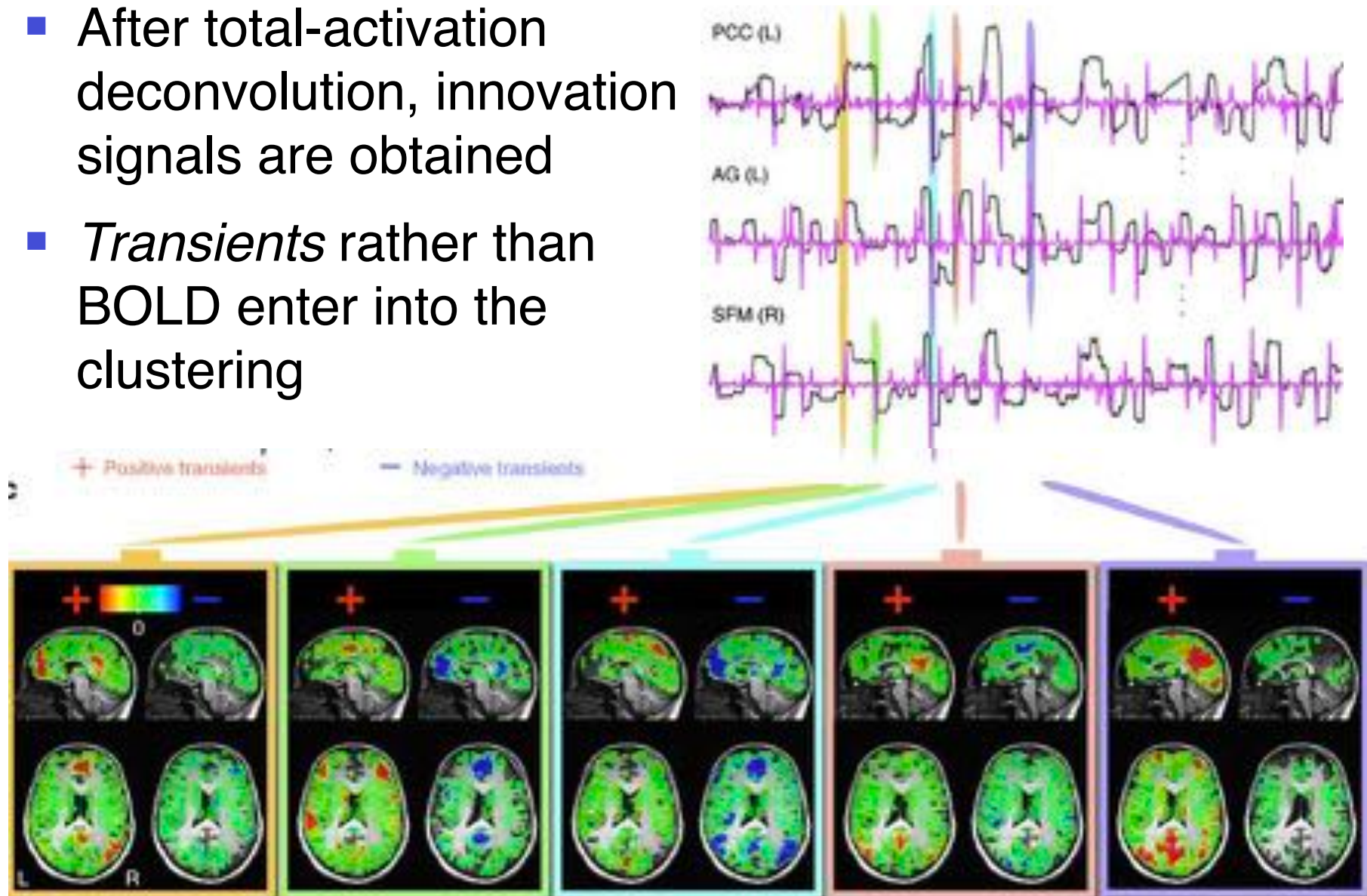
$$H^{-1} = \frac{\prod_{i=1}^4 (D - \alpha_i I)}{D - \gamma I}$$

$$\vec{\alpha} = \left\{ -\frac{1}{\tau_0}, -\frac{1}{\alpha\tau_0}, -\frac{1}{2\tau_s} \left(1 \pm j\sqrt{\frac{4\tau_s^2}{\tau_f} - 1} \right) \right\},$$

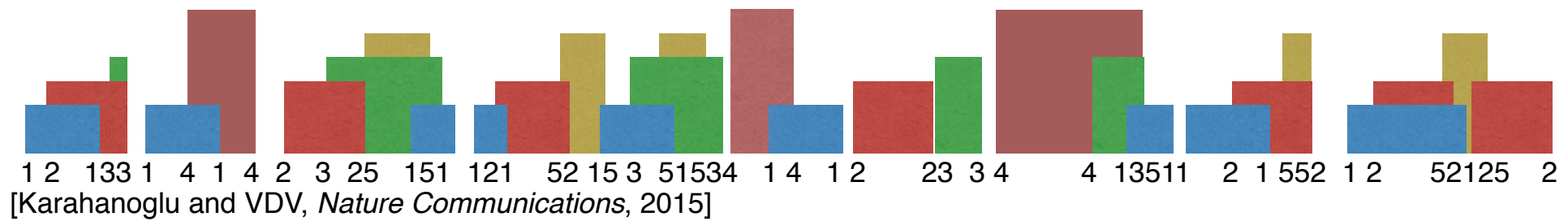
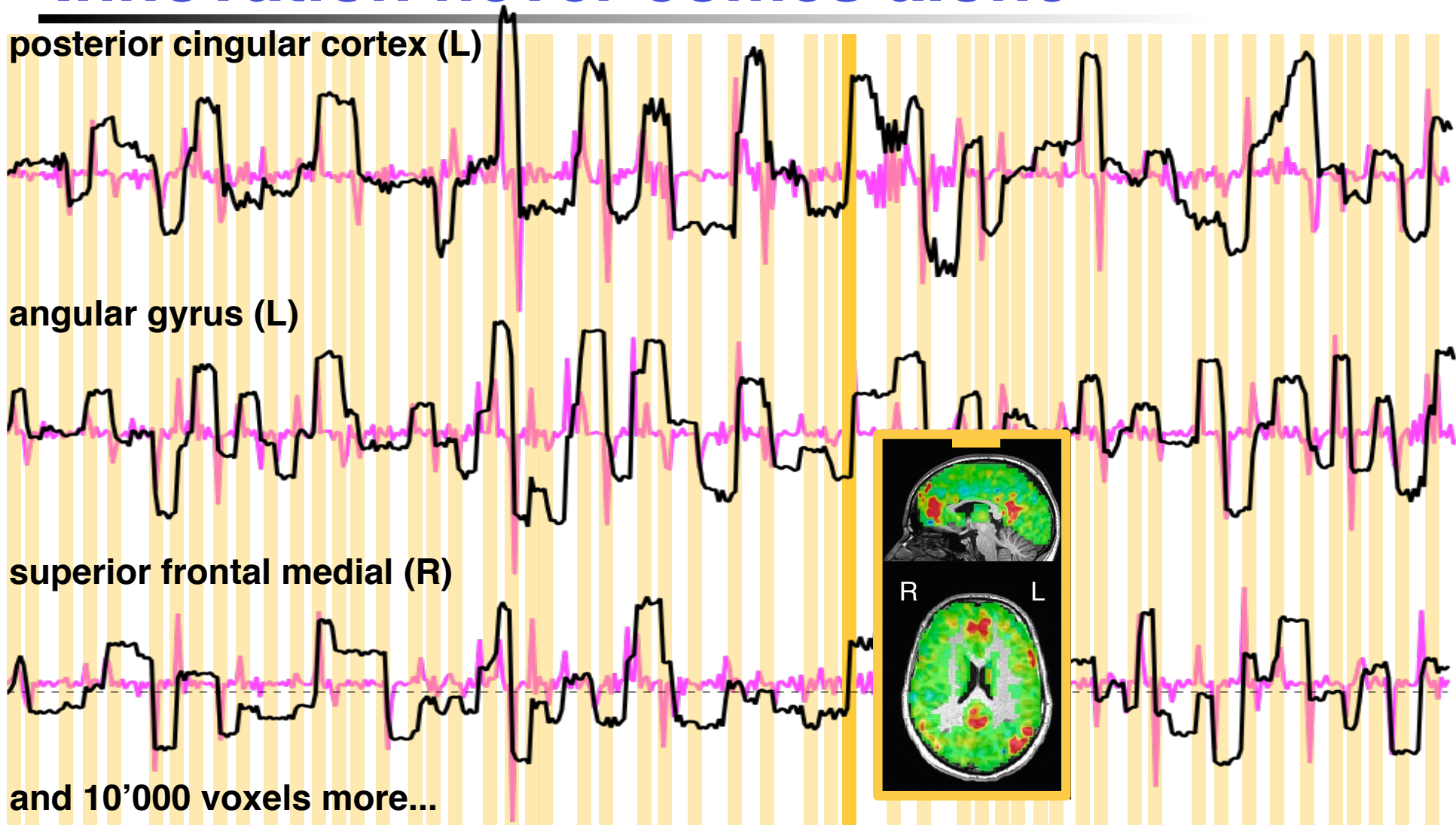
$$\vec{\gamma} = \left\{ \frac{(k_1 + k_2)\left(\frac{1-\alpha}{\alpha\tau_0} - \frac{c}{\alpha}\right) - (k_3 - k_2)\frac{1}{\tau_0}}{-(k_1 + k_2)c\tau_0 - k_3 + k_2} \right\}.$$

Innovation-driven co-activation patterns

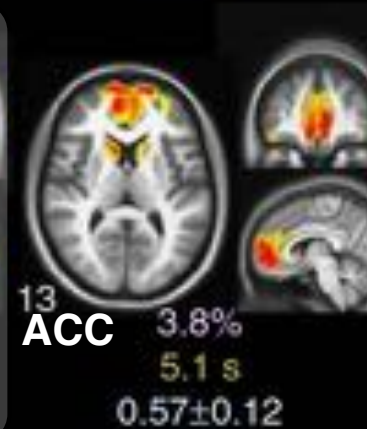
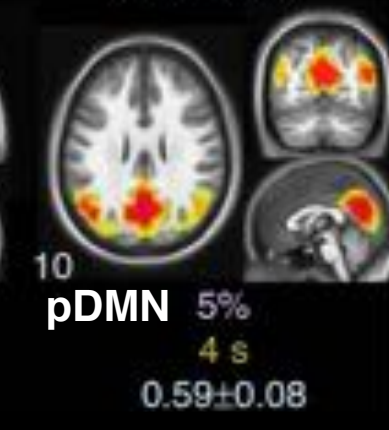
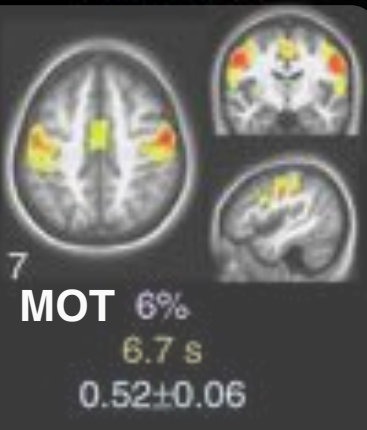
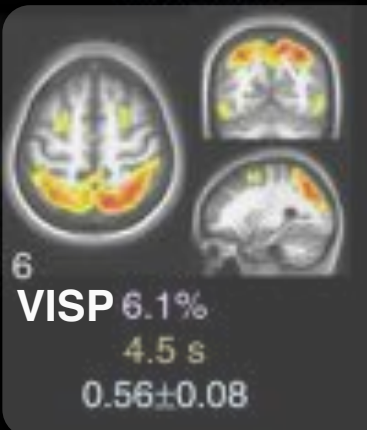
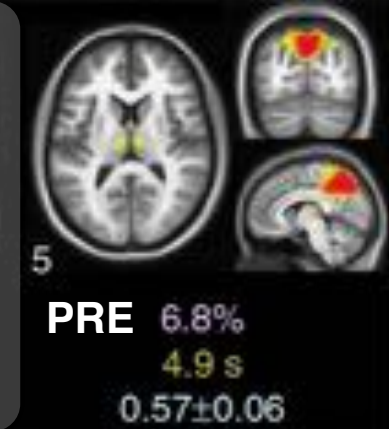
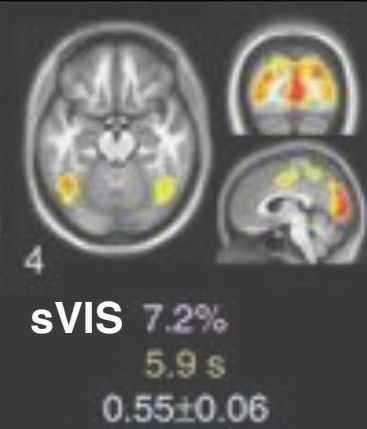
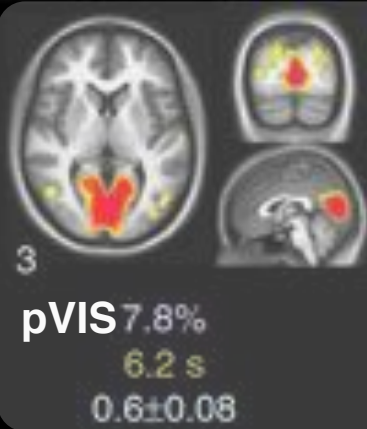
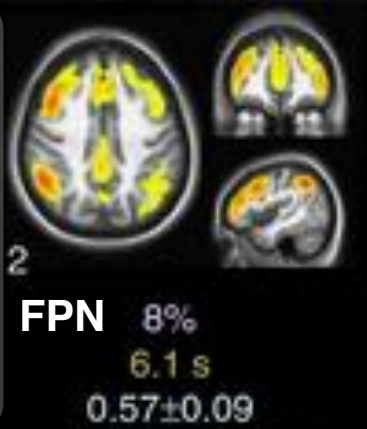
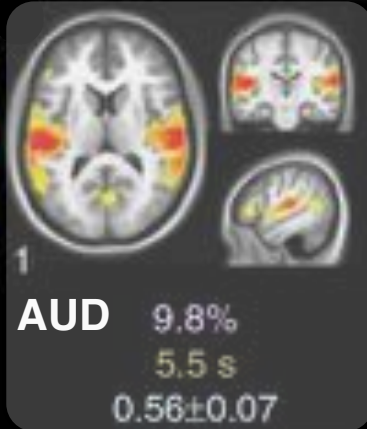
- After total-activation deconvolution, innovation signals are obtained
- *Transients* rather than BOLD enter into the clustering



Innovation never comes alone



Brain's repertoire of functional networks

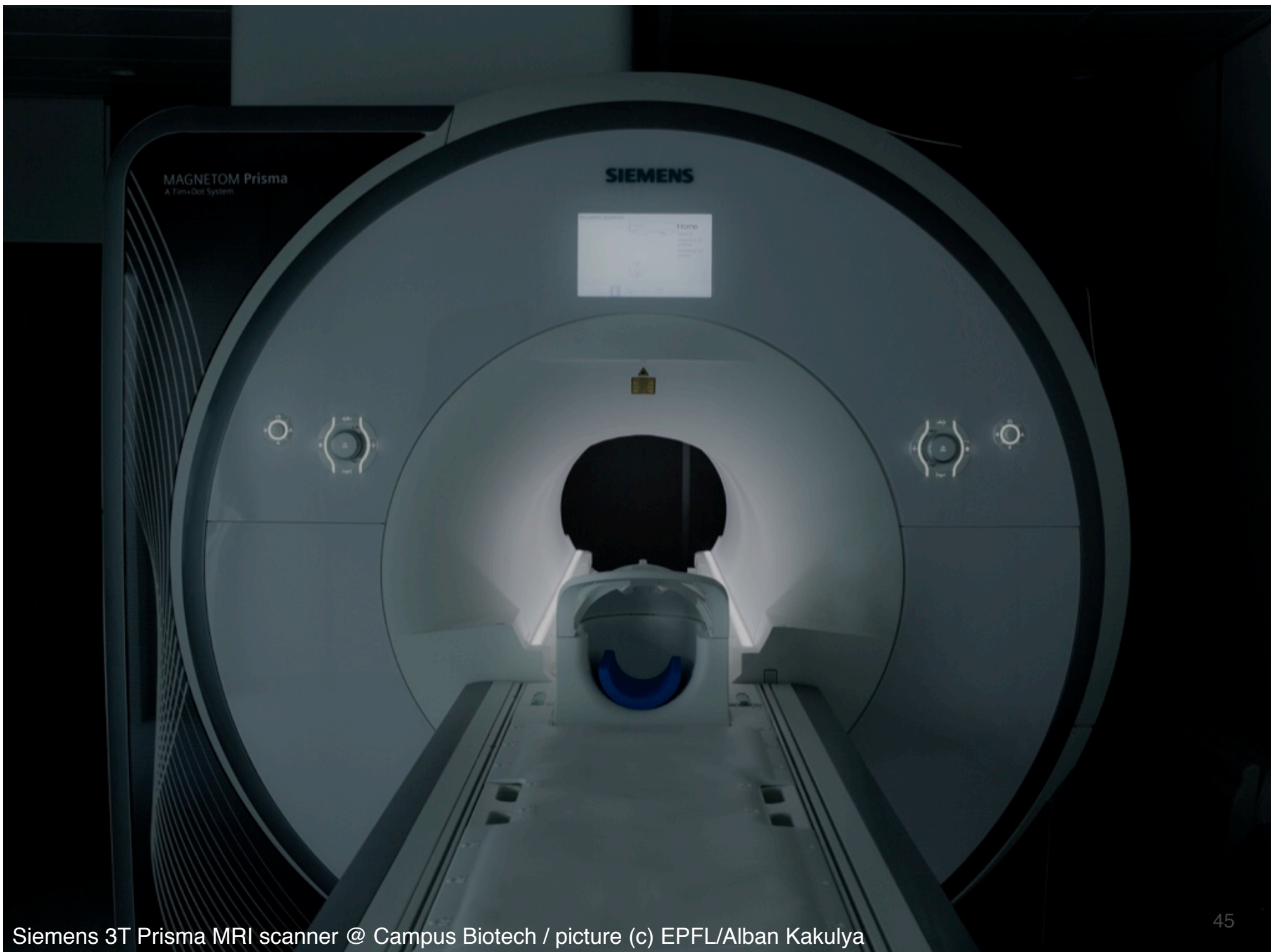


iCAPs ordered in terms of occurrence

sensory components

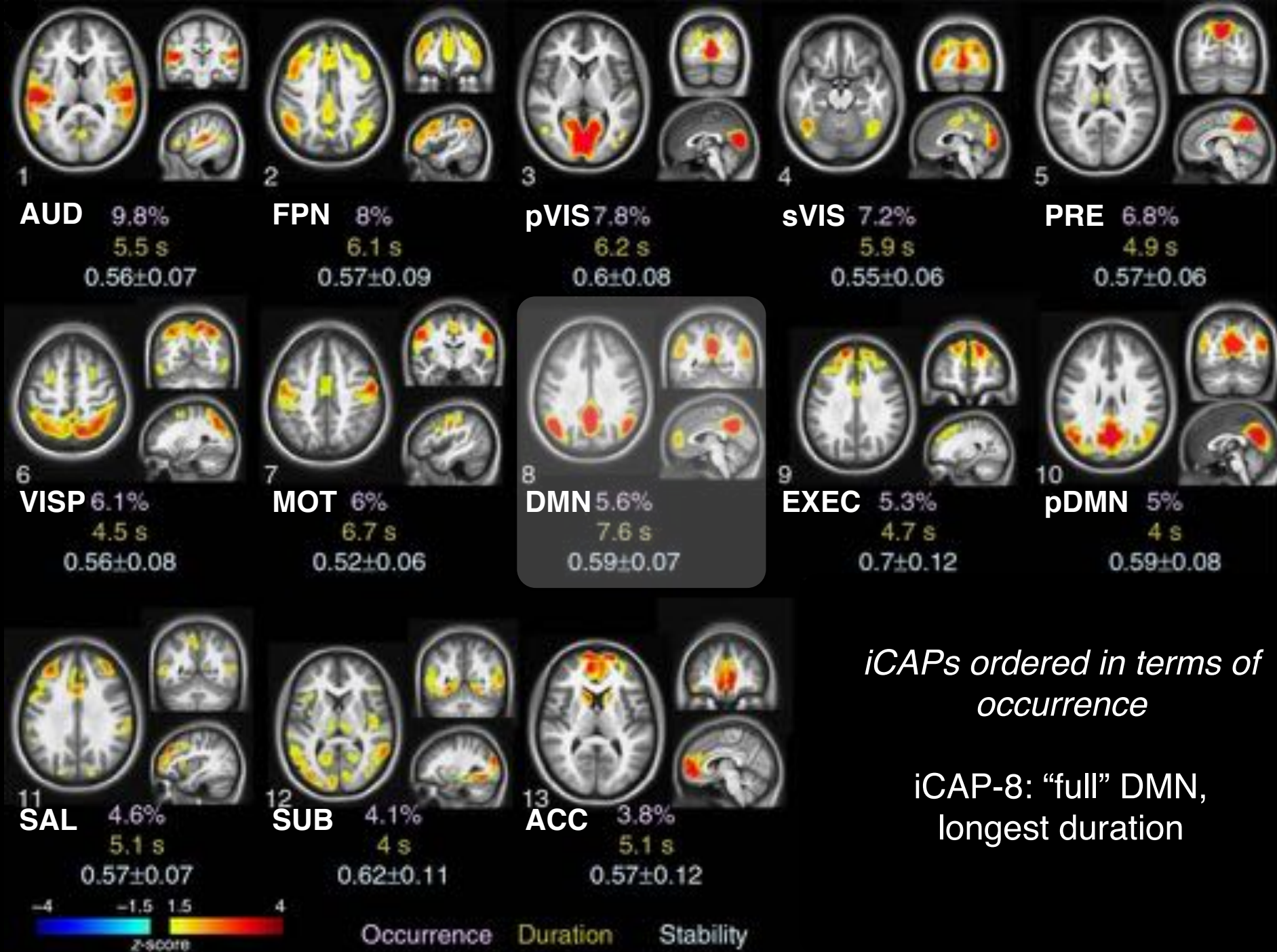


Occurrence Duration Stability



Siemens 3T Prisma MRI scanner @ Campus Biotech / picture (c) EPFL/Alban Kakulya

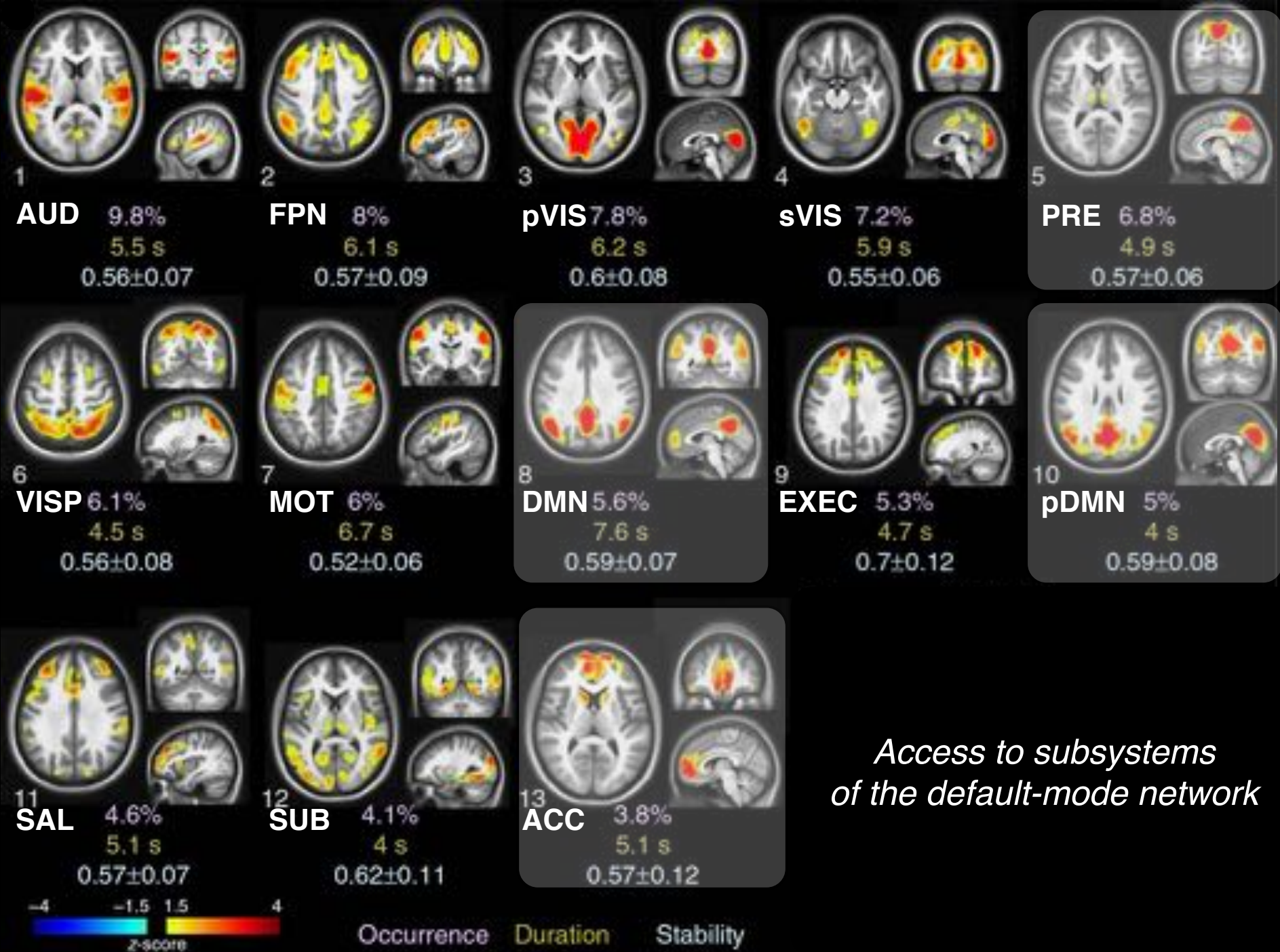
Brain's repertoire of functional networks



iCAPs ordered in terms of occurrence

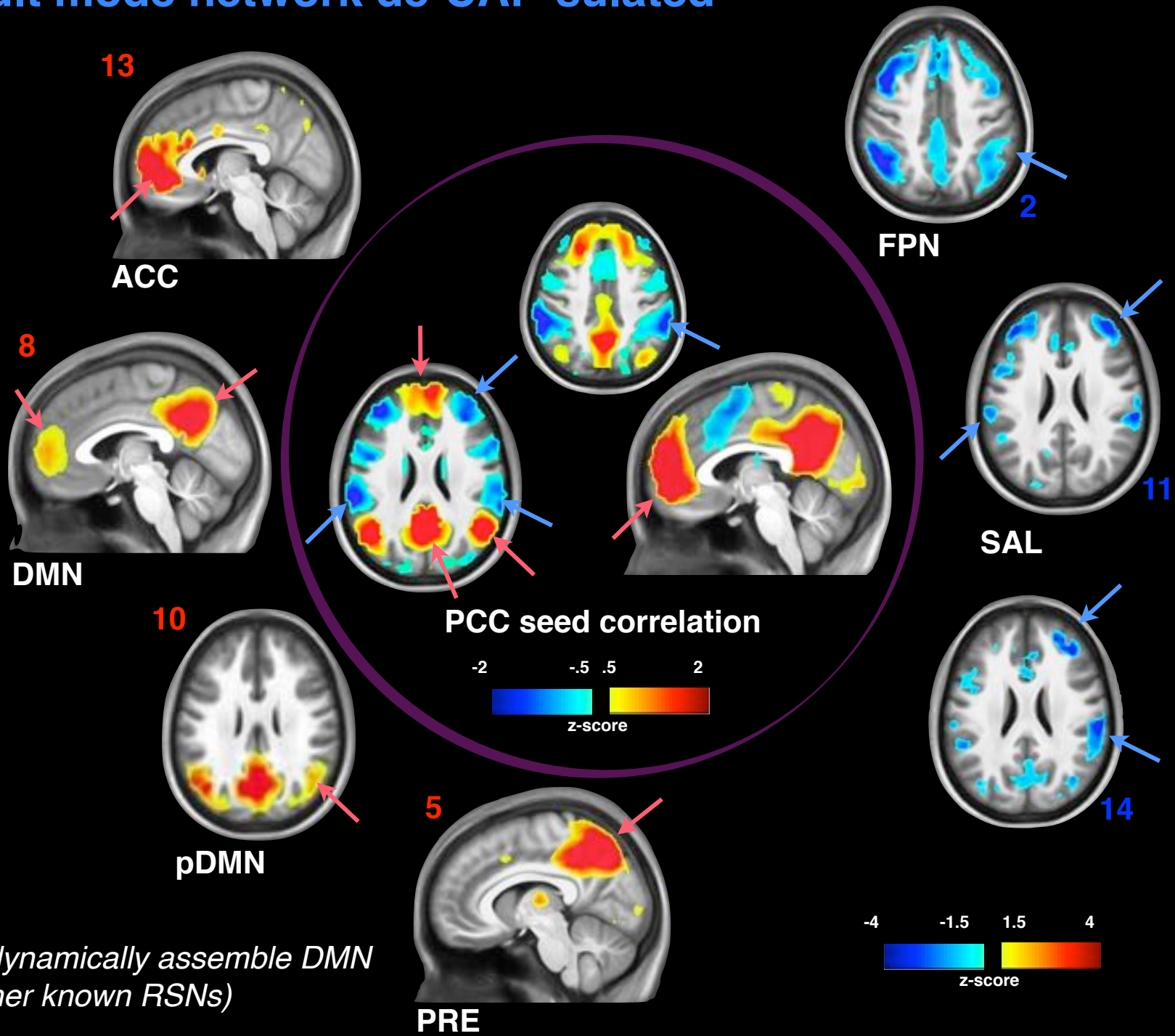
iCAP-8: "full" DMN, longest duration

Brain's repertoire of functional networks



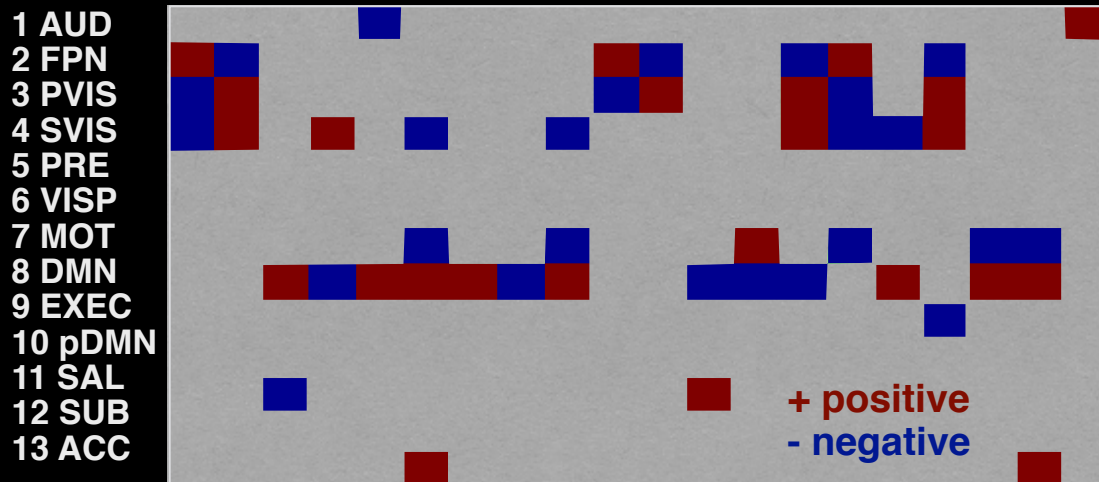
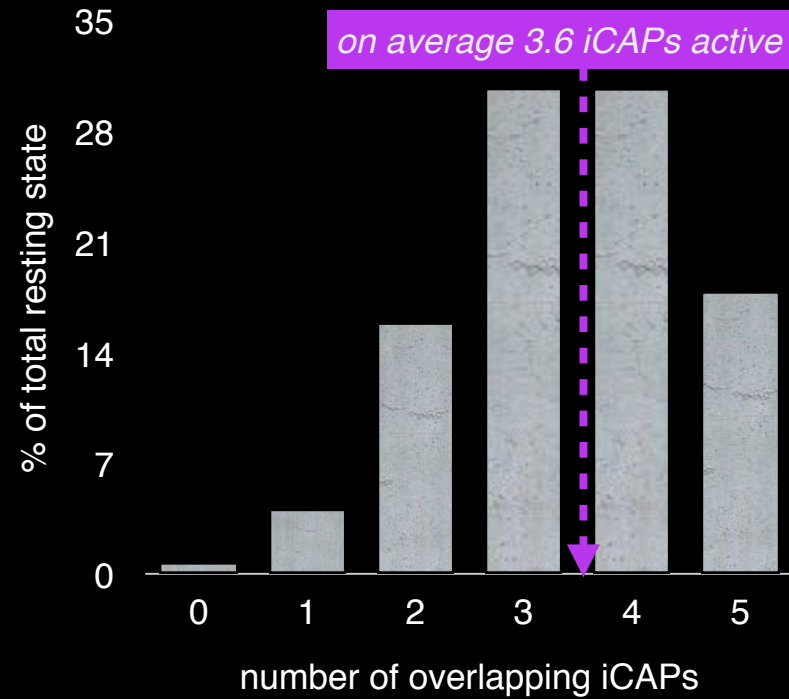
Access to subsystems of the default-mode network

Default mode network de-CAP-sulated

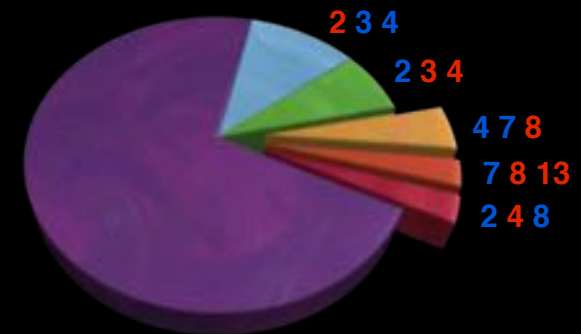


*iCAPs dynamically assemble DMN
(and other known RSNs)*

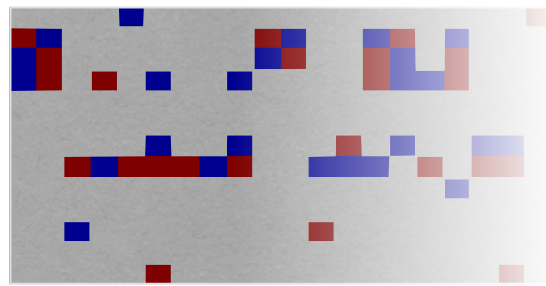
Temporal overlap of iCAPs



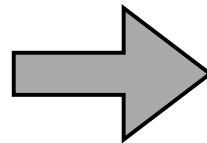
20 most frequent combinations



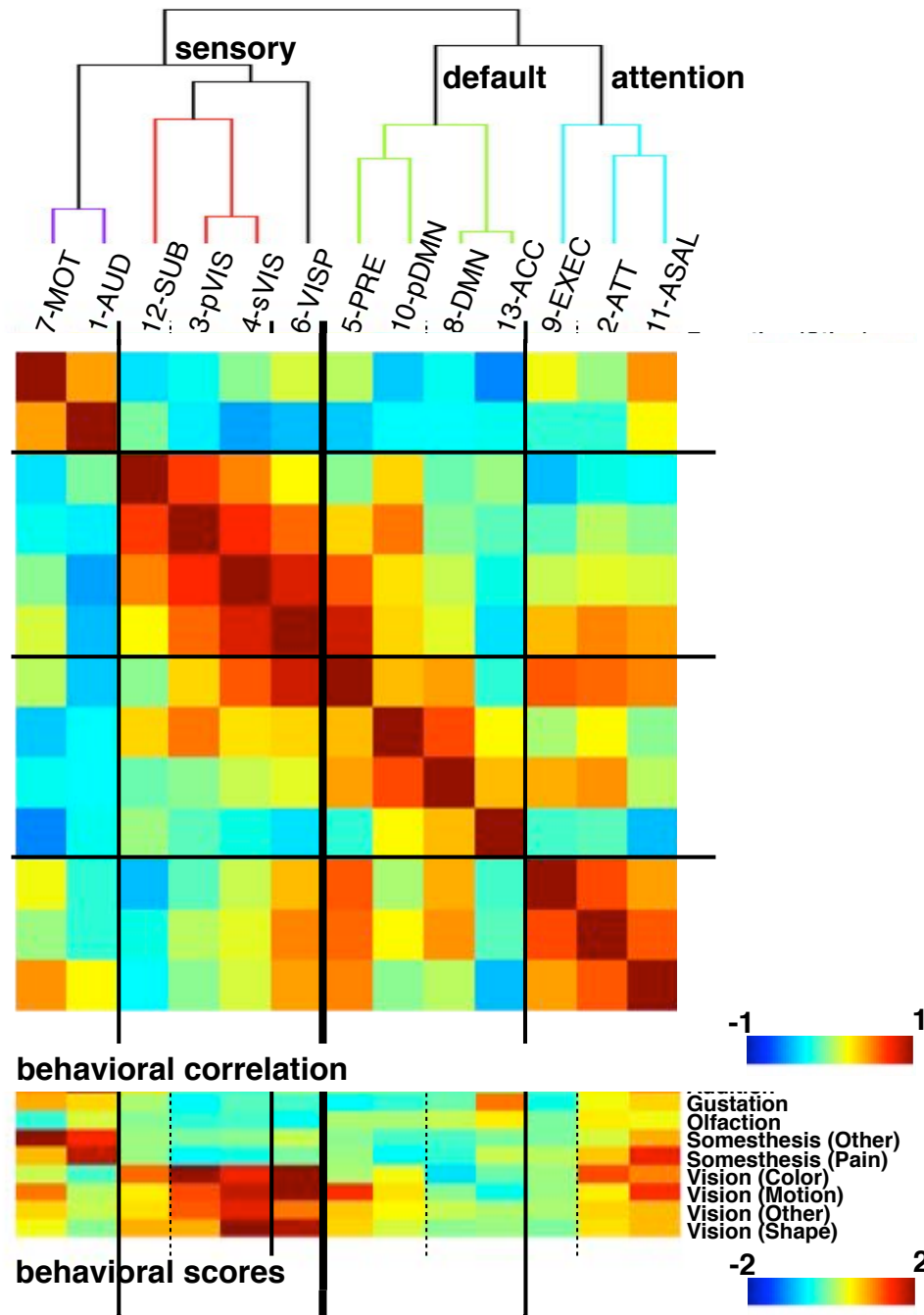
Relationship between iCAPs and behavior



2098 iCAPs combinations



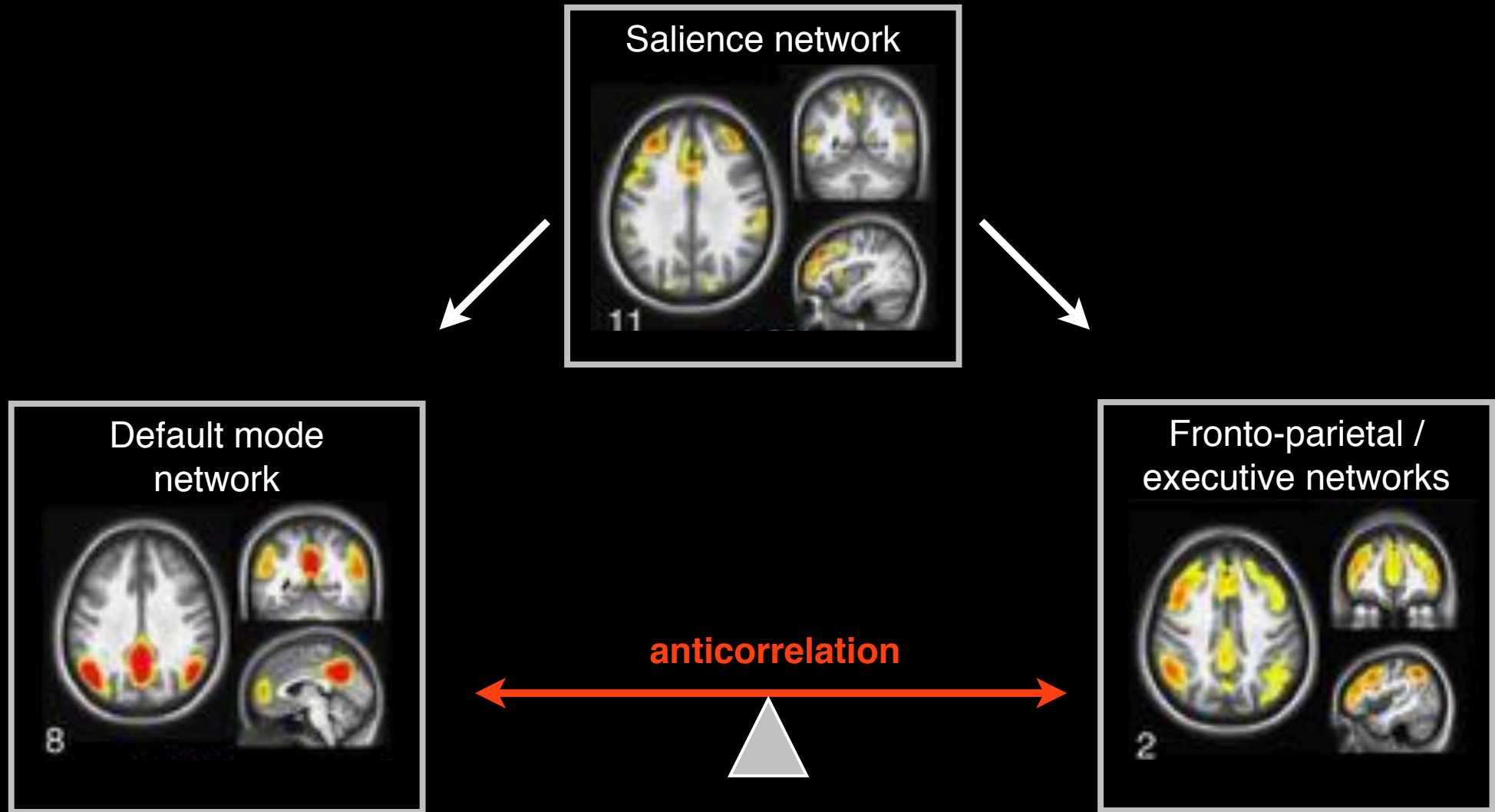
hierarchical clustering



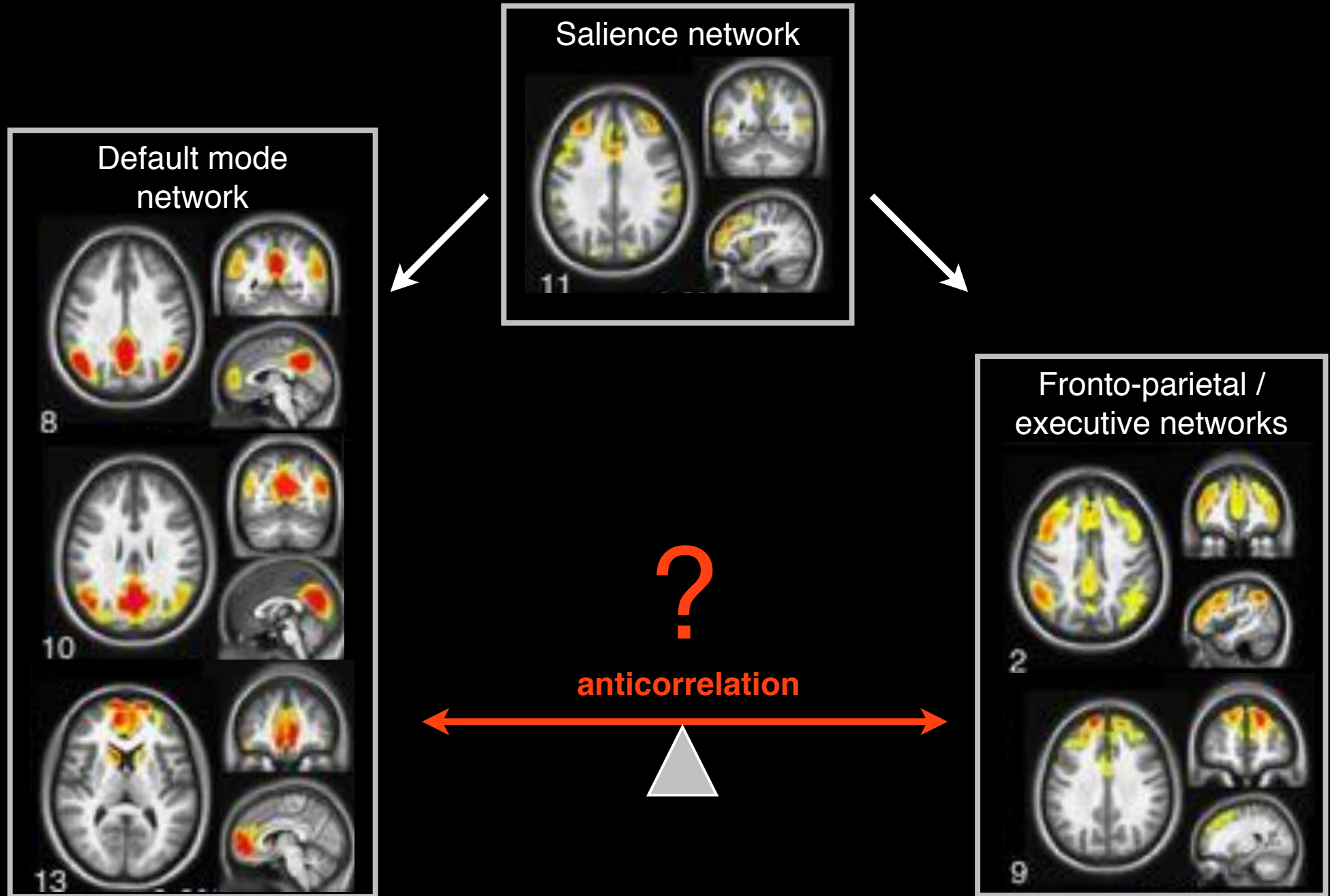
- Highest level of hierarchy: sensory / DMN / attention
- Behavioral profiles can be determined (BrainMap)
- ... and form consistent groupings as driven by iCAPs' combinations

BrainMap:
[Lancaster et al, Frontiers Neuroinformatics, 2012]

Time to rethink our models?

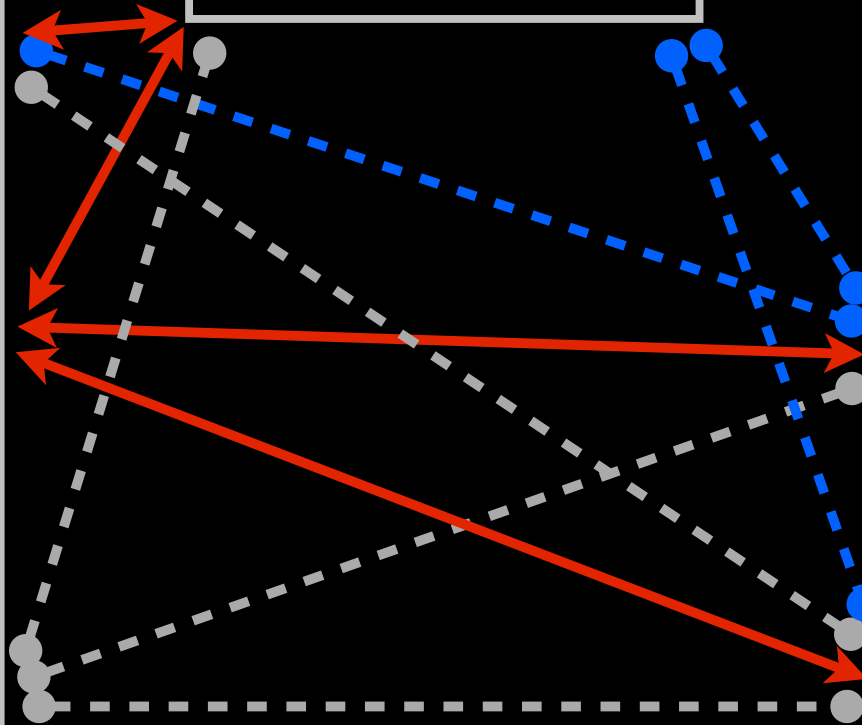
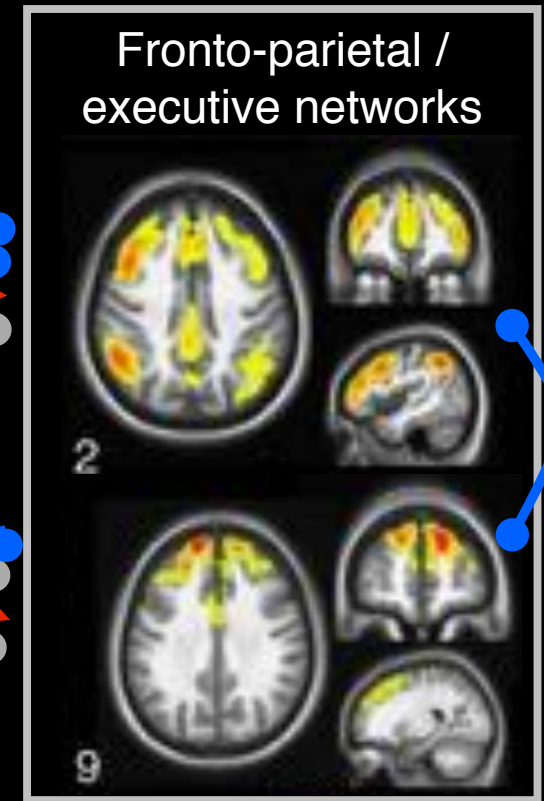
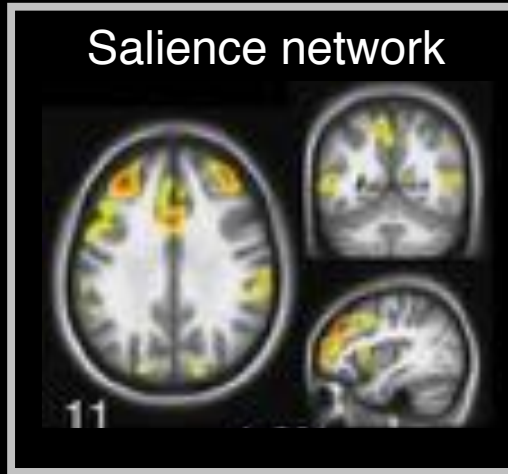
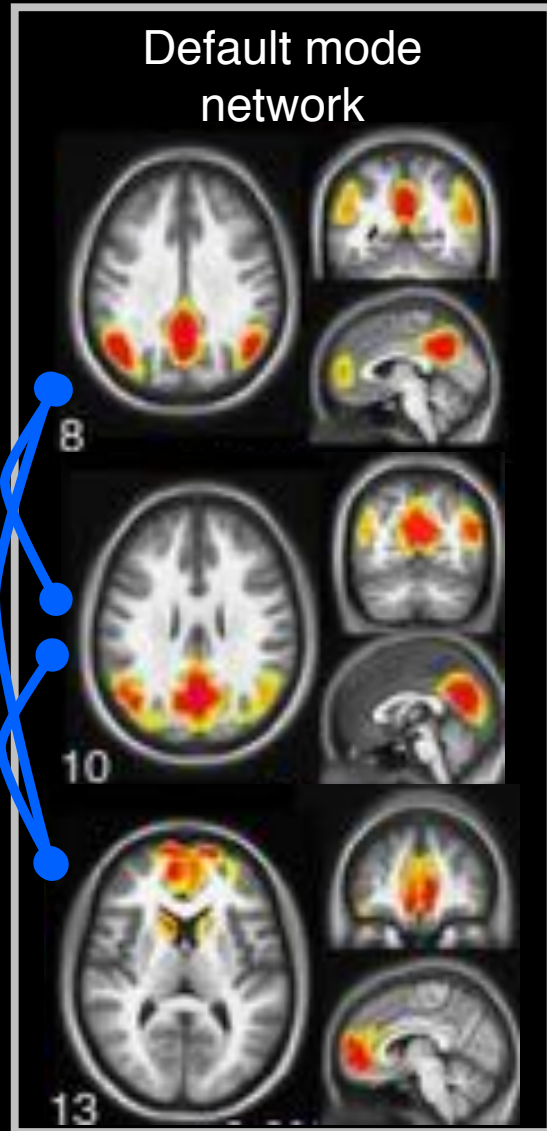


Time to rethink our models?



Time to rethink our models?

↔ opposite signs
● - - ● same signs
● - - ● mixed signs



Unraveling crosstalk of ongoing spontaneous activity

ICAPs

ATT

SVIS

ADMM

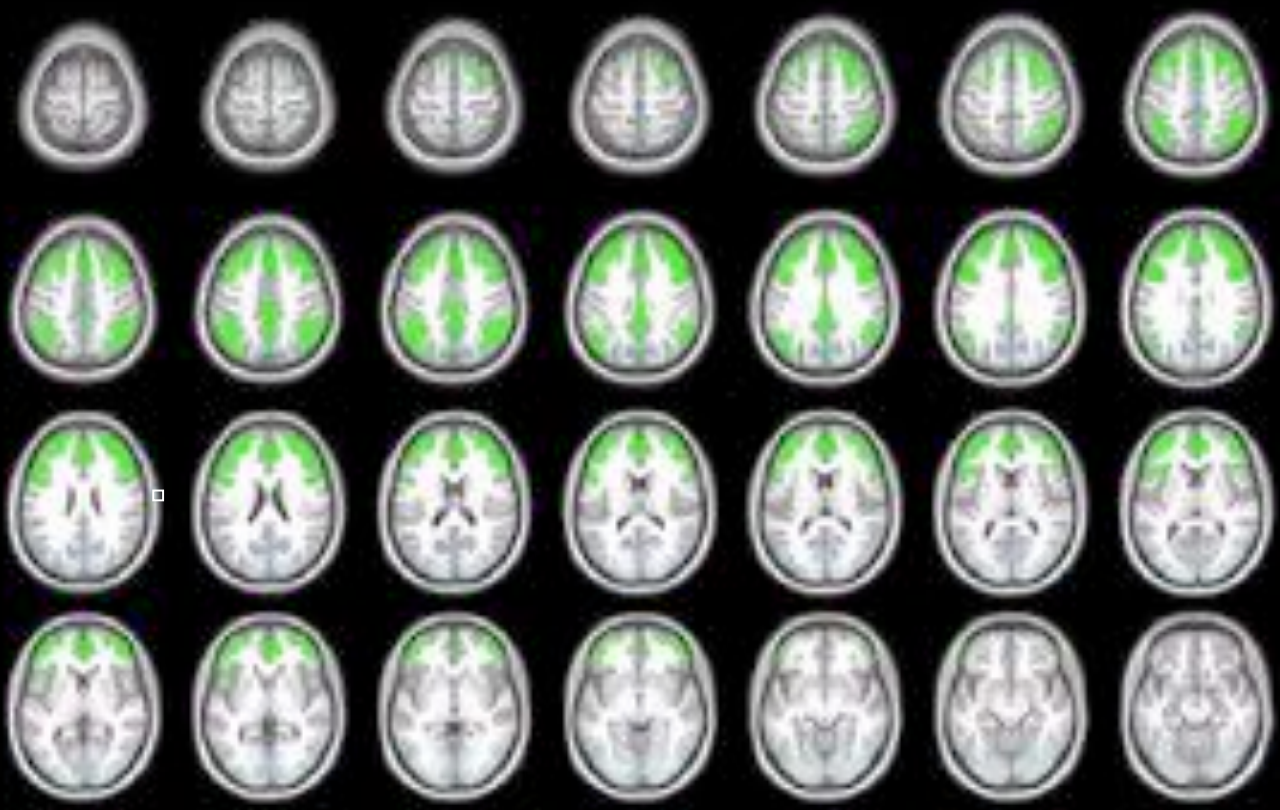
behavior

Action

Cognition

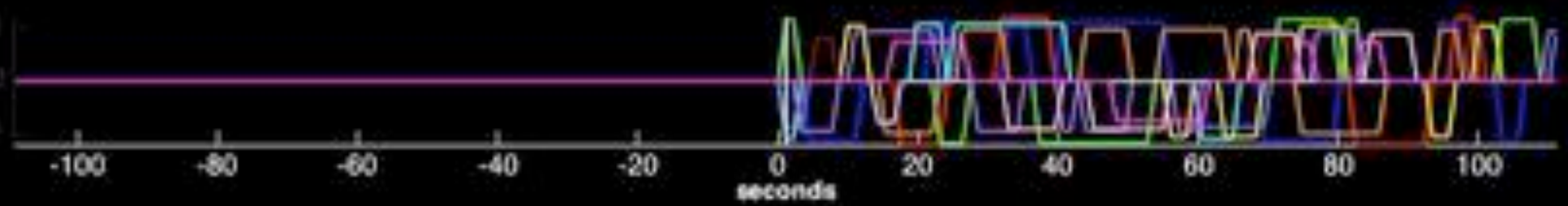
Emotion

Perception

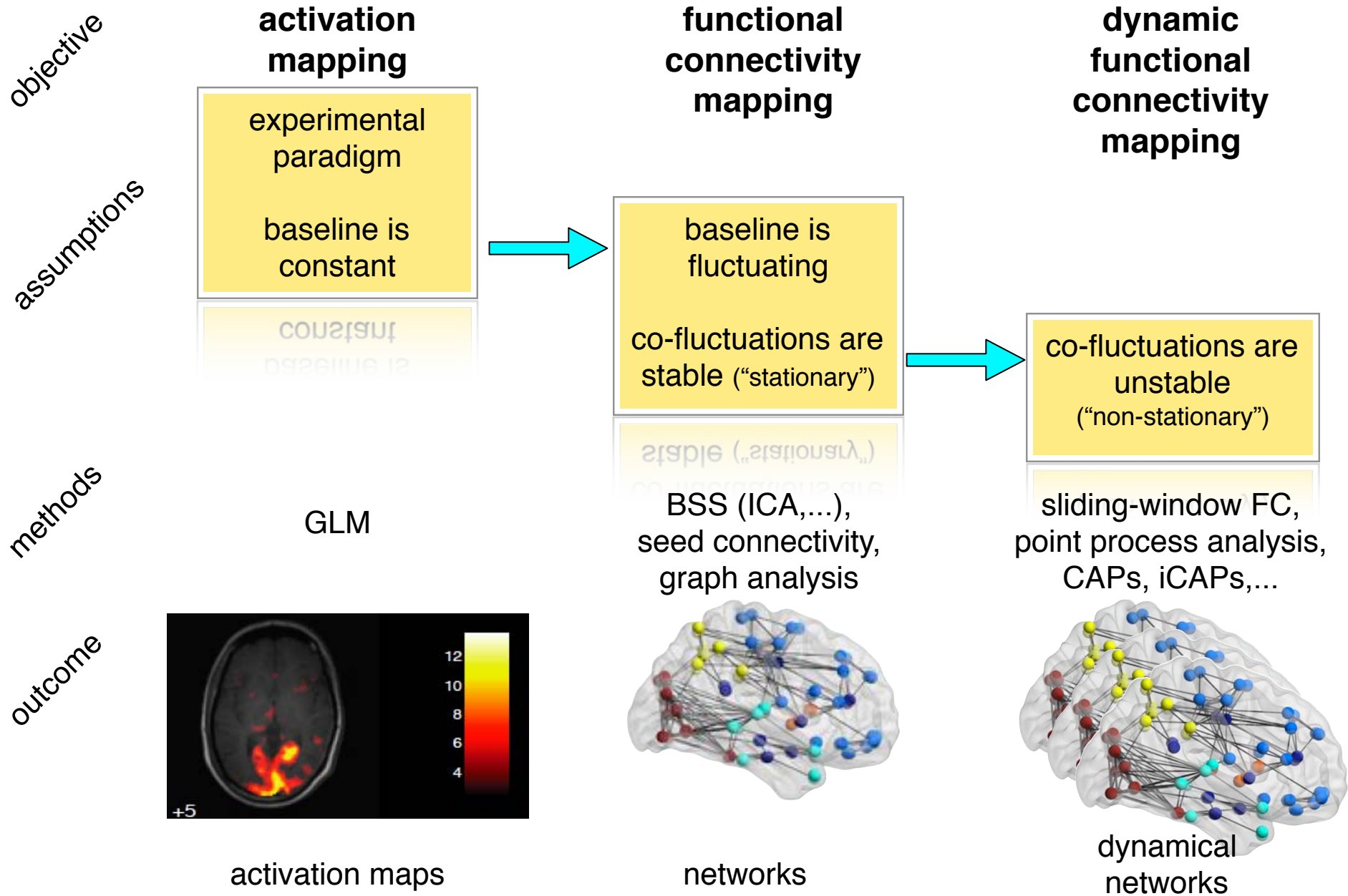


active networks

active behavior



Summary



Take home message



- Seeing the brain in action, at any *moment*, and *systems* level
 - *Multidisciplinary* endeavor where *computational* approaches are essential
 - Understand how it all fits together
- Perspectives
 - Tracking of brain states
 - Naturalistic stimuli and tasks
 - Neurofeedback
 - Graph signal processing: connect function (signal) with structure (graph)
 - Towards new models and markers of brain (dys)function
 - Benefit from “big MRI data” in health and disease

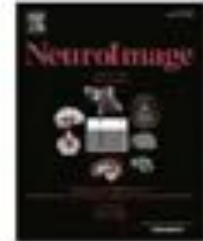
Two recent review papers



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The dynamic functional connectome: State-of-the-art and perspectives



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Current Opinion in
Biomedical Engineering

Dynamics of large-scale fMRI networks: Deconstruct brain activity to build better models of brain function

Fikret Işık Karahanoğlu^{a,b} and Dimitri Van De Ville^{c,d}

MIP:lab @ Campus Biotech

<http://miplab.epfl.ch>



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