

Network theory and brain disorders

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Disclosures

Employment

- **Paid Employment**
University of Cambridge (50% FTE)
GlaxoSmithKline (50% FTE)
- **Editorial Roles**
Biological Psychiatry – Deputy Editor
Network Neuroscience – Senior Editor
- **National Health Service (HCP) Role**
Hon Consultant Psychiatrist and Director of R&D,
Cambridgeshire & Peterborough NHS FT
- **Stock Equity (>\$10,000)**
GlaxoSmithKline
- **Speaker's Bureau**
None

Sources of Research Support

- Medical Research Council
- Wellcome Trust
- National Institute for Health Research
- National Institutes of Health,
Graduate Partnership Program

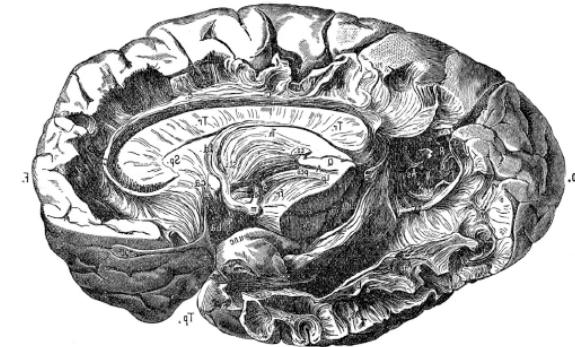
Brain networks and psychiatry go back a long way



Theodor Meynert

(1833-1892)

Anatomy of white matter



Carl Wernicke

(1848-1905)

Aphasia and psychosis as disorders of large scale brain connectivity

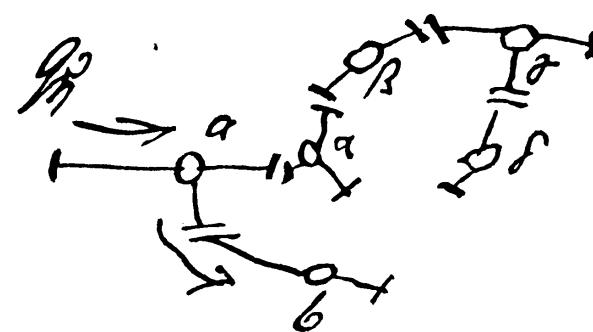


Sigmund Freud

(1856-1939)

Project for a Scientific Psychology

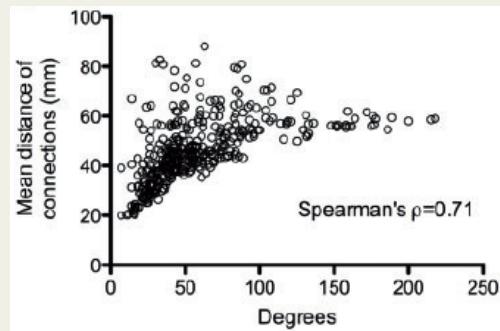
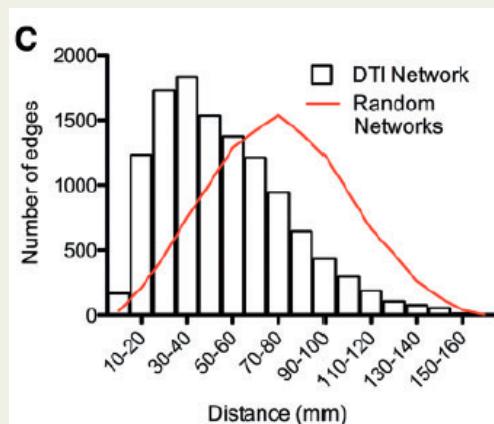
Mental states represented by flow of libido through cellular circuits



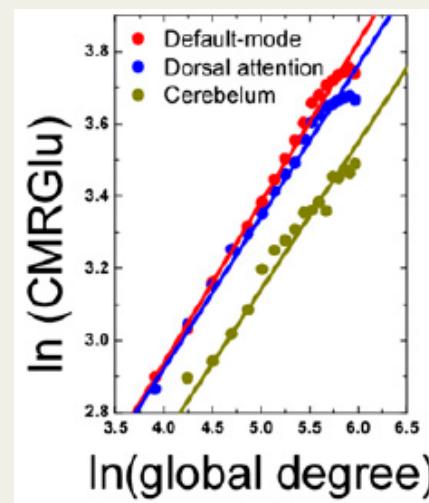
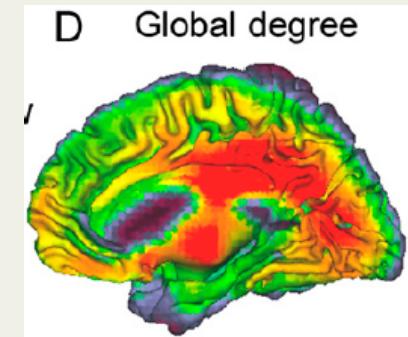
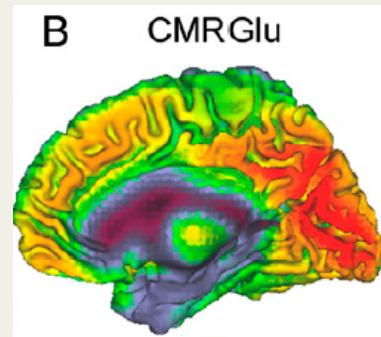
Economics of brain networks

A trade-off between “biological cost” and “topological value”

wiring cost of DTI networks



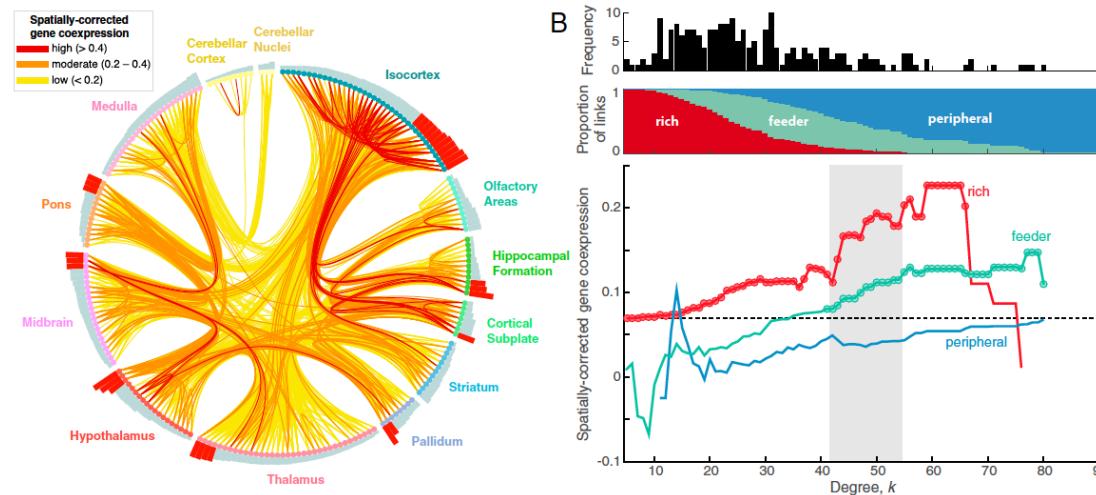
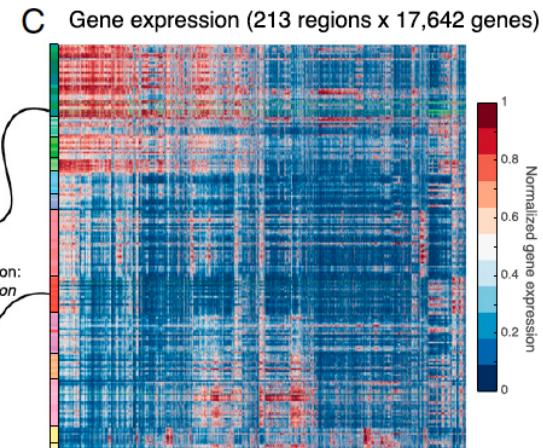
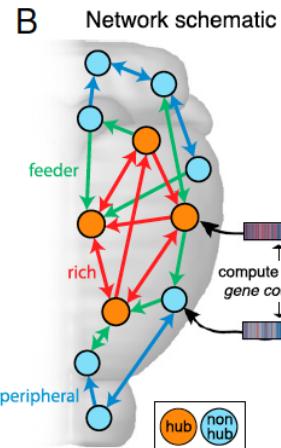
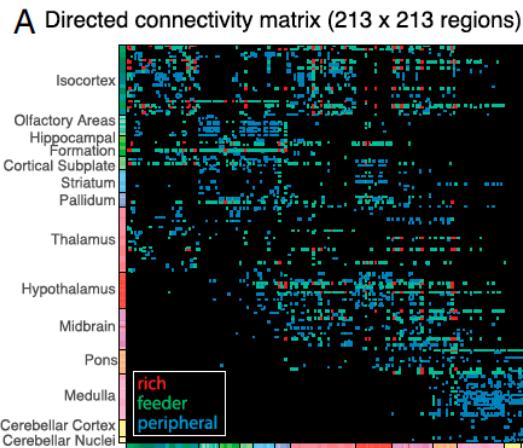
metabolic cost of fMRI networks



Crossley et al (2014) *Brain*

Tomasi et al (2013) *Proc Natl Acad Sci (USA)*

Mouse connectome topology and gene expression: connected hubs co-express genes for oxidative metabolism



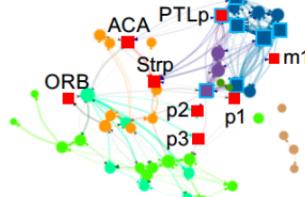
Anatomical connections between hub regions of the mouse tract-tracing connectome constituted a rich club.

Hub-hub connections linked brain regions that strongly co-expressed a set of genes significantly enriched for oxidative metabolism.

Genomics meets connectomics (in the mouse brain)

Reducing the dimensionality of a high dimensional problem

$O(10)$ nodal topology metrics



?

$O(10^4)$ mRNA measurements



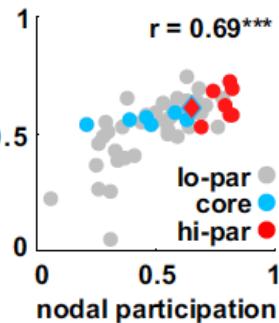
$O(10^2)$ regions

$O(10^2)$ regions

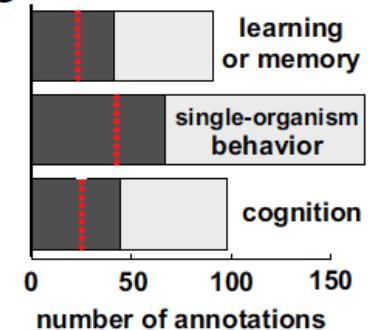
Partial Least Squares (PLS) is a family of multivariate methods for finding a few components that maximise the covariance between a set of response variables and a larger set of collinear predictor variables

Response variables = nodal topology (participation)
Predictor variables = nodal gene expression

B
gene-expression predictor



C

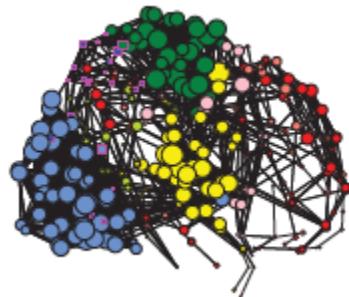


McIntosh & Bookstein (1996) *NeuroImage*

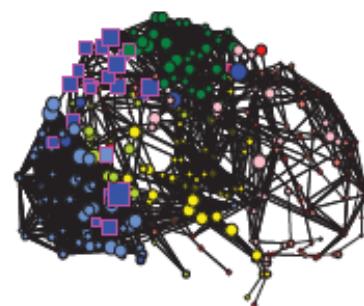
Rubinov, Ypma et al (2015) *Proc Natl Acad Sci USA*

Genomics meets connectomics (in the human brain)

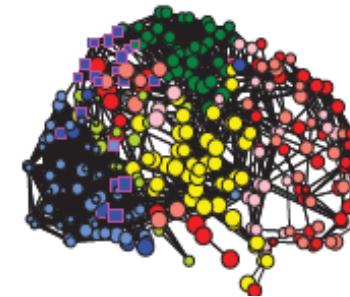
intra-modular degree



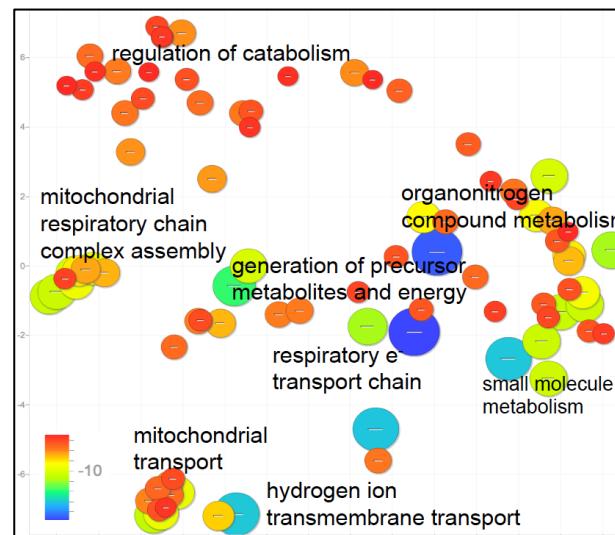
inter-modular degree



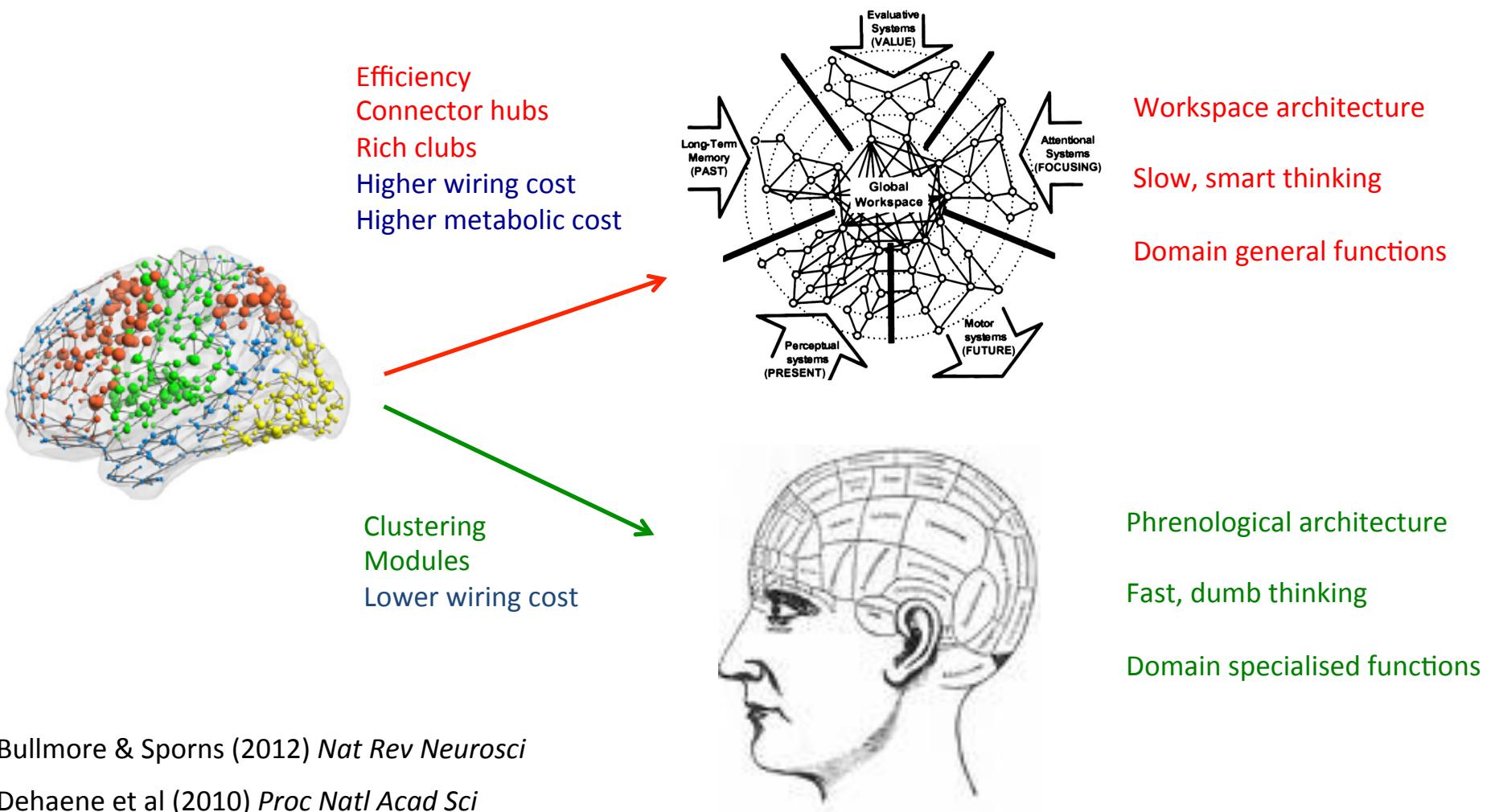
connection distance



	k	PC	k_{intra}	k_{inter}	d
PLS1	0.50	-0.26	0.59	0.07	-0.34
PLS2	0.26	0.45	-0.01	0.48	0.28



Cartoon of the economical connectome



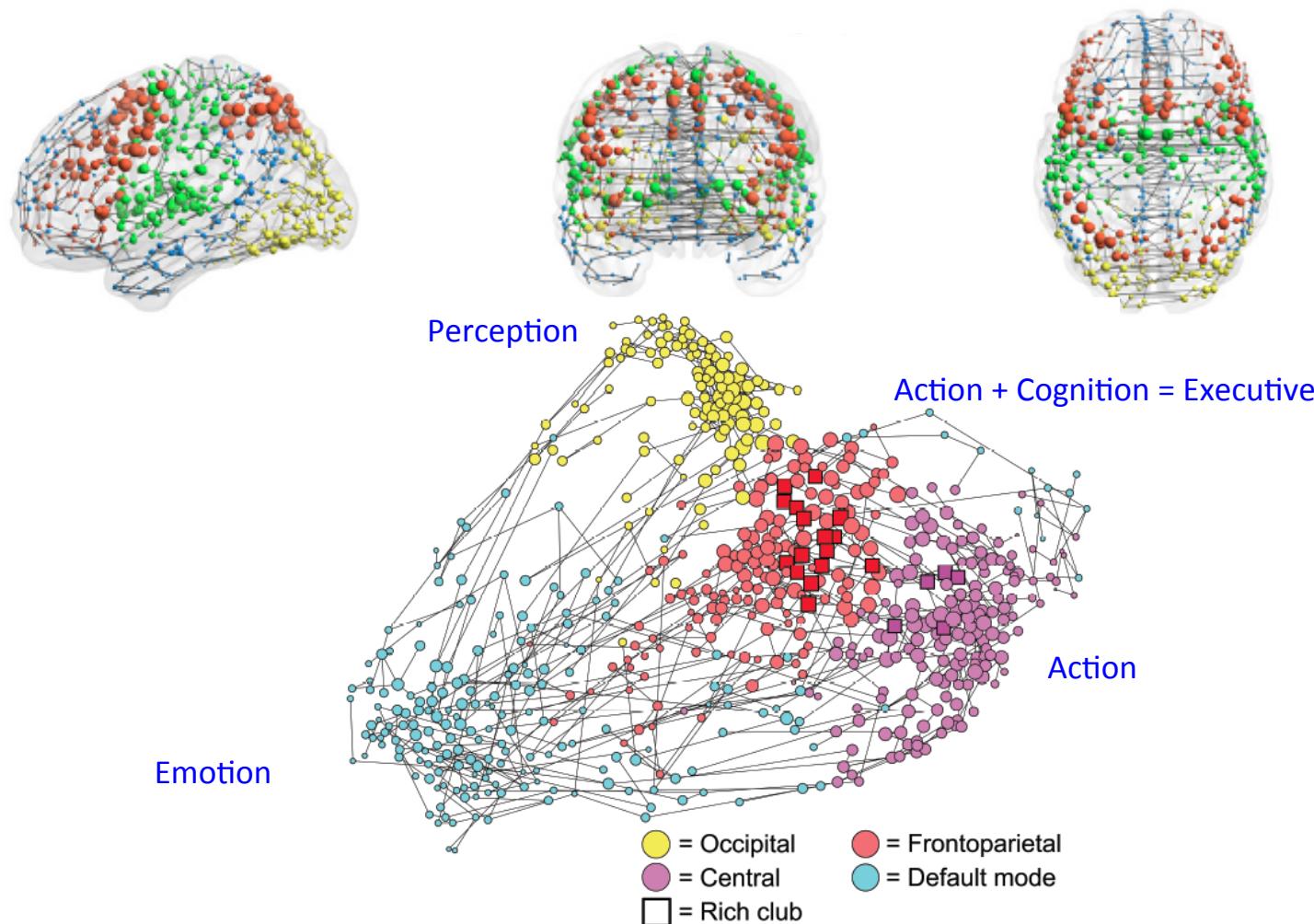
Bullmore & Sporns (2012) *Nat Rev Neurosci*

Dehaene et al (2010) *Proc Natl Acad Sci*

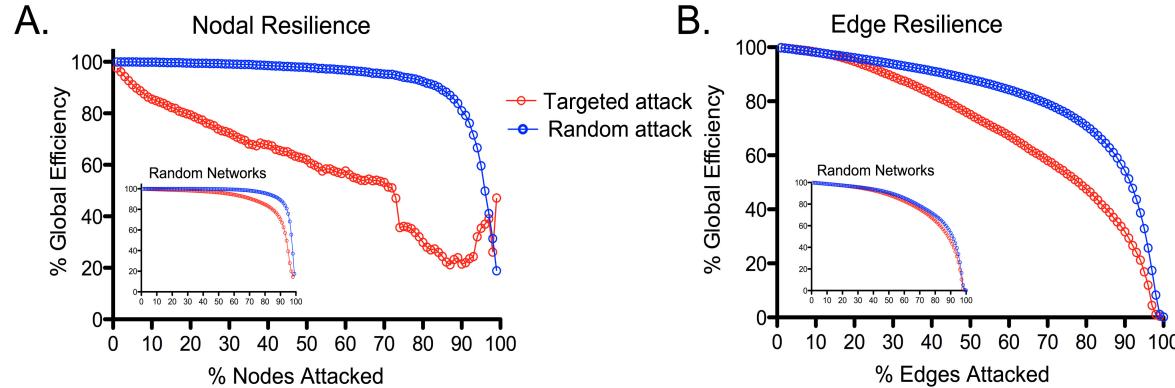
Fodor (1983) *Modularity of Mind*

Cognitive value of topological integration in human brain

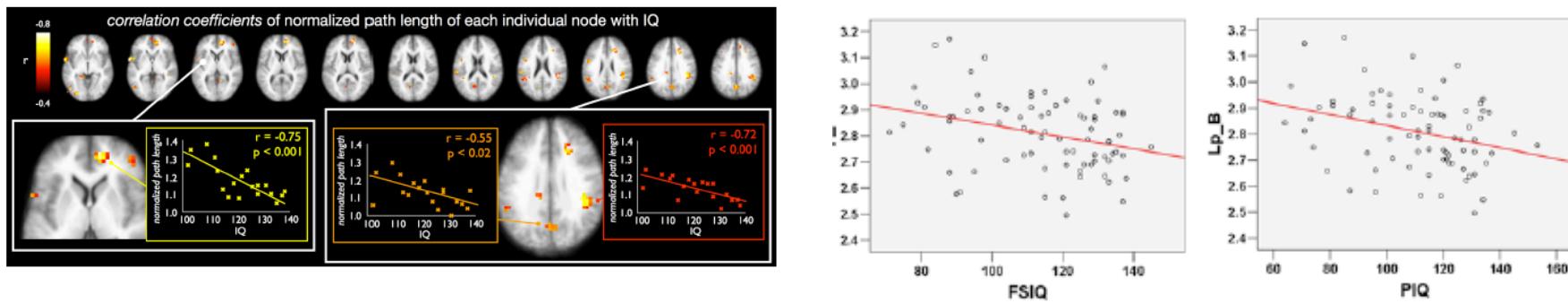
Meta-analysis > 1500 fMRI primary studies



Biologically expensive hubs may be “worth it” because they enable network integration, which is cognitively valuable



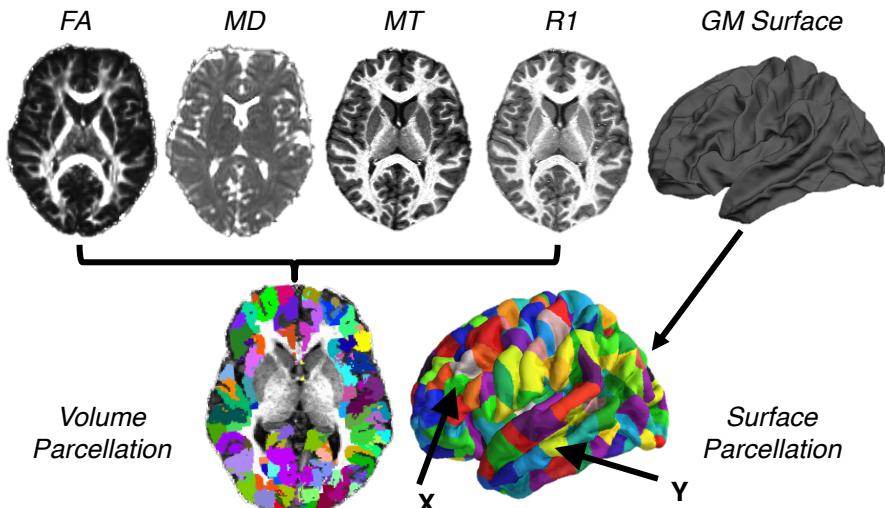
Hubs are important for integrative network configuration – targeted attack on hubs (or long distance edges) degrades global efficiency much faster than random attack



Integrative network configuration is important for adaptive behaviour and cognition – higher IQ goes with shorter network path length (greater global efficiency)

Morphometric similarity mapping

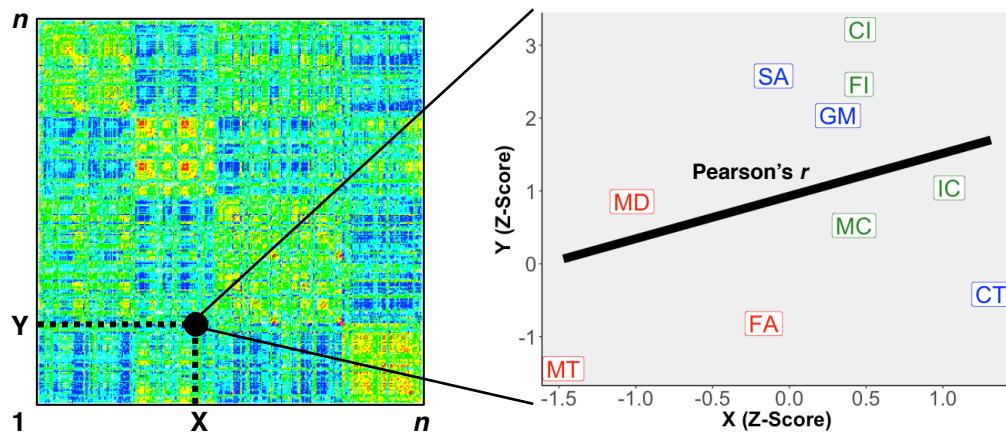
A new approach for single subject structural network analysis



Multiple morphometric features per regions

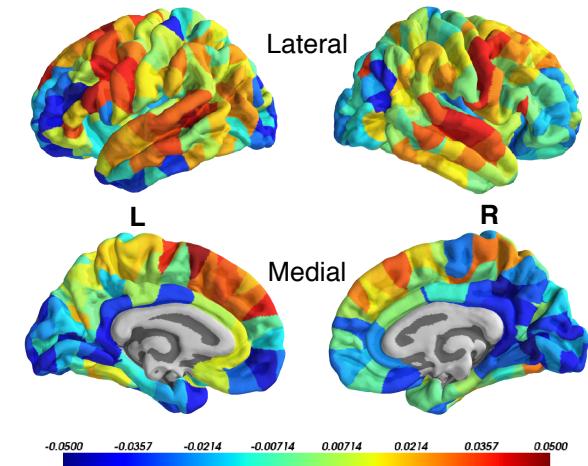
Myelination			Gray Matter			Curvature			
FA	MD	MT	GM	SA	CT	IC	MC	CI	FI

Morphometric Similarity Matrix (MSM, N=1)



Morphometric Similarity Network (MSN, N=1)

Weighted degree



MSN Modules and Rich Club (10% Connection Density)

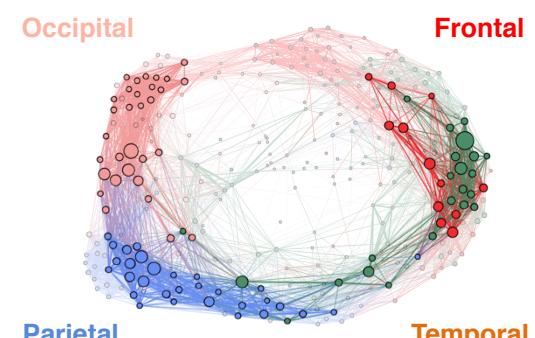


Occipital

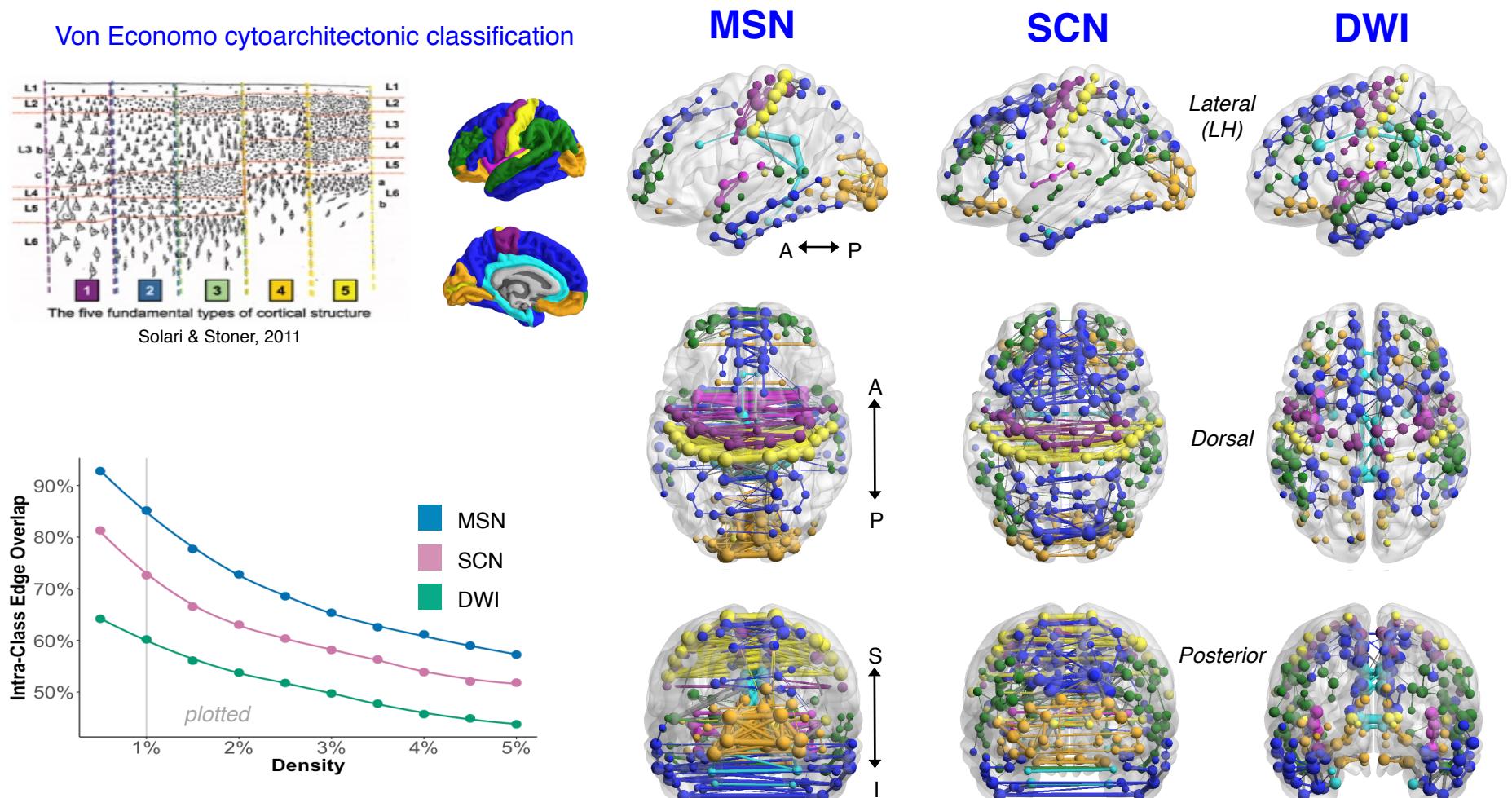
Frontal

Parietal

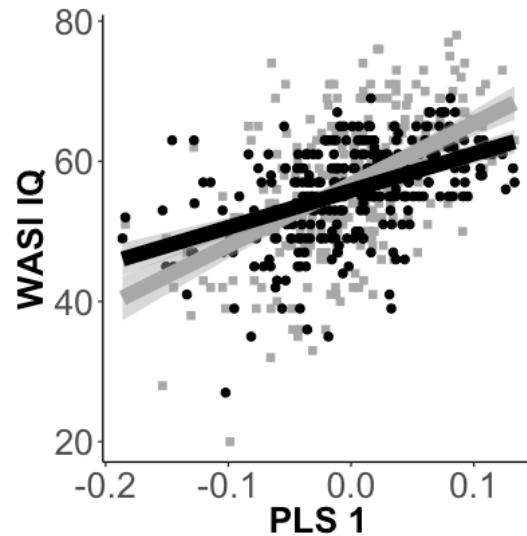
Temporal



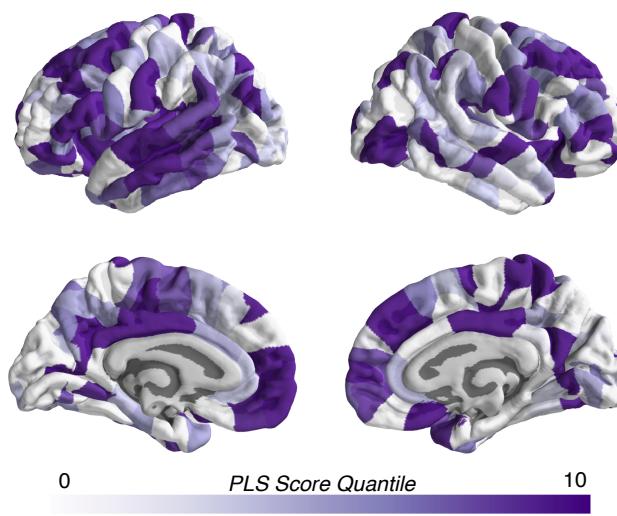
Morphometric similarity networks are more consistent with cytoarchitectonics than structural covariance networks (SCN) or diffusion weighted imaging (DWI)



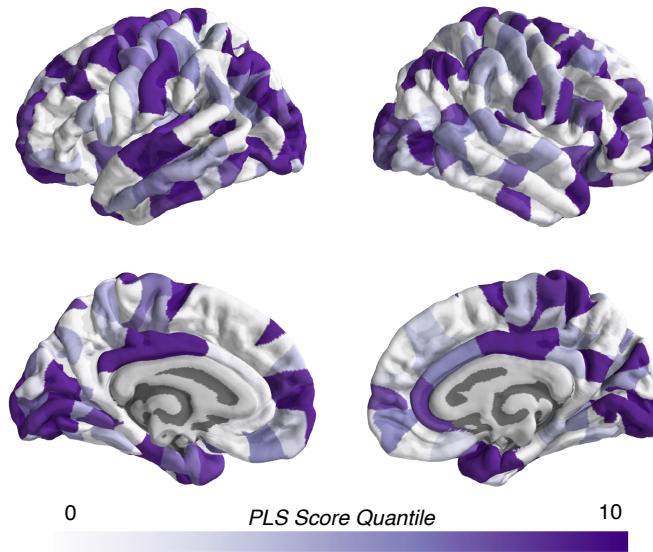
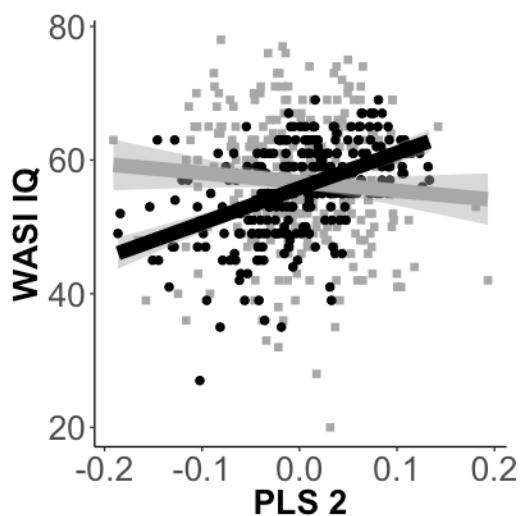
Hubs of morphometric similarity networks predict about 40% of between-subject variance in IQ



- Matrix Reasoning
- Vocabulary

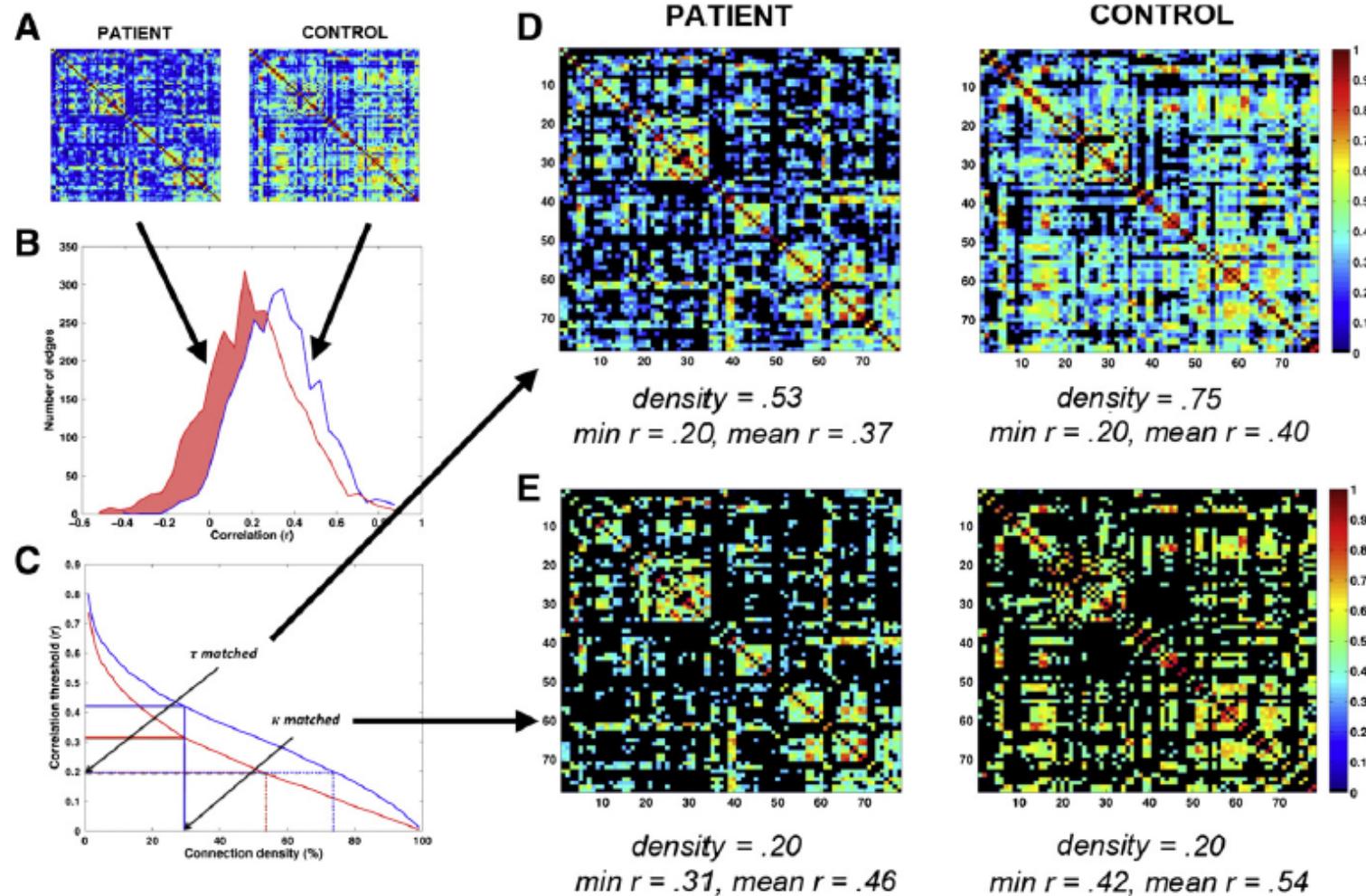


spectral acoustic comprehension language network phonological articulatory musical english production heard syntactic sentence semantically spoken broca self speech sounds music speech production noise referential default listened lexical pitch personal semantic information



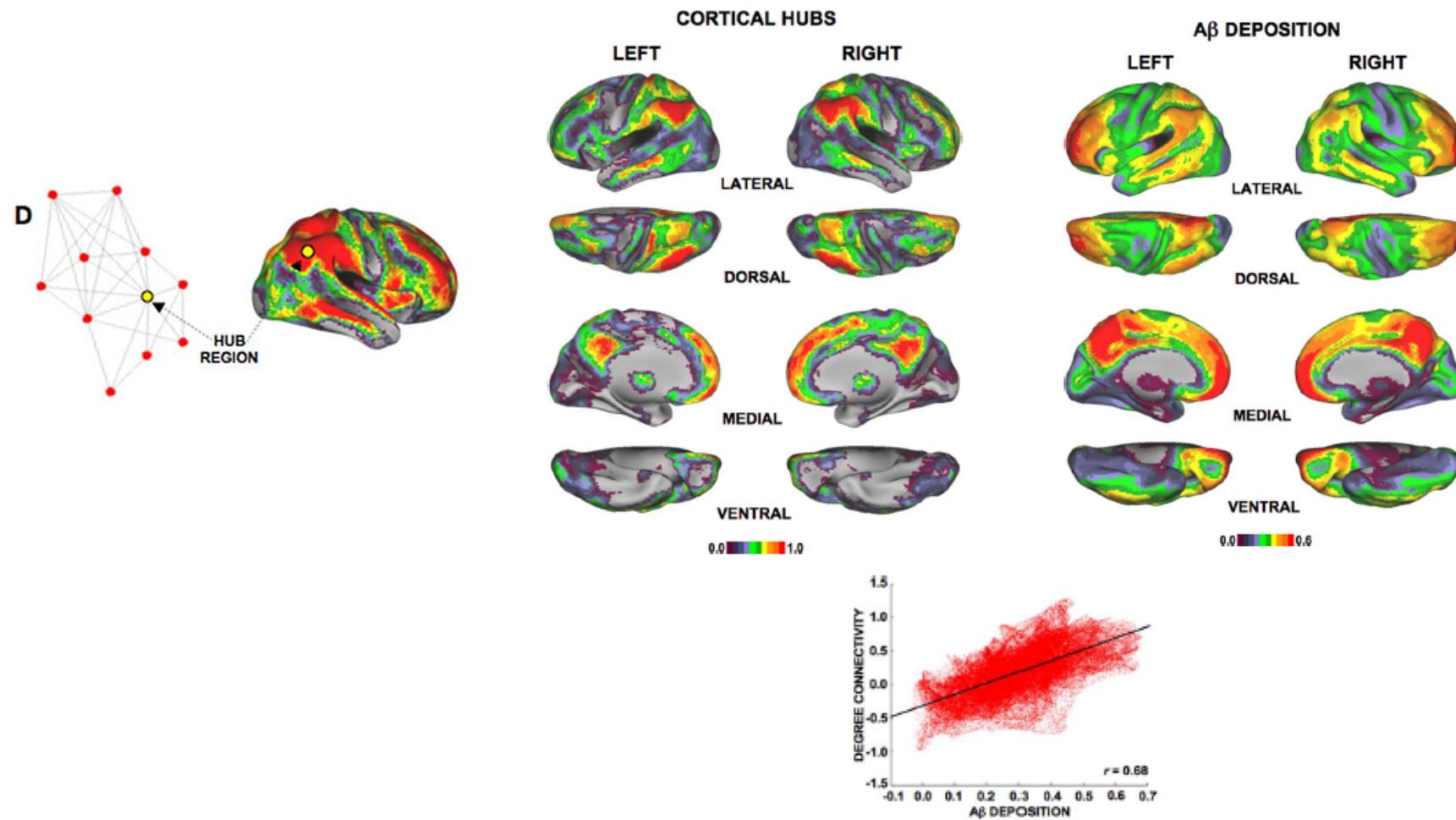
alzheimer disease repetition suppression faces written adaptation formation categories repetition face word recognition word form selectivity imagery semantic memory vision object reach sighted color visual word consolidation recognition identity objects navigation encoding concepts congruent eye movements letters form recall episodic memory category

Case-control comparisons can be tricky in fMRI network studies of clinical disorders



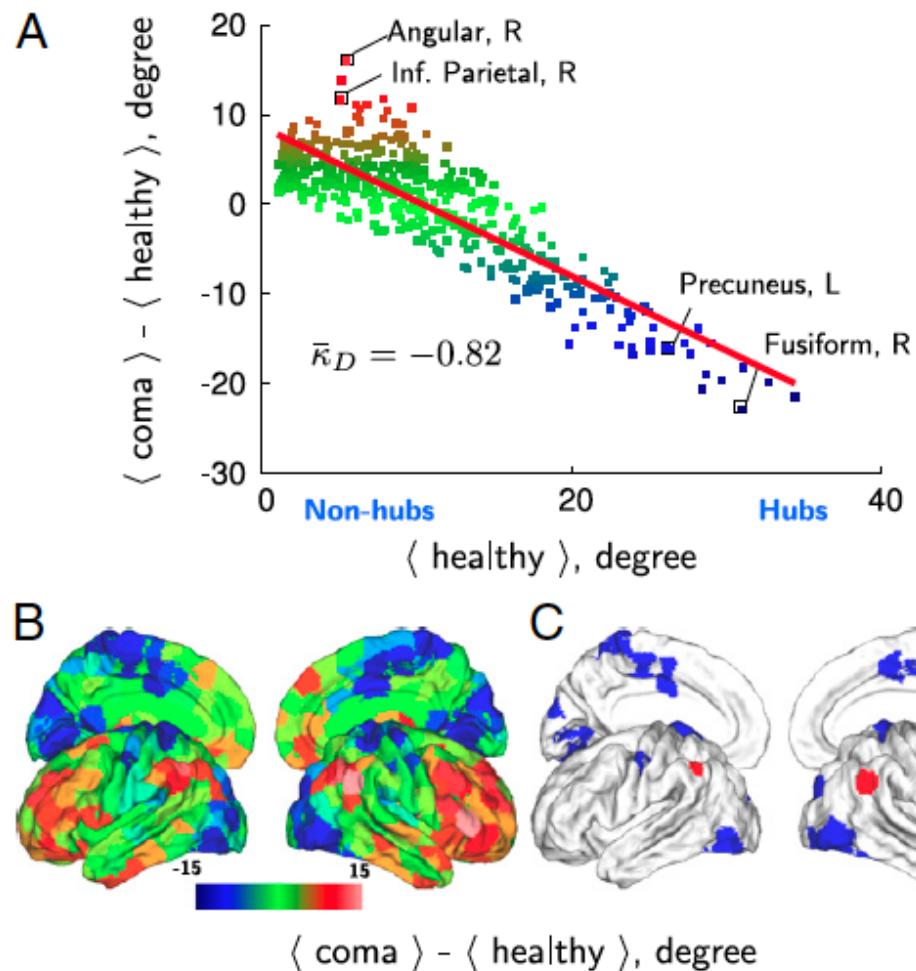
Fornito et al (2012) *NeuroImage*

The hubs of brain networks are most vulnerable to amyloid deposition in Alzheimer's disease



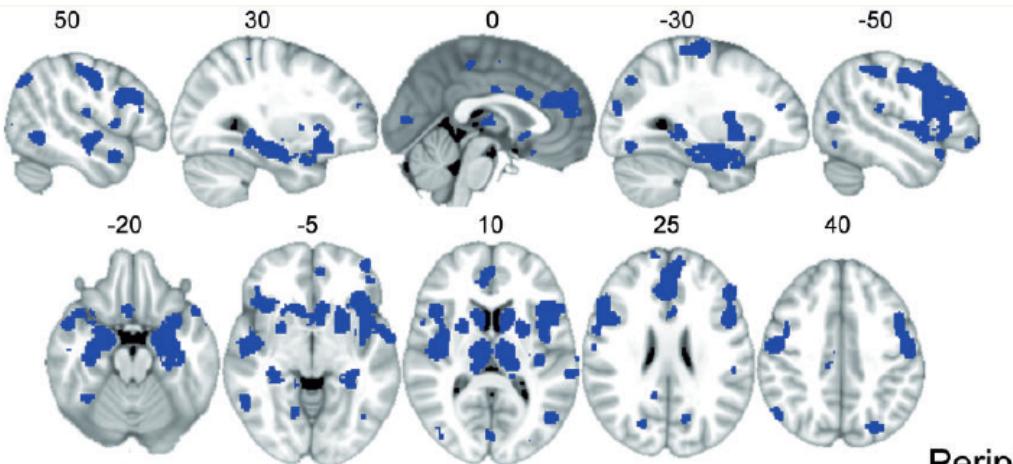
Buckner et al (2009) *J Neurosci*; Buckner et al (2008) *Ann NY Acad Sci* [review]

Coma due to acute brain injury is associated with radical disruption of hubs

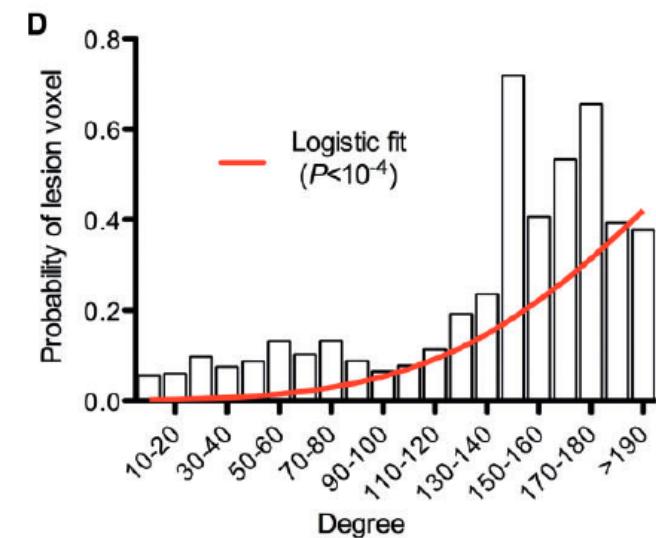
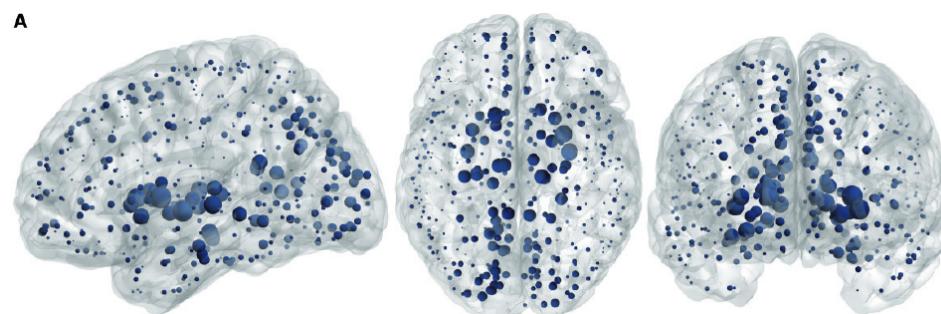


High cost / highly central hubs are brain disease “blackspots”

Meta-analysis of MRI data on 20,000 patients, 26 disorders



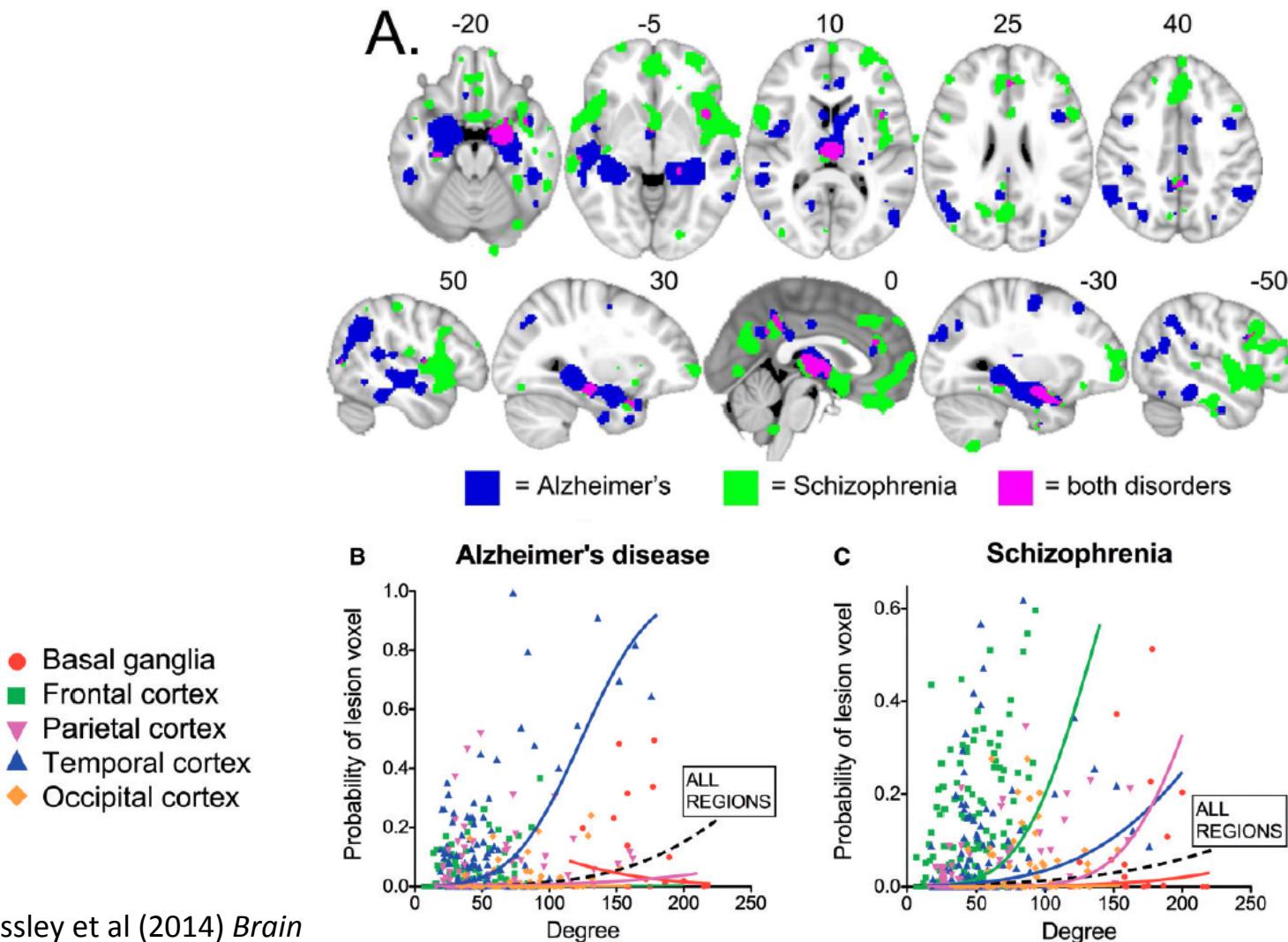
A



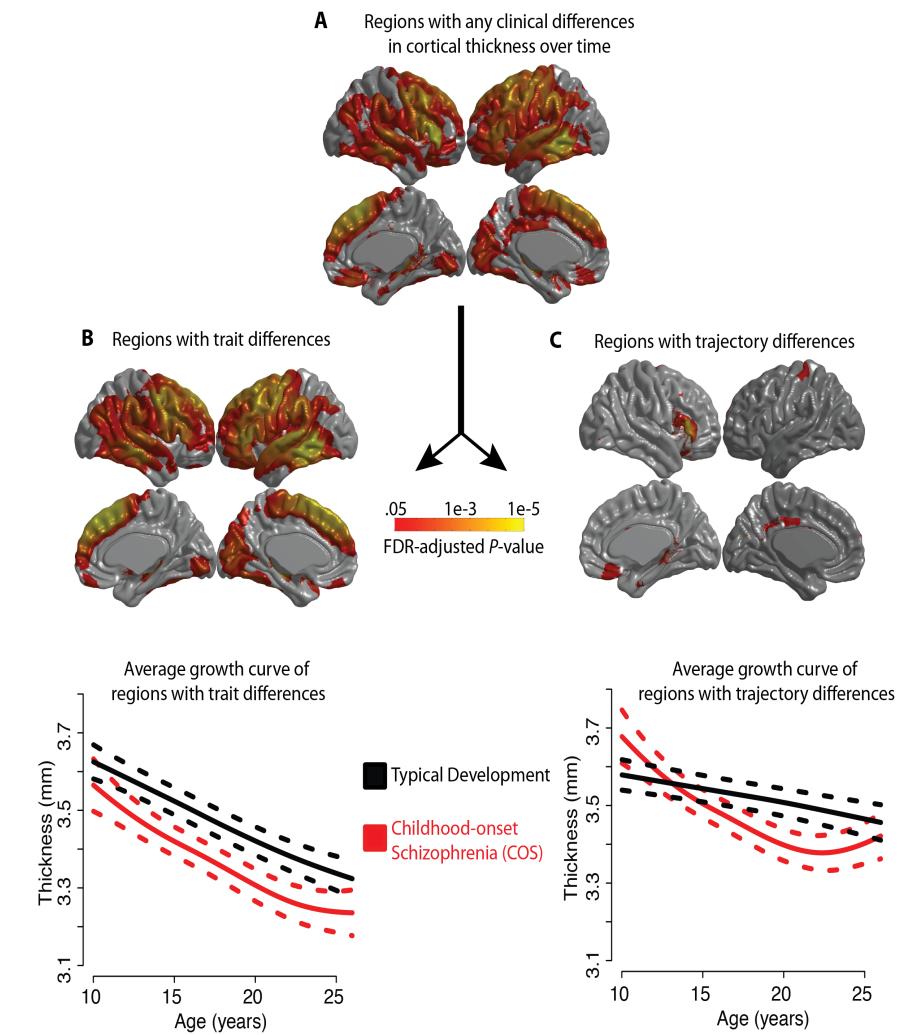
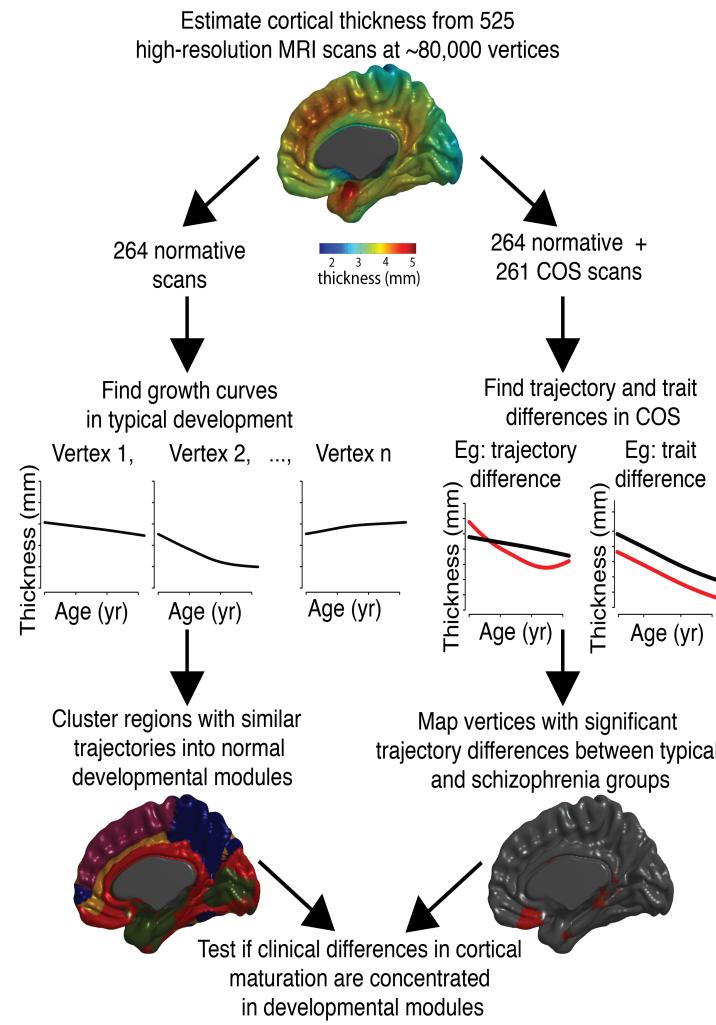
High cost could make hubs more vulnerable to disease processes

High centrality could make lesioned hubs more likely to be symptomatic

Alzheimer's disease and schizophrenia both preferentially impact hubs, but not the same hubs



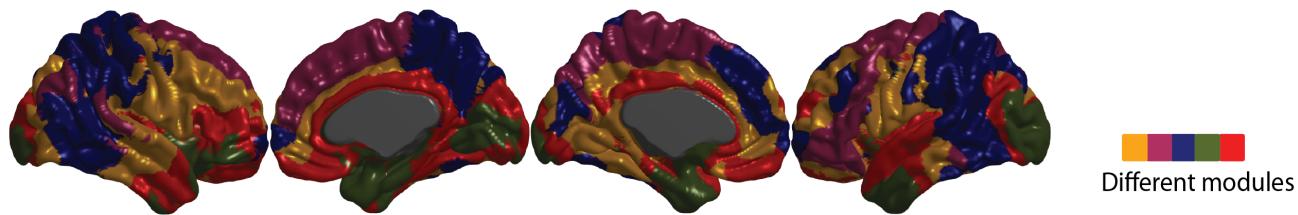
Linking abnormal cortical maturation in schizophrenia to the modular community structure of the anatomical connectome



Abnormal trajectories of cortical shrinkage in adolescent schizophrenia are concentrated within a single module of the normal connectome

A

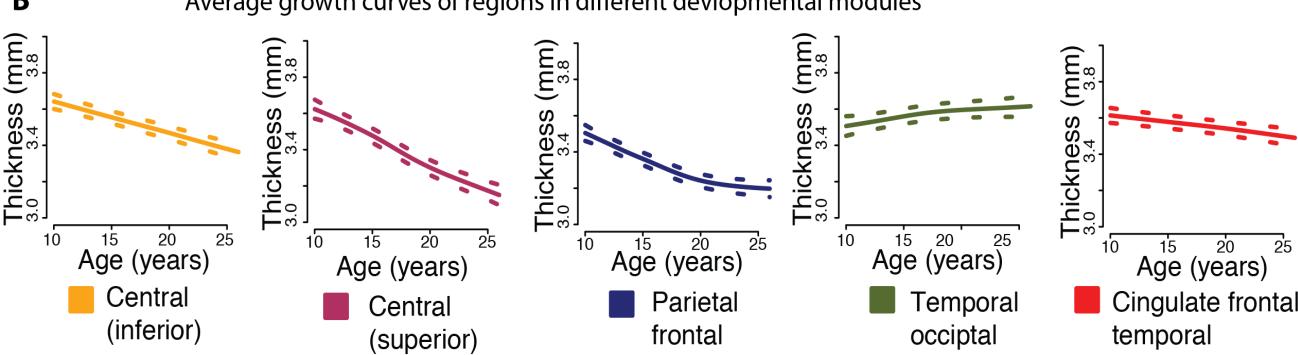
Developmental modules in typical development



Different modules

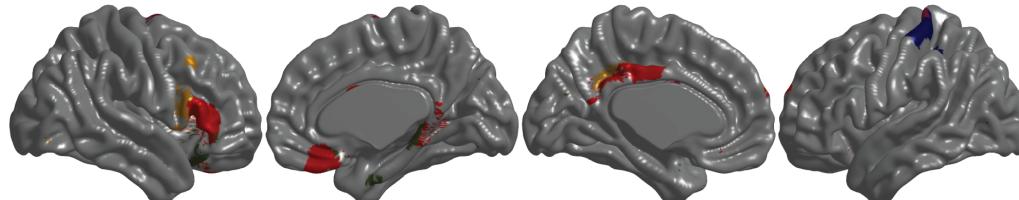
B

Average growth curves of regions in different developmental modules



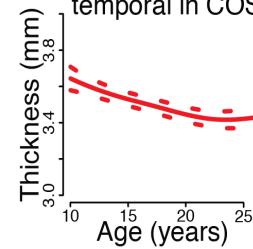
C

Regions with trajectory abnormalities in COS, color-coded by developmental module



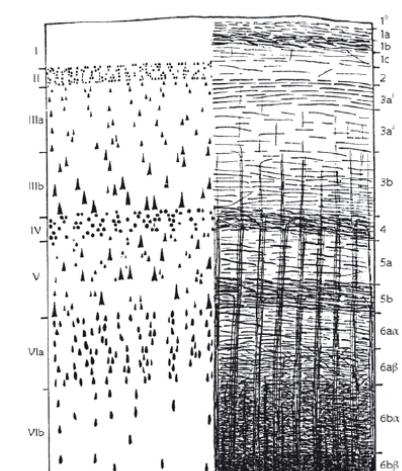
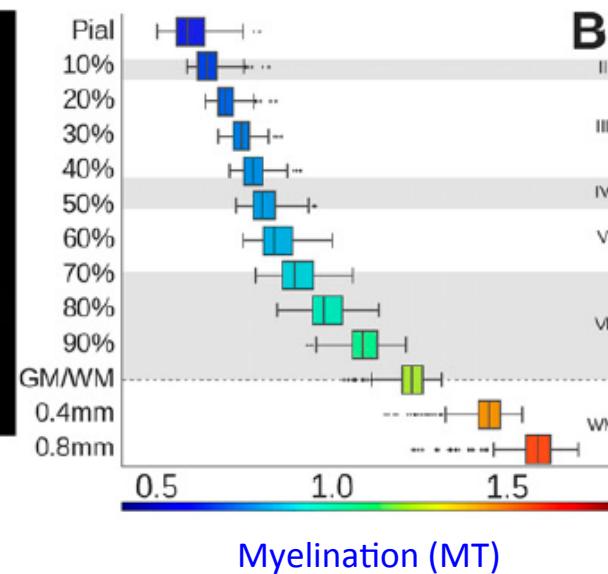
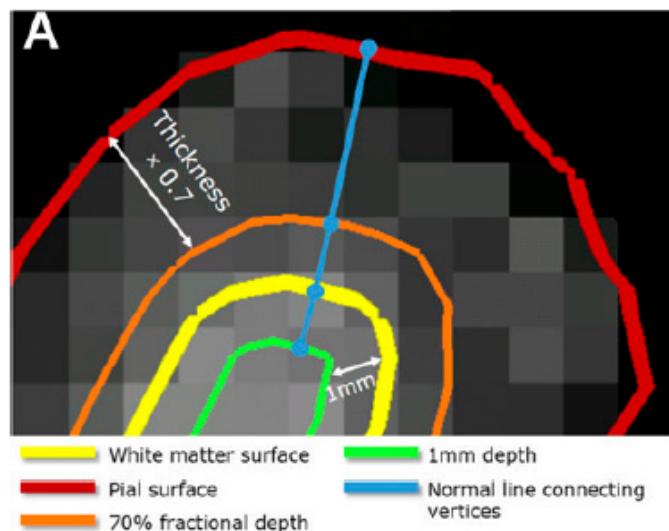
D

Cingulate frontal temporal in COS



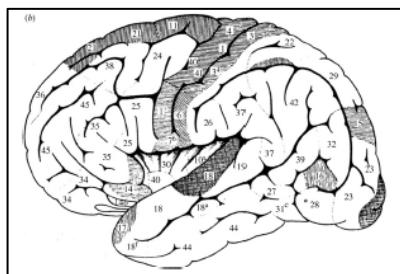
MRI is great for measuring the difference between fat and water

(in the brain that's the difference between myelin and everything else)

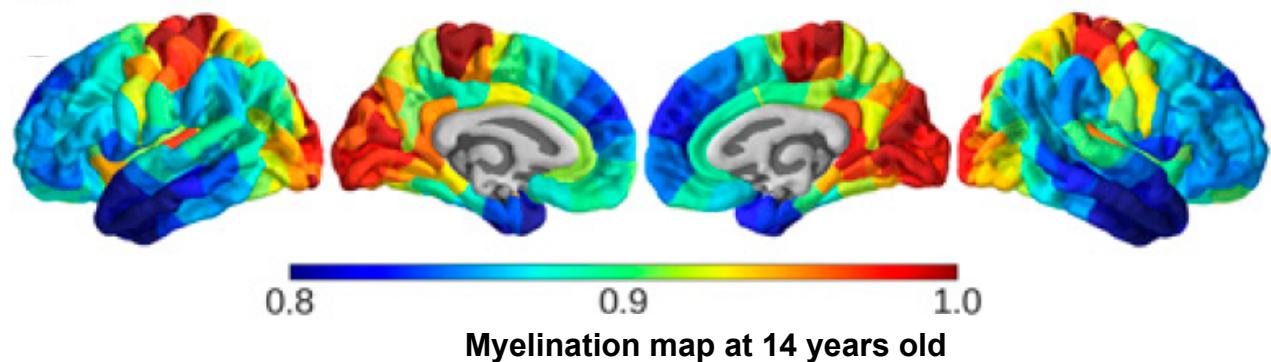


Myelination (Vogt,1903)

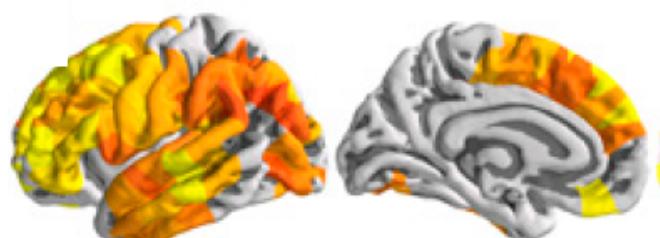
Adolescent shrinkage of association cortex is highly correlated with intracortical myelination



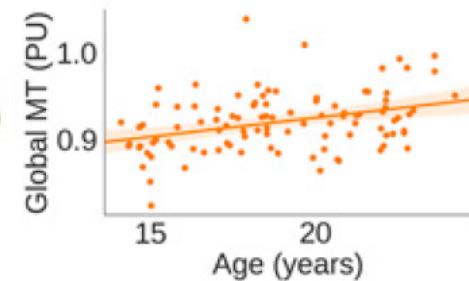
Paul Flechsig (1847-1929)



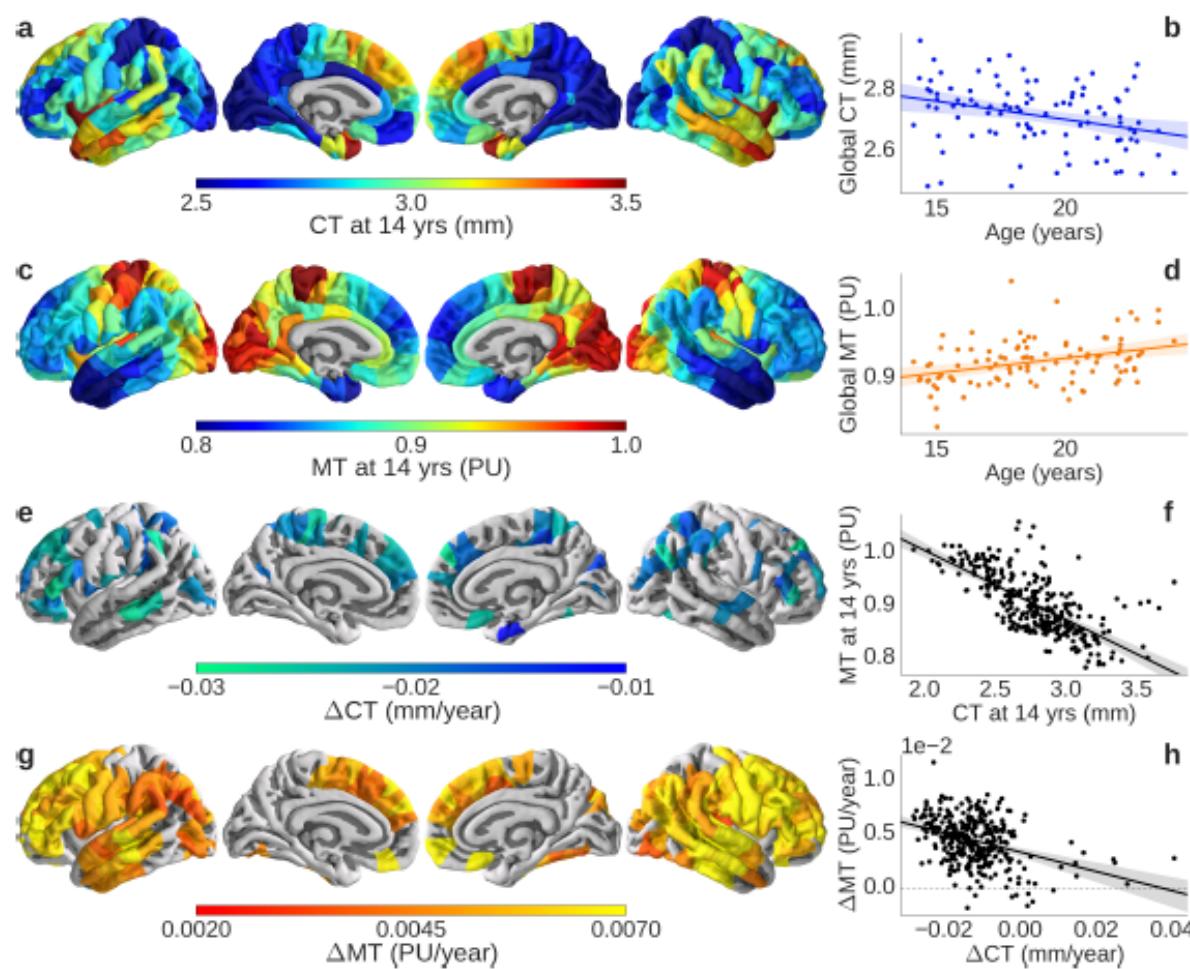
Myelination map at 14 years old



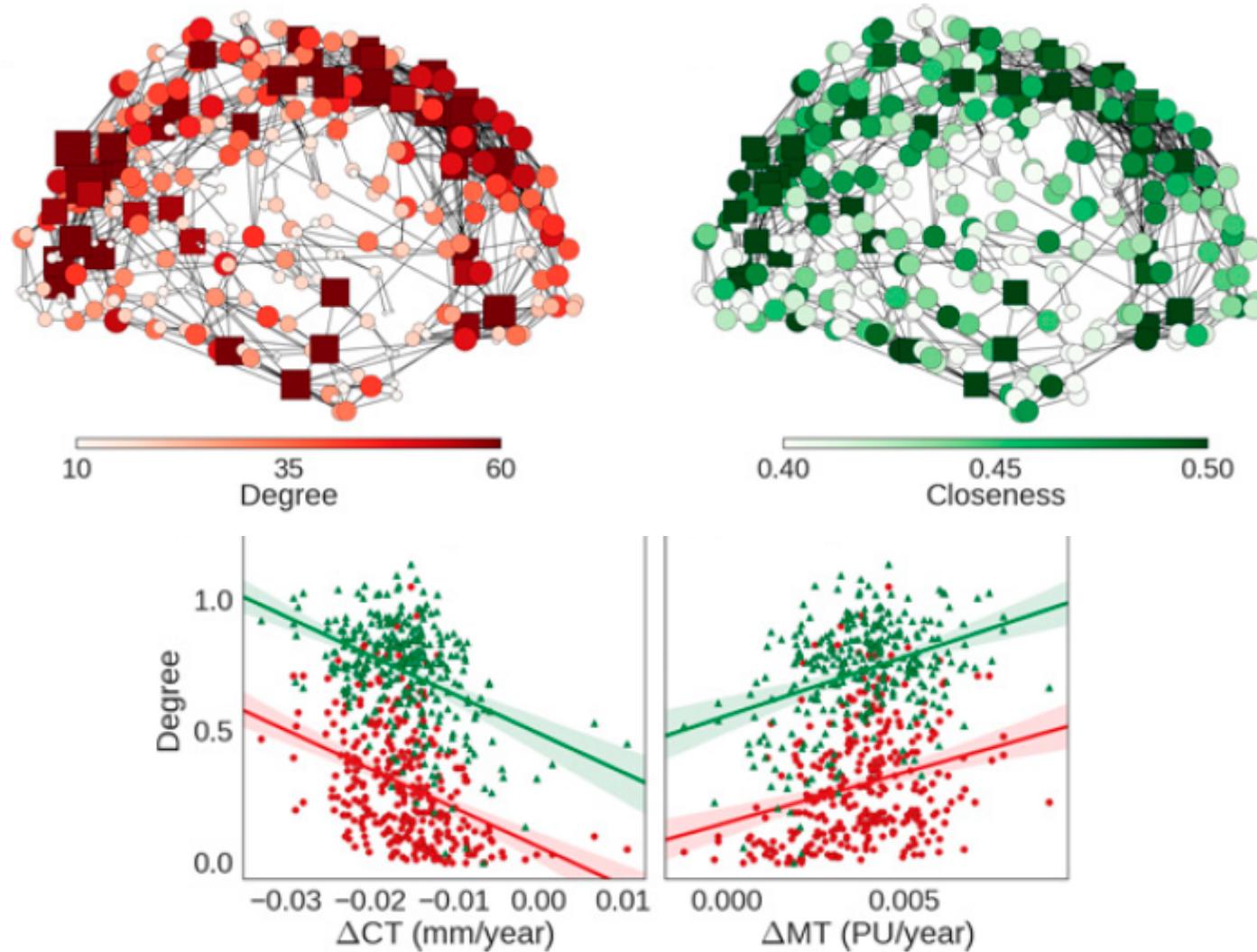
Map of increasing myelination
during adolescence, 14-25 years



Adolescent shrinkage of association cortex is highly correlated with intracortical myelination

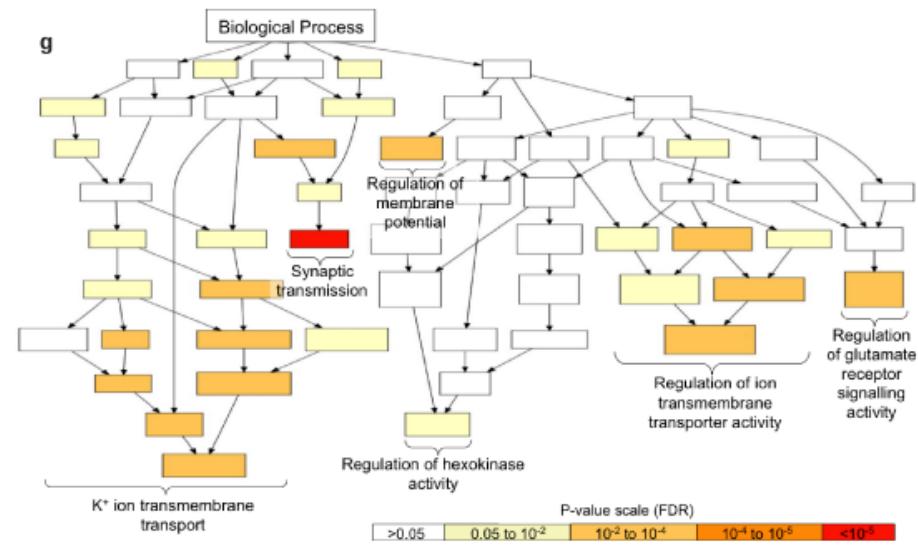
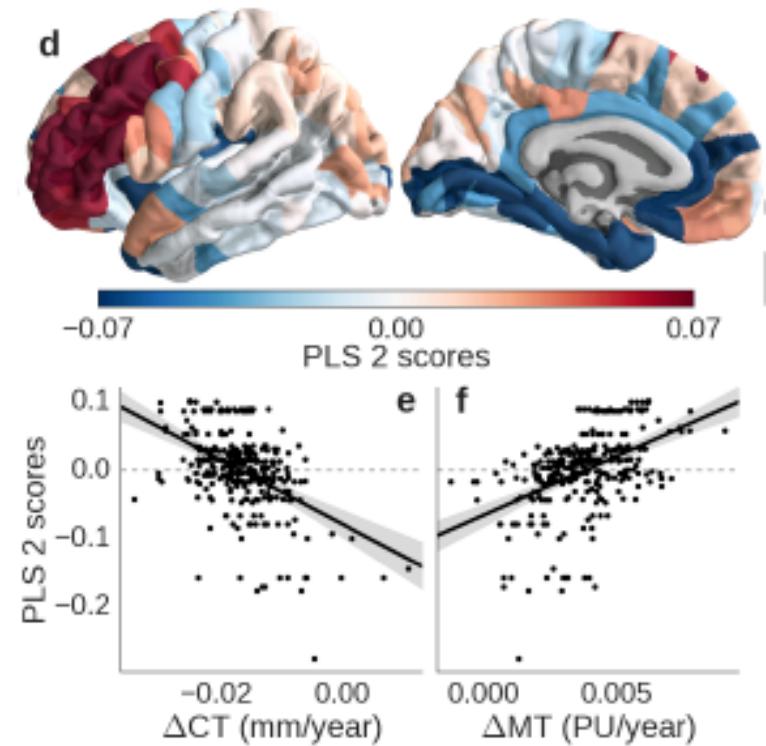


Connectome hubs have fastest rates of cortical myelination (and shrinkage) during adolescence

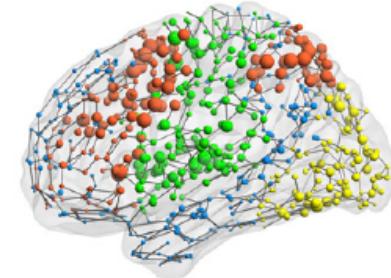


Whitaker, Vértes et al (2016) *Proc Natl Acad Sci USA*

Adolescent myelination of connectome hubs is associated with expression of risk genes for schizophrenia



Review



- Psychiatry wanted to understand the connectome long before it was possible
 - Our history is rooted in ideas of brain network disorganization and development
- Connectomes represent a trade-off between biological cost and topological integration
 - Hubs and clubs are biologically expensive but “worth it” for cognitive value added
- High cost / high value network hubs are blackspots for brain disorders
 - Brain network hubs are likely more vulnerable to disease and more symptomatic if lesioned
- Schizophrenia is a disorder of adolescent consolidation of connectome hubs
 - Linking connectomics to genetics can lead to a more mechanistic understanding of abnormal brain network development in mental health disorders

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