








The brain at 'rest': investigating spontaneous activity in BOLD fMRI using Independent Component Analysis

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The 'resting' condition

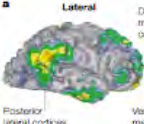
"The fact that the body is lying down is no reason for supposing that the mind is at peace. Rest is far from restful"

 Seneca the younger, ~65A.D.

- *Poorly characterised experimental condition:*
"a behavioural condition characterised by quiet repose usually with eyes closed but occasionally, in the experimental setting, with eyes open with or without visual fixation."

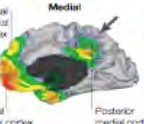
 Raichle, M, Trends Cogn Sci 2010

'Default Mode' of Brain Activity



Lateral

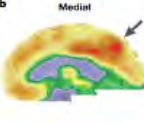
Posterior lateral cortex



Dorsal medial cortex




Ventral medial cortex

Posterior medial cortex



Medial

- Meta-analysis of 9 functional PET studies (N=132): consistent decreases during performance of attention demanding cognitive tasks
- Glucose metabolism in N=22 subjects under rest measured using FDG-PET

 Shulman et al., J Cog Neurosci 1997
  Gusnard et al., NRN 2001
  Raichle et al., PNAS 2001

Brain connectivity

Functional Connectivity

–temporal correlations between
spatially remote
neurophysiological events



Resting-State Networks

–Multiple spatial patterns of
correlated temporal dynamics,
resembling activation map



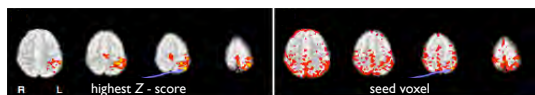
RSN analysis

Need to estimate whole-brain temporal
correlations...

Two popular approaches for RSN analysis:

- Seed-based correlation (Biswal, Raichle/Fox)
- ICA-based analysis (Kiviniemi, Beckmann, Calhoun)

Seed-based analysis



Activation maps from a
finger tapping experiment


Correlation maps from a
resting state experiment

- Biswal (1995) first studied resting correlations through
reference time course:

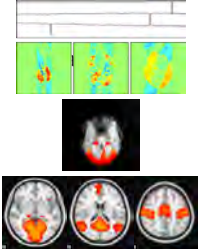
Functional Connectivity maps

Biswal et al. (1995) MRM

ICA

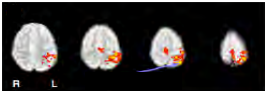


- Data-driven multivariate analysis.
- Good for finding:
 - Scanner and physiological
 - Activation with unknown characteristics
 - Resting state networks

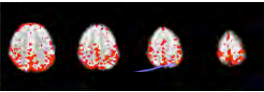


RSN analysis

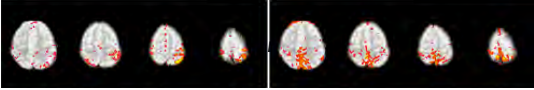
- Biswal (1995) first studied resting correlations through reference time course



Activation maps from a
finger tapping experiment



Correlation maps from a
resting state experiment

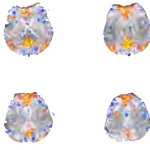


Spatial maps from an ICA decomposition

“Resting-State” Networks

Multiple spatial patterns of correlated temporal dynamics, resembling activation maps

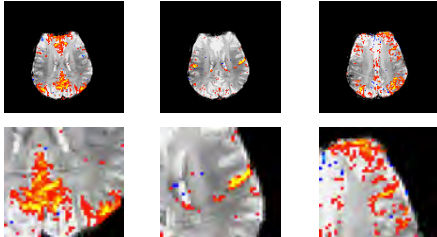
- can be found in fMRI data (BOLD & ASL) obtained at rest *and* in activation data
- characterised by (apparent) low frequency power spectra...
- seen when awake/sleep/anaesthesia, human/animals



Also referred to as: “low-frequency correlations”, “default activity”, “default mode”, “spontaneous network correlations”, “intrinsic connectivity networks” ...

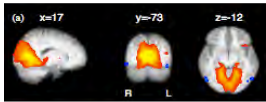
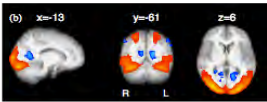

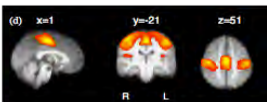
Spatial characteristics

- RSNs are grey-matter networks



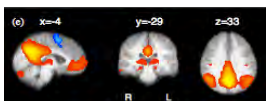
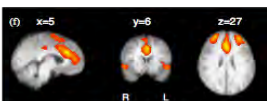
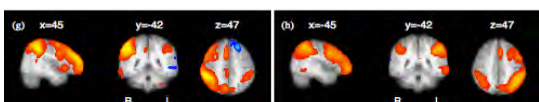
FSL

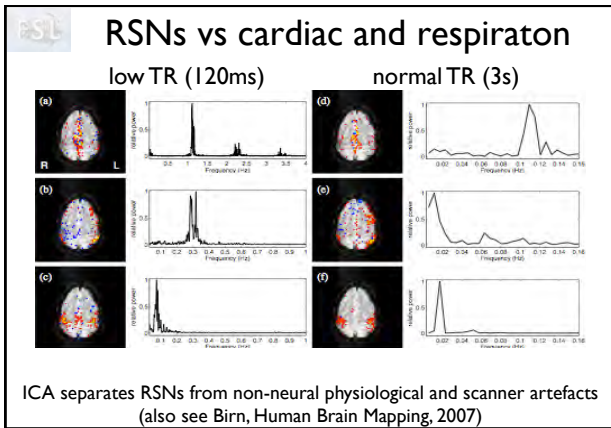
Spatial characteristics

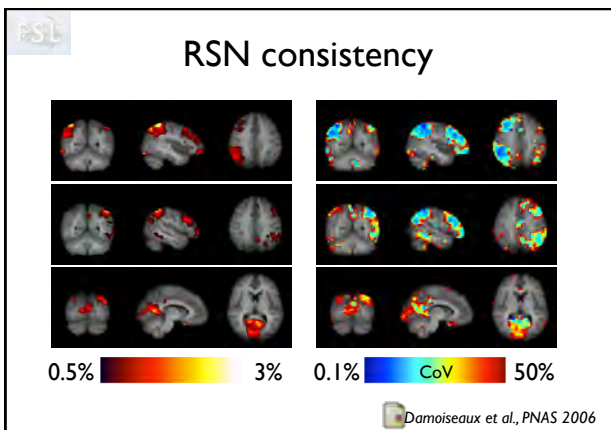
 <p>(a) $x=17$ $y=73$ $z=12$</p>			 <p>(b) $x=13$ $y=61$ $z=6$</p>		
Medial visual cortex			Lateral Visual Cortex		
 <p>(c) $x=3$ $y=-17$ $z=1.5$</p>			 <p>(d) $x=1$ $y=-21$ $z=51$</p>		
Auditory system			Sensory-motor system		

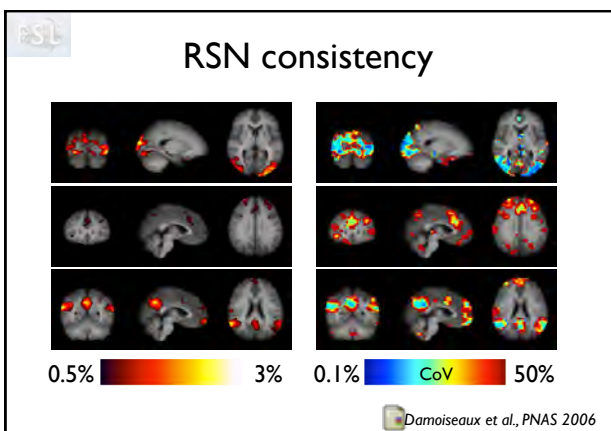
FSL

Spatial characteristics

 <p>(e) $x=-4$ $y=29$ $z=33$</p>			 <p>(f) $x=5$ $y=6$ $z=27$</p>		
'default mode network'			Executive control		
 <p>(g) $x=45$ $y=42$ $z=47$ (h) $x=45$ $y=42$ $z=47$</p>					
fronto-parietal					







functional & structural conn.

control

- absent contra-lateral connectivity in agenesis of CC

task seed R seed L

Quigley et al., Am J Neuroradiol, 2003

Parcellation of Thalamus

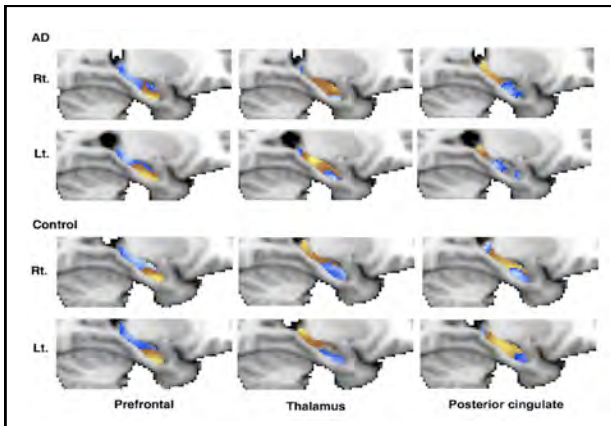
Behrens et al. Nat Neurosci 2003

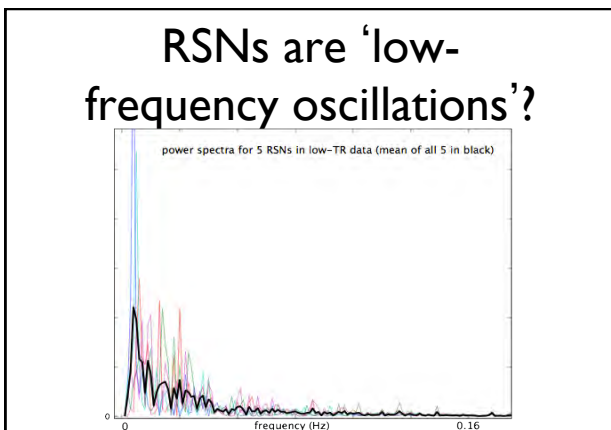
Zhang et al. J Neurophysiol 2008

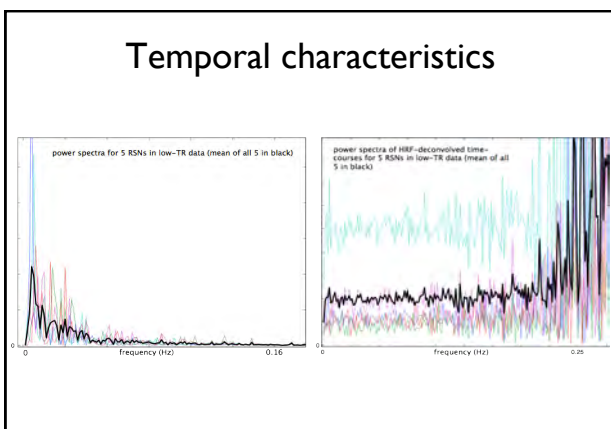
Target masks

- Visual (MTV)
- Sup. temporal auditory
- Somatosensory
- Motor and premotor
- Posterior parietal
- PFC

O'Reilly et al. Cereb Cort 2009

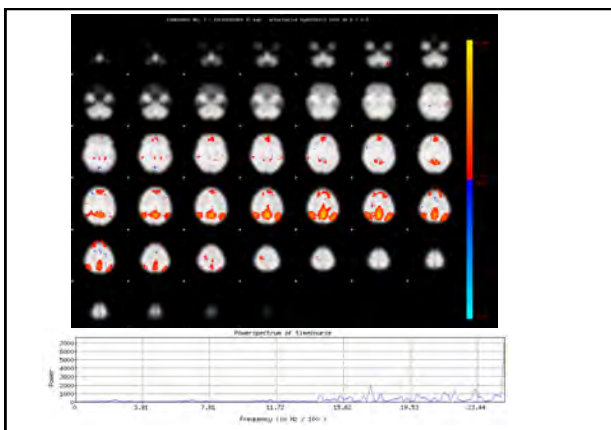






- Resting-state Networks are broadband processes!
- no 1/f characteristics!

Niazy, R et al. (2010)



The functional architecture of the human brain

Correspondence of the brain's functional architecture during activation and rest

Stephan M. Smith^{1,2}, Peter T. Fox³, Karla L. Miller⁴, David C. Glahn^{1,2}, P. Mickie Fox³, Clare E. Mackay³, Nicola Filippini¹, Kate E. Watkins¹, Roberto Toro⁵, Angela R. Laird⁶, and Christian F. Beckmann^{1*}


¹Centre for Functional MRI of the Brain, University of Oxford, Oxford, UK; ²Centre for Cognitive Neuroimaging, Research Imaging Centre, University of Texas Health Science Center, San Antonio, TX, USA; ³VA Medical Research Center, Institute of Living, Yale University, New Haven, CT, USA; ⁴Human Genetics and Cognitive Function, Medical Research Council, 225A Park Road, Farnham and Clinical Neuroimaging Research, Imperial College London, London SW7 2AZ, United Kingdom

Abstract Neural connections, providing the substrate for functional networks, exist whether or not they are functionally active at any given moment. However, it is not known to what extent brain regions are continuously interacting when the brain is at rest. In this work, we identify the major explicit activation networks by carrying out an image-based activation network analysis of thousands of separate activation maps derived from the BrainMap database of functional imaging studies, involving nearly 30,000 human subjects. Independently, we extract the major covering networks in the resting brain, as imaged with functional magnetic resonance imaging in 36 subjects at rest. The sets of major brain networks, and their dependencies into subnetworks, show close correspondence between the independent analyses of resting and activation brain dynamics. We conclude that the full repertoire of functional networks utilized by the brain in action is continuously and dynamically "active" even when at rest.

Brain connectivity: BrainMap: 1468; functional connectivity: resting state: network

Native images from the list of activation peak locations (13, 15) in all those available in coordinates space: coordinates of all those

The functional architecture of the human brain:
Correspondence between resting FMRI and task-activation studies



BrainMap, RIC, San Antonio

Data


- 1687 FMRI / PET activation studies (19% of all published activation imaging studies)
- 7342 separate activation conditions/contrasts
- 29,671 human subjects
- 66 "Behavioural domains" (gross paradigm classifications)

Resting FMRI data

Data

- 36 healthy subjects, age 20-35
- Subjects at rest, eyes open, fairly dark scanner room
- 6 minutes FMRI, standard BOLD, 3x3x3.5mm

Data & Methods



BrainMap, RIC, San Antonio

space ('unwrapped' voxels)

time (experiment-condition-ID 1/7342)

BrainMap data

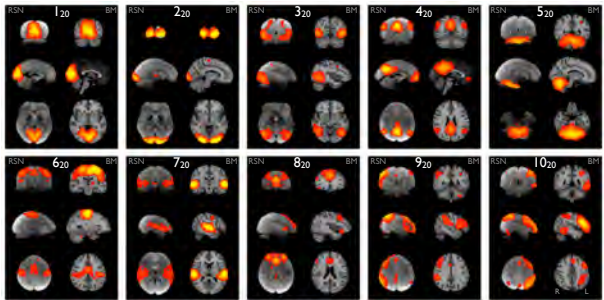
Resting FMRI data

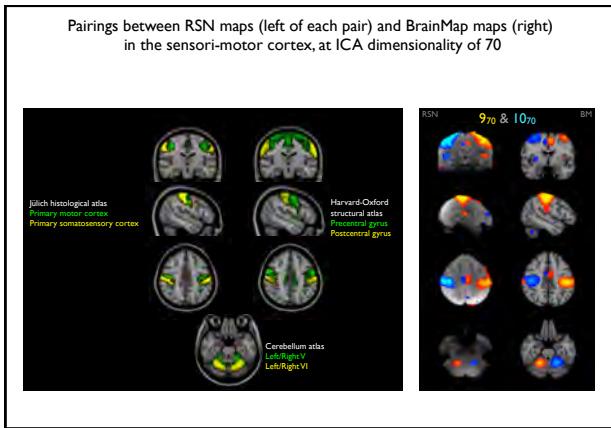
space ('unwrapped' voxels)

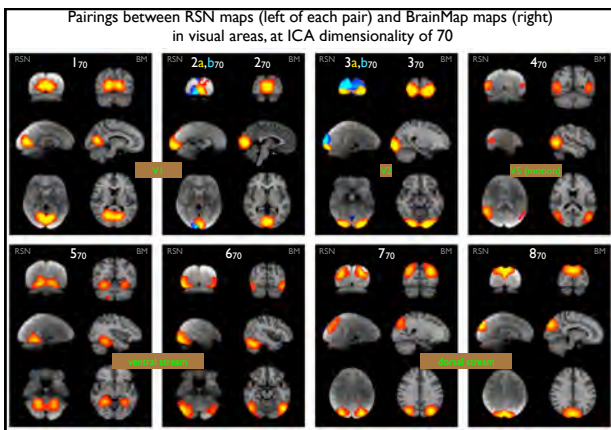
time

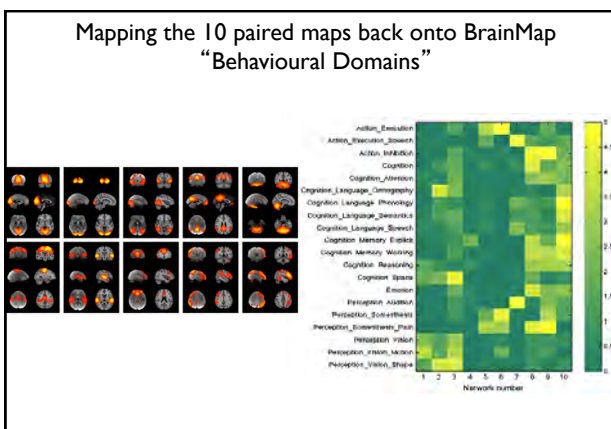
FMRI data

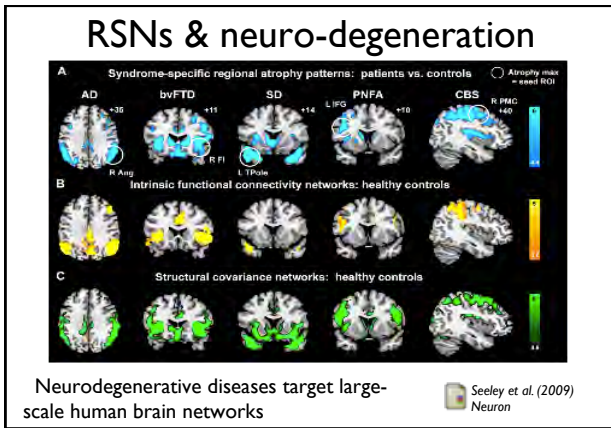
Pairings between RSN maps (left of each pair) and BrainMap maps (right) at ICA dimensionality of 20

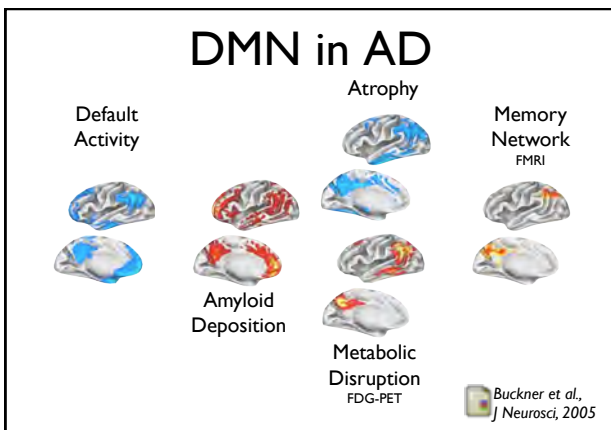


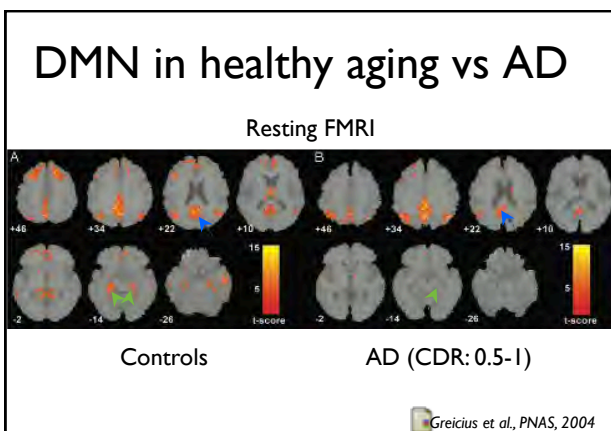












Altered functional connectivity in young, healthy carriers of APOE-ε4

Distinct patterns of brain activity in young carriers of the APOE-ε4 allele

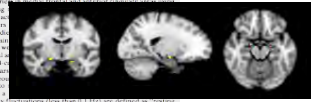
Nicola Filippi^{1,2*}, Bradley J. Macintosh³, Morgan G. Hough⁴, Guy M. Goodwin⁵, Giovanni B. Frisoni⁶, Stephen M. Smith⁷, Paul M. Matthews⁸, Christian F. Beckmann⁹, and Clare E. Mackay^{1,10}

¹University Department of Psychiatry and Functional Magnetic Resonance Imaging of the Brain Centre, University of Oxford, Oxford OX4 2DQ, United Kingdom; ²Department of Psychiatry, Massachusetts General Hospital, Harvard Medical School, Boston, MA 02114, USA; ³Department of Psychology, University of Cambridge, Cambridge CB2 3RQ, United Kingdom; ⁴Department of Clinical Neuroscience, Imperial College, Hammersmith Campus, London W12 0BQ, United Kingdom

*Edited by Robert D. Hoge, The Journal of Neuroscience, September 14, 2009; accepted March 10, 2009. This article includes supplemental material. Access to this article is provided at www.jneurosci.org. DOI: 10.1523/JNEUROSCI.1211-09.2009

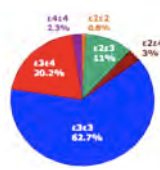
The APOE ε4 allele is a risk factor for late-life pathological changes that is also associated with anatomical and functional brain changes in midlife and elderly healthy subjects. We investigated structural and functional effects of the APOE polymorphism in 18 young healthy APOE ε4 carriers and 18 matched noncarriers (age range 20–35 years). Brain activity was studied both at rest and during an encoding memory paradigm using blood oxygen level-dependent (BOLD) resting fMRI revealed increased “default mode network” (DMN) activity in hippocampal, medial temporal, and medial prefrontal cortex at rest in ε4 carriers relative to noncarriers. The encoding task produced greater hippocampal activation in ε4 carriers relative to noncarriers. Further research could be explained by differences in memory performance, brain morphology, or resting cerebral blood flow. The APOE ε4 allele modulates brain function decades before any clinical or neurophysiological expression of neurodegenerative processes.

Keywords: memory; neuroimaging; resting connectivity



apolipoprotein E - APOE

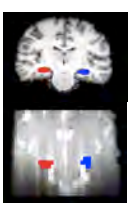
- lipid transport protein coded on chromosome 19
- ε 4 allele first studied as risk factor for cardiovascular disease
- ε 4 Identified as risk for AD in 1993
 - ▶ Increased prevalence
 - ▶ decreased age of onset
 - ▶ gene dose effect

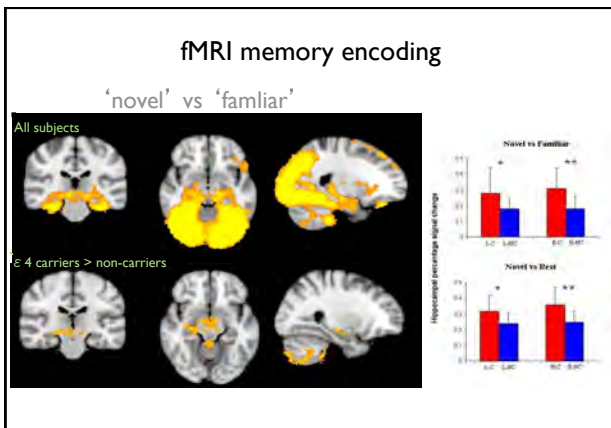


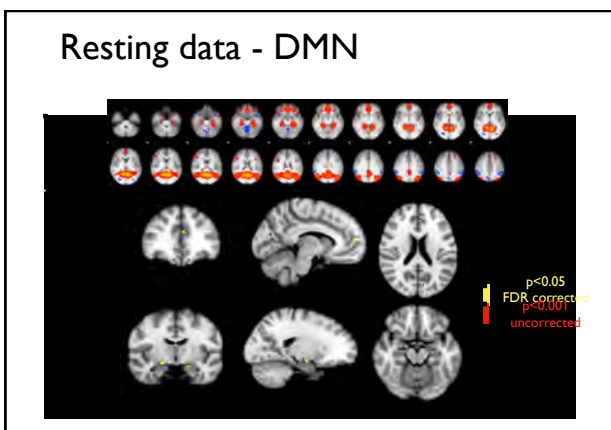
Menzel et al., 1983

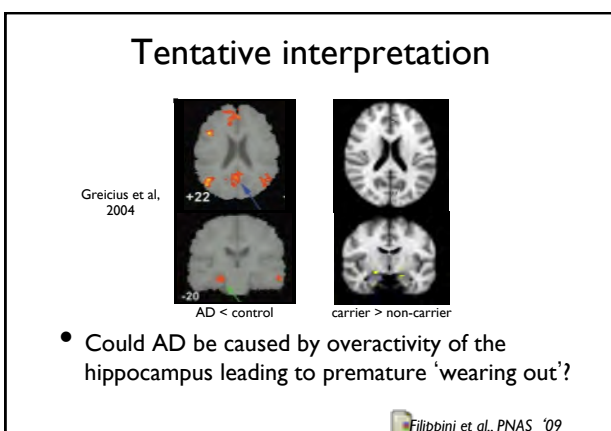
What is NOT different

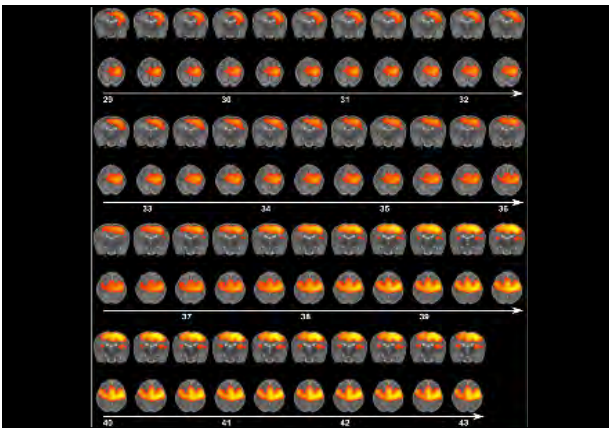
- Structure - Whole brain GM, WM and hippocampal ROIs & VBM not different
- Cerebral blood flow - ASL in hippocampal and lobe ROIs and whole brain not different
- Memory performance and reaction time not different

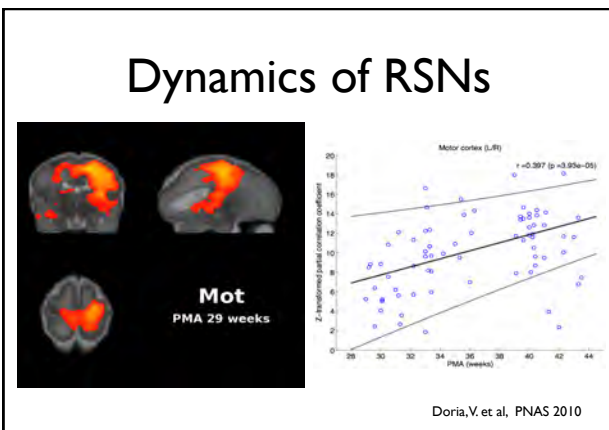


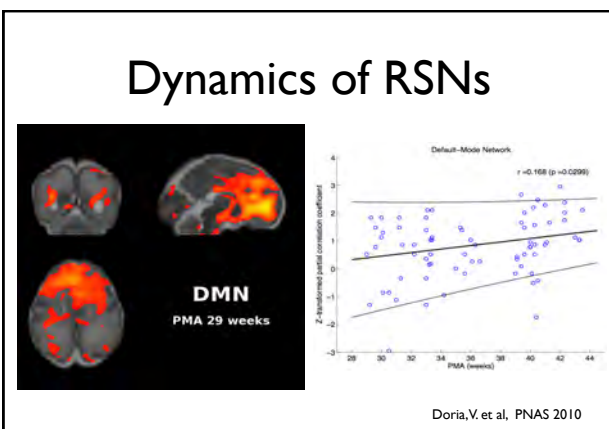










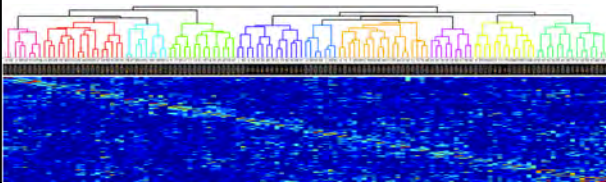


Hierarchy of patterns

- RSNs typically characterised by 2-10 components/clusters/networks
- relatively low-dimensional 'splitting' of the data
- consider more detailed hierarchy of functional connectivity by increasing ICA model order

Resting-FMRI Network

- 36-subjects' matrices (ICOV: direct functional connections)
- One-group t-test to give population-significant network matrix
- Will threshold at $p < 0.05$ (corrected for all possible connections)
- Feed into hierarchical clustering to reorder matrix and find (e.g) 10 functional sub-networks



"One man's noise is another man's signal"

- Resting-state FMRI promising technique, particularly in the clinical domain:
- does not require task compliance
- does not only probe specific cognitive functions but simultaneously investigates major *functional networks* which link to behaviour

Acknowledgements

- Steve Smith, Clare Mackay, Nicola Filippini, Mark Woolrich, Tim Behrens, Heidi Johansen-Berg (FMRIB Oxford)
- David Cole, Tomoki Arichi, David Edwards, Valentina Doria (Imperial College London)
- Serge Rombouts (LUMC Leiden)
- Rami Niazy (Rijad)
- Peter Fox, Angie Laird (RIC San Antonio)
