

# Introduction of fMRI Analysis

## 功能性磁振造影：影像分析簡介

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Specialization  
Representation  
Integration

## Answers to the Questions

**Brain activation** to the corresponding task.

- Where is the motion area?
- Where is the face recognition area?

**Brain encoding** to the representation of stimulus classes.

- What are the varying brain states in an area?
- How do brain cortices encode different types of information?

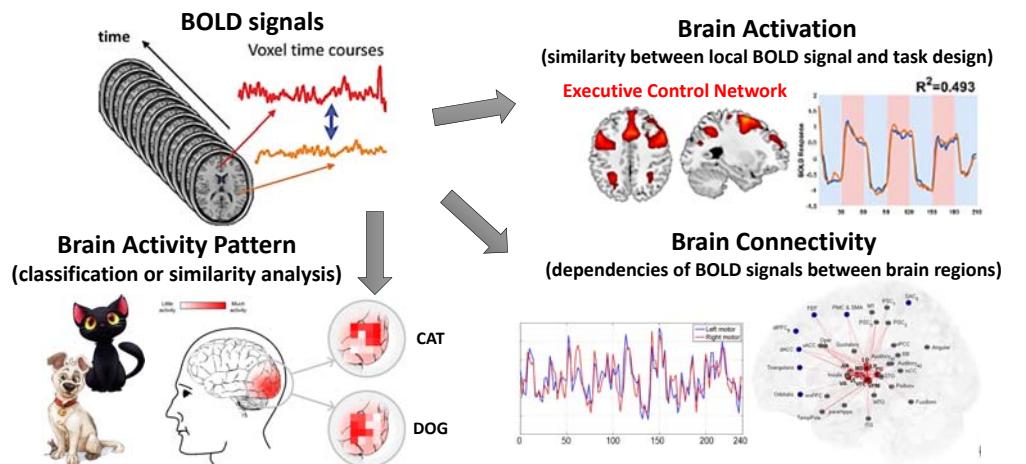
**Brain connectivity** to the integration of neural networks.

- How do neurons and neural networks process information?

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## fMRI Analysis



## fMRI Analysis

### • Brain Activation Analysis

- General Linear Model (GLM)

### • Brain Decoding:

- Multivariate Pattern Analysis (MVPA)
- Classifier-based MVPA, pattern similarity analysis

### • Brain Connectivity:

- Statistical dependency
- Independent Component Analysis (ICA)
- Network analysis

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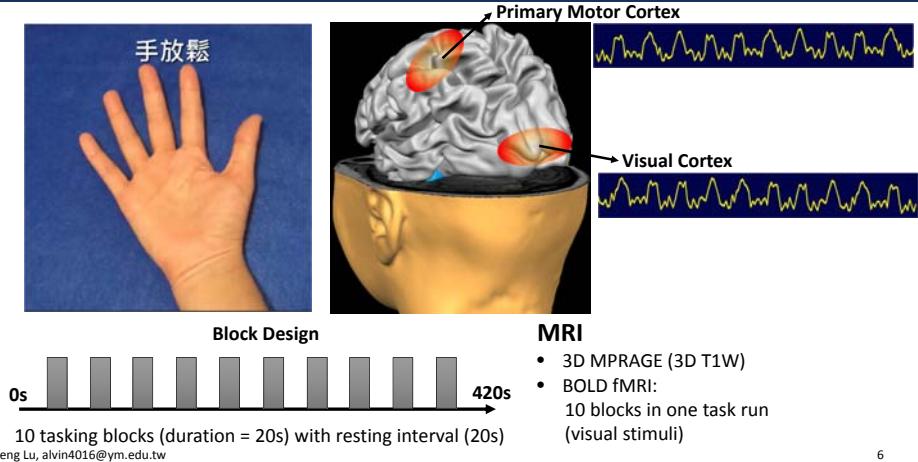
## Brain Activation Analysis

### General Linear Model (GLM)

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## Data Acquisition

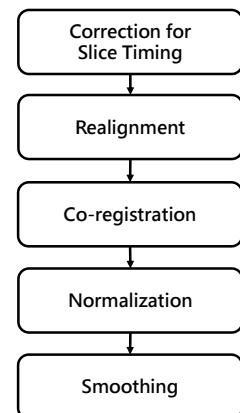


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## BOLD-fMRI Preprocessing



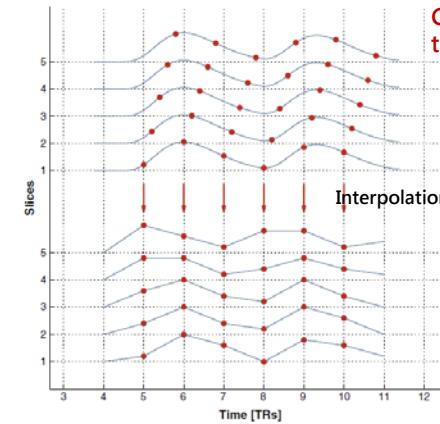
- Standard preprocessing steps for fMRI
  - Slice timing
  - Realignment
  - Co-registration (with anatomical images)
  - Normalization
  - Smoothing
  - Segment (tissue classification; optional)



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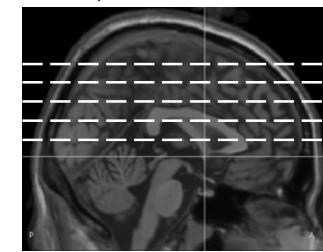
## Correction for Slice Timing



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Correction for slightly different imaging timing for multi-slice acquisition in a TR.

For example:  
Acquire 5 slices in 1 TR  
→ Temporal offset between slices



Sladky et al, NeuroImage 2011, 58:588-594.

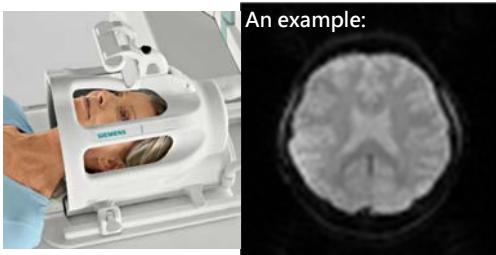
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## Realignment of head motion

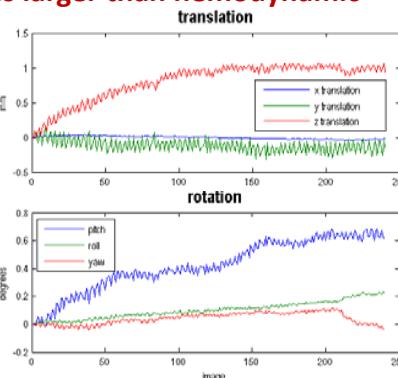


- The signal variation from movement is larger than hemodynamic response.

6-parameter Rigid body registration  
& transformation (align to the 1<sup>st</sup> volume)  
→ 6 co-variates for rs-fMRI analysis



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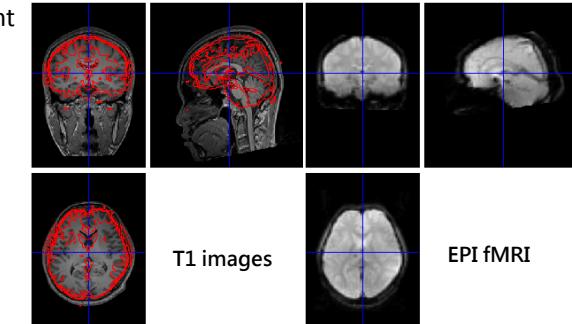
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## Co-registration



- Align fMRI (EPI) data to structural (T1) images.

- Rigid body transformation using mutual information
- Manual adjustment



T1 images

EPI fMRI

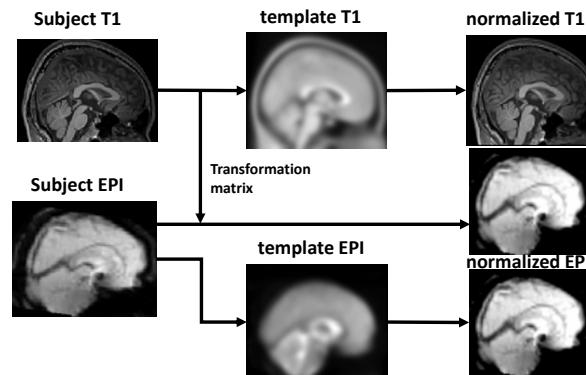
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## Spatial Normalization



- Perform spatial normalization using either anatomical (T1) images or fMRI (EPI) data.



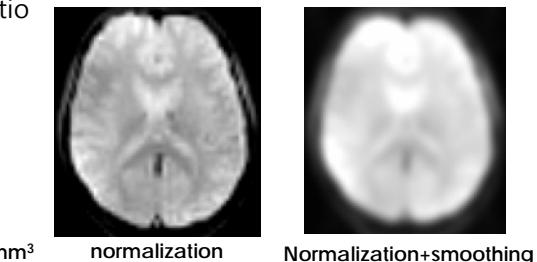
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## 3D Gaussian Smoothing



- Each voxel becomes weighted average of surrounding voxels.
- Render the data more normally distributed.
- Compensate for inaccuracies in normalization between individuals.
- Increase signal-to-noise ratio



2 x 2 x 2 mm<sup>3</sup> normalization

Normalization+smoothing

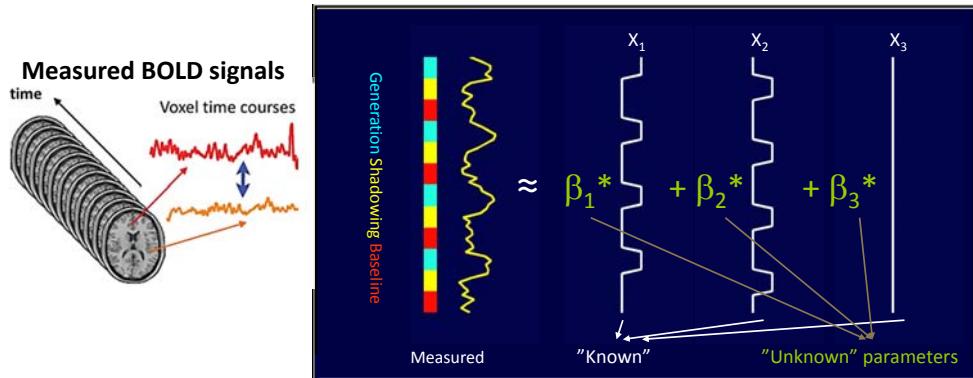
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# The Model of GLM



Finding the linear combination of these hypothetical time series "best" fits the data.



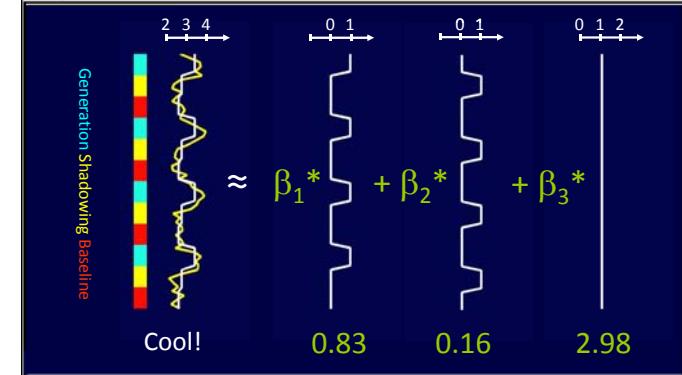
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# Parameter Estimation



Beta value represents the association between a condition design and the measured BOLD signal.



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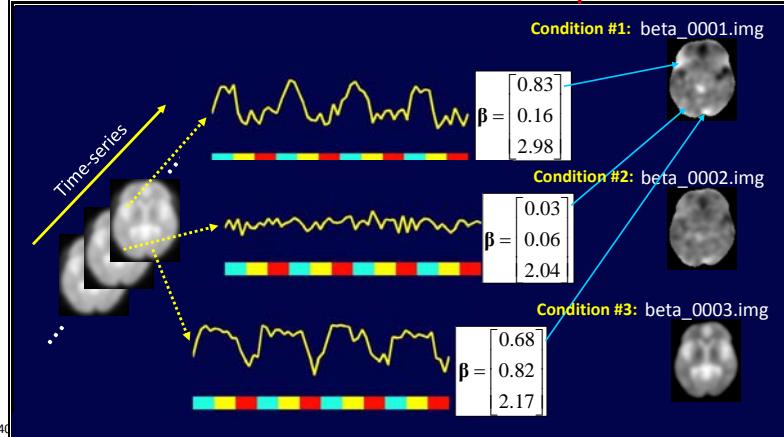
# Parameter Estimation



Same model for all voxels.

Different parameters for each voxel.

Beta map for each task condition



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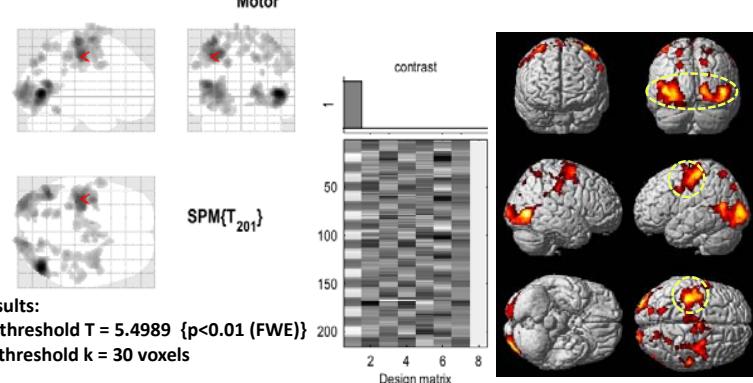
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# T-test: a simple example



Right hand grasping task

Q: activation during grasping?



$$t = \frac{c^T \hat{\beta}}{Std(c^T \hat{\beta})}$$

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## Available Softwares

- Statistical Parametric Mapping, SPM
  - <https://www.fil.ion.ucl.ac.uk/spm/>
- Analysis of Functional NeuroImages, AFNI
  - <https://afni.nimh.nih.gov/>
- FSL-FEAT
  - <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/FEAT>



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## Brain Decoding

### Multivariate Pattern Analysis (MVPA)

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## Decoding Activity Pattern of Brain

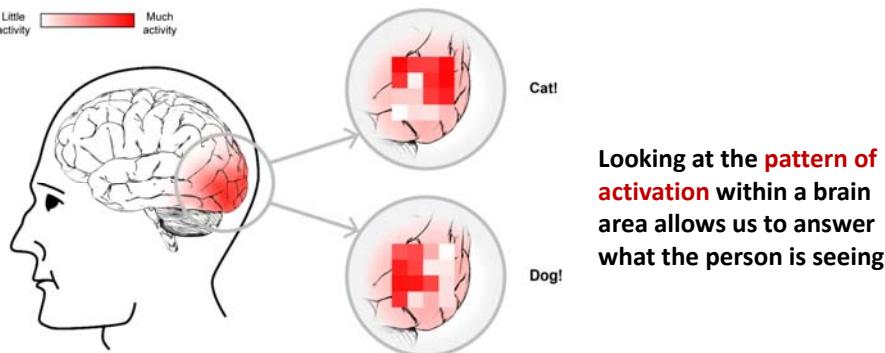


Illustration by Pim Mostert <http://blog.donders.ru.nl/?p=4361&lang=en>

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## Brain Activation → Brain Decoding

### • Mass-univariate model-based analysis

- Analyze every voxel (~50,000) one at a time
- General Linear Model, GLM (since 1995)

### • Multivoxel Pattern Analysis, MVPA

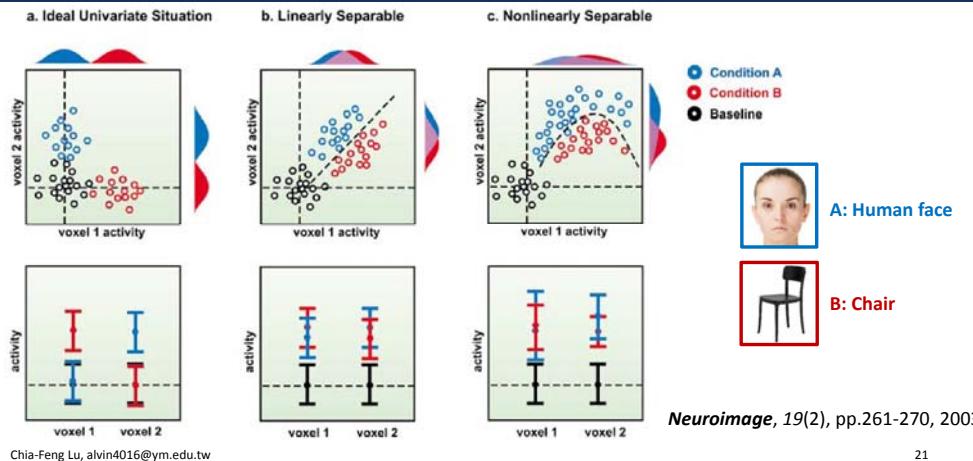
#### Multivariate Pattern Analysis, MVPA

- Original version: correlation analysis
- Machine learning: Support Vector Machine, SVM

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## Why we need multivariate analysis?



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## Major limitations of GLM

- The basic assumption that the covariance across neighboring voxels is not informative about the cognitive function under examination.
- Such covariance is considered as uncorrelated noise and normally reduced using spatial filters that smooth BOLD signals across neighboring voxels.
- Additionally, the GLM approach is inevitably limited by the model used for statistical inference.

→ Fail to capture “distributed” neural codes.

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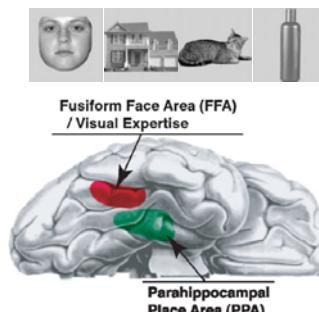
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## Origin of MVPA



Three hypothesis made by **James V. Haxby**, 2001.

- Each object category would evoke a distinct pattern of response in ventral temporal cortex.
- These distinctive patterns would not be restricted to category-selective regions, such as the FFA (face) and PPA (other objects).
- Neural activity patterns within category-selective regions would carry information that discriminates between categories.



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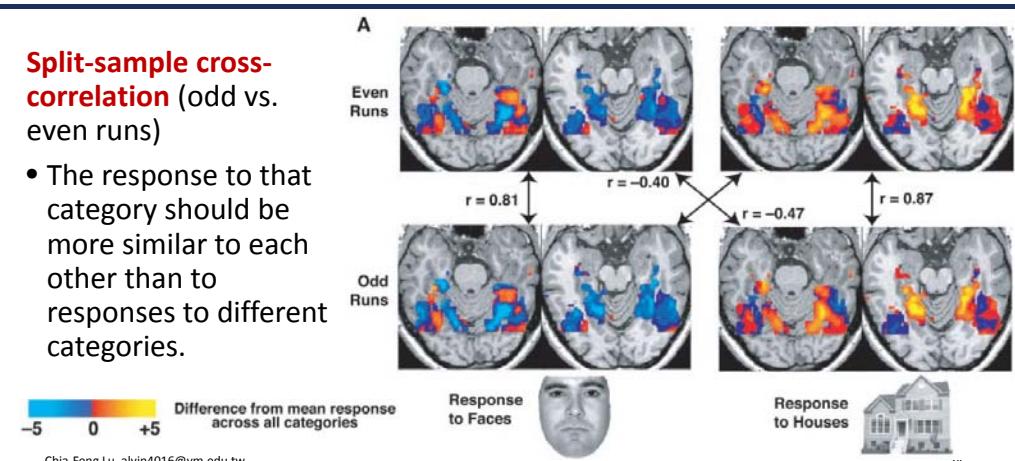
## Origin of MVPA



*Science*, 293(5539), 2425-2430, 2001.

### Split-sample cross-correlation (odd vs. even runs)

- The response to that category should be more similar to each other than to responses to different categories.



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## MVPA: A Classification Problem

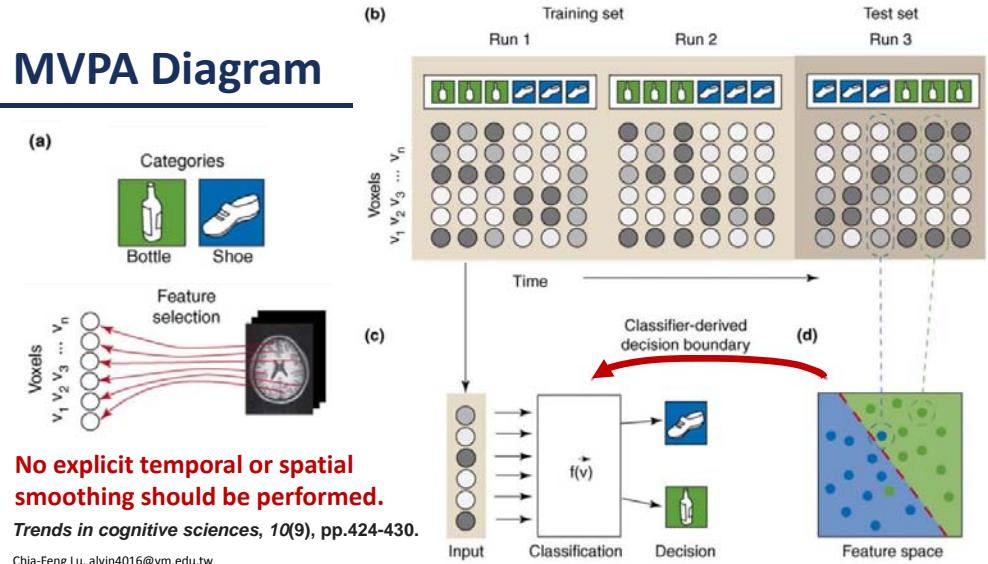
- Classification consists in determining a decision function  $f$  that takes the values of various “features” in a data “example”  $x$  and predicts the class of that “example.”
- An “example” may represent a given trial in the experimental run.
- The “features” may represent the corresponding fMRI signals in a cluster of voxels.
- The experimental conditions may represent the different “classes”.

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## MVPA Diagram

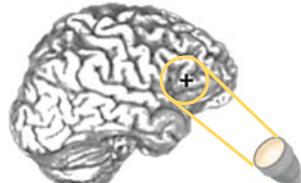


## Searchlight Approach

- Kriegeskorte, N., Goebel, R. and Bandettini, P., 2006. Information-based functional brain mapping. *Proceedings of the National Academy of Sciences*, 103(10), pp.3863-3868.

- Where in the brain does the activity pattern contain information about the experimental condition?

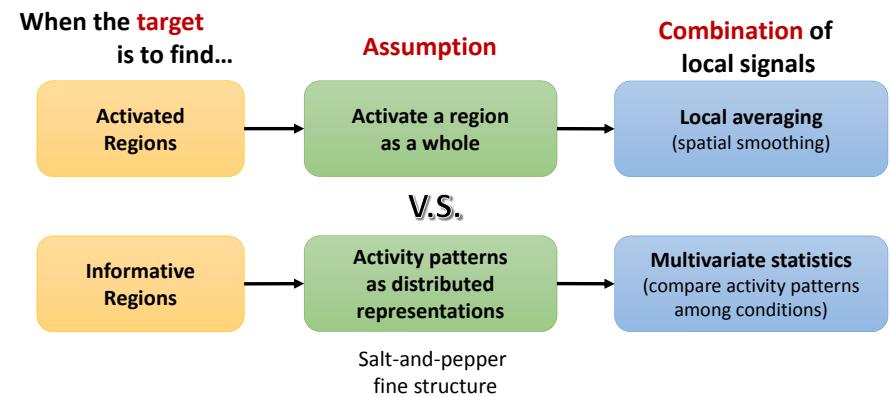
- Rather than asking where in the brain does the average activity changes across experimental condition.



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## Searchlight Approach



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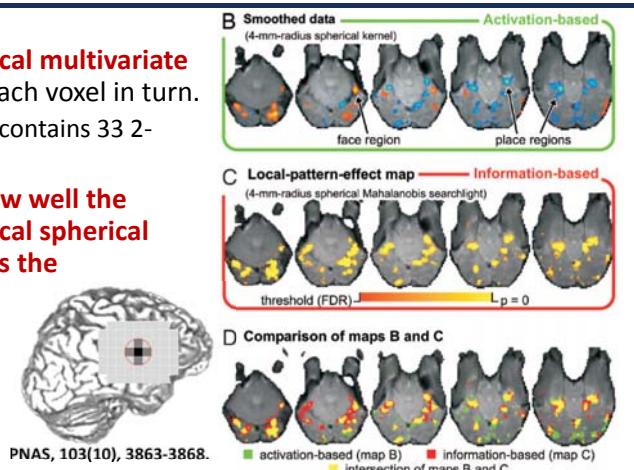
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## Searchlight Approach



- Scan the brain with a **spherical multivariate “searchlight”** centered on each voxel in turn.
    - An optimal radius of 4mm contains 33 2-mm-isotropic voxels.
  - The resulting map shows **how well the multivariate signal in the local spherical neighborhood differentiates the experimental conditions.**
    - Average absolute t value
    - Mahalanobis distance



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PNAS, 103(10), 3863-3868

activation-based (map B) ■ information-based (map C)  
● intersection of maps B and C

## Toolbox of MVPA



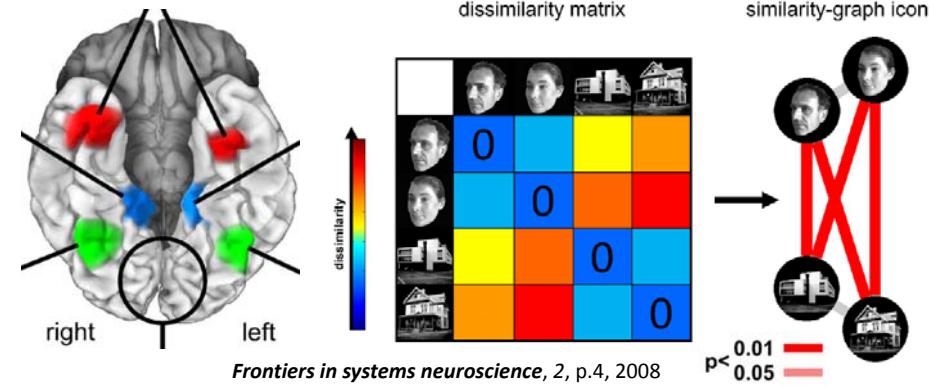
- Hanke, M., Halchenko, Y.O., Sederberg, P.B., Hanson, S.J., Haxby, J.V., Pollman, S., 2009. PyMVPA: a Python toolbox for multivariate pattern analysis of fMRI data. *Neuroinformatics* 7, 37–53.
  - An MVPA toolbox using Matlab (the Princeton MVPA toolbox) (<http://code.google.com/p/princeton-mvpa-toolbox/>).
  - Oosterhof, N.N., Connolly, A.C. and Haxby, J.V., 2016. CoSMoMVPA: multi-modal multivariate pattern analysis of neuroimaging data in Matlab/GNU Octave. *Frontiers in neuroinformatics*, 10, p.27.

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# Pattern Similarity Analysis



Compared between regions, species, mental states, and diseases.



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## Brain Connectivity

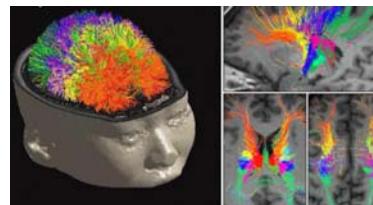
# Statistical Dependency and Network Analysis

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## Brain Connectivity

- **Structural connectivity**
  - Anatomical links (neuronal fibers)
- **Functional connectivity**
  - Statistical dependency
- **Effective connectivity**
  - Causal interactions between distinct units

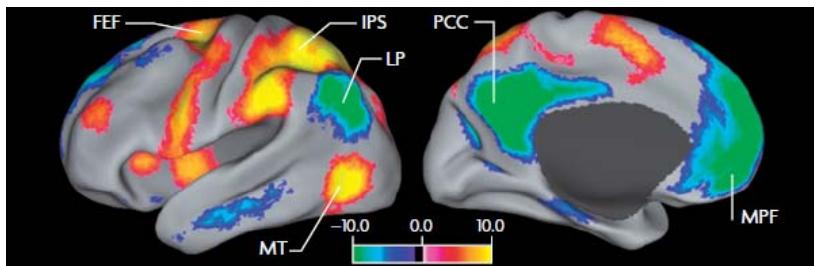


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## Default Mode Network

- During resting state scans, there are two networks in which areas are correlated with each other and anticorrelated with areas in the other network



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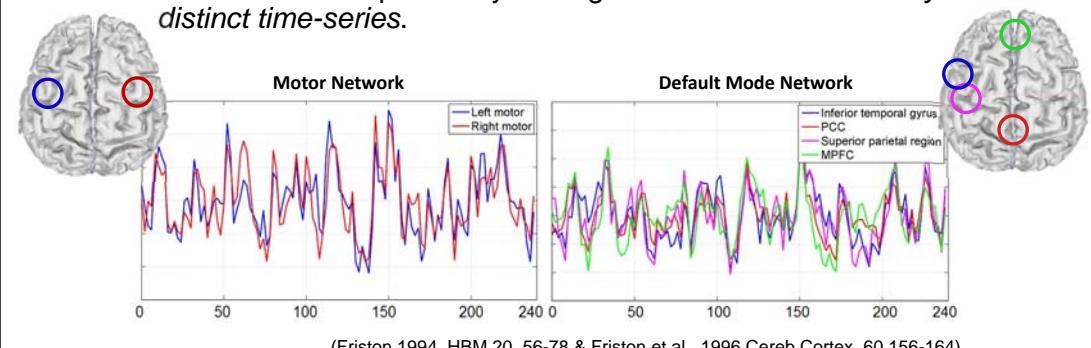
Fox and Raichle, 2007, Nat. Rev. Neurosci.

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## Functional connectivity

- Functional connectivity (FC) is defined as the statistical association or dependency *among two or more anatomically distinct time-series*.



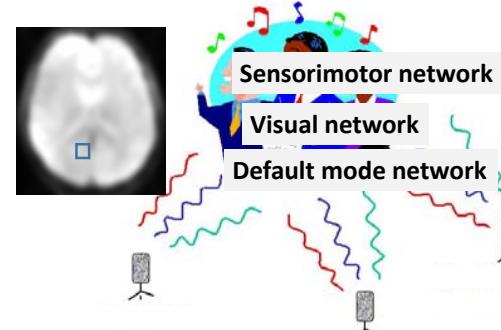
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## Independent Component Analysis

- **A cocktail-party problem**



Number of microphones  $\geq$  number of speakers

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$$\begin{bmatrix} x_1(t) \\ x_2(t) \\ \vdots \\ x_m(t) \end{bmatrix} = \mathbf{A} \begin{bmatrix} s_1(t) \\ s_2(t) \\ \vdots \\ s_k(t) \end{bmatrix}$$

$$\begin{bmatrix} x_1(t) \\ x_2(t) \\ \vdots \\ x_m(t) \end{bmatrix} = \mathbf{A} \begin{bmatrix} s_1(t) \\ s_2(t) \\ \vdots \\ s_k(t) \end{bmatrix}$$

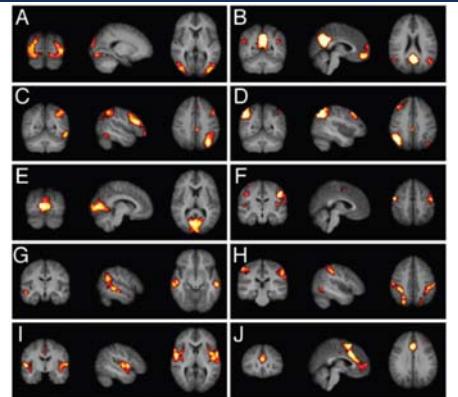
$\mathbf{A}$  is the mixing matrix



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## Independent Components

- A. parts of visual cortex
- B. default mode network
- C & D. left and right memory function
- E. visual cortex
- F. sensorimotor cortex
- G. occipitotemporal pathway (ventral stream)
- H. superior parietal cortex
- I. auditory cortex
- J. executive control & working memory

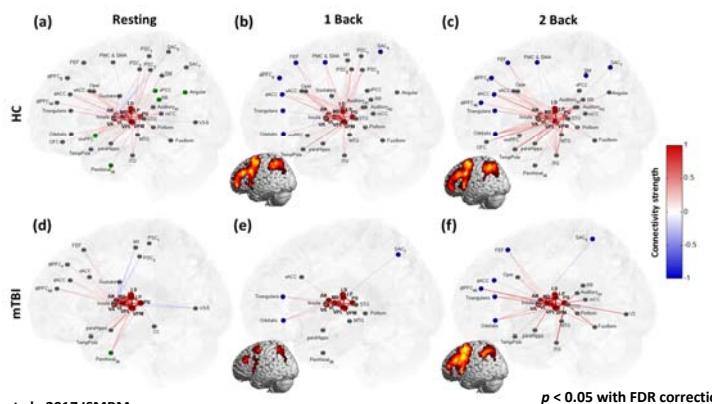


Consistent resting-state networks across healthy subjects. PNAS 2006, 103(37): 13848-13853.

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## Concussion-related alterations of FC



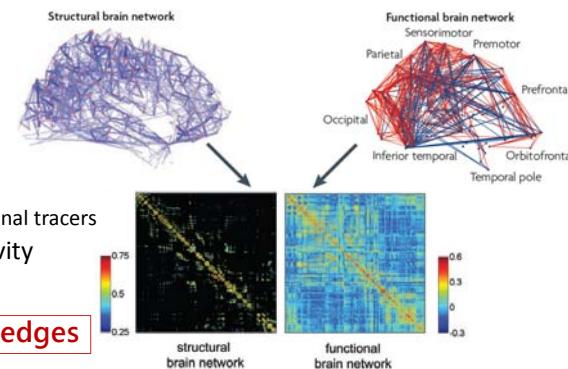
Lu et al., 2017 ISMRM.

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## Network construction

- Nodes**
  - Cortical regions
- Edges**
  - Cortical thickness correlations
  - Fiber connections
    - DSI, DTI, transneuronal tracers
  - Functional connectivity
    - fMRI, EEG, MEG



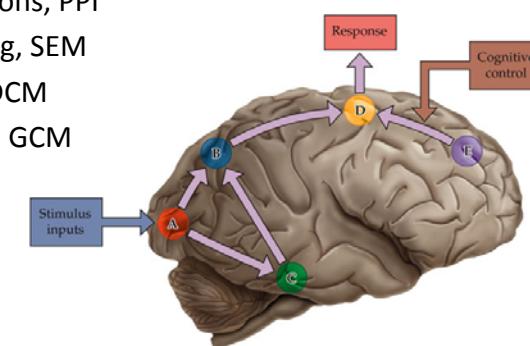
Network = nodes + edges

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## Correlation ≠ Causality

- Psychophysiological Interactions, PPI
- Structural Equation Modelling, SEM
- Dynamic Causal Modelling, DCM
- Granger Causality Modelling, GCM



Functional Magnetic Resonance Imaging 2e, Figure 11.10

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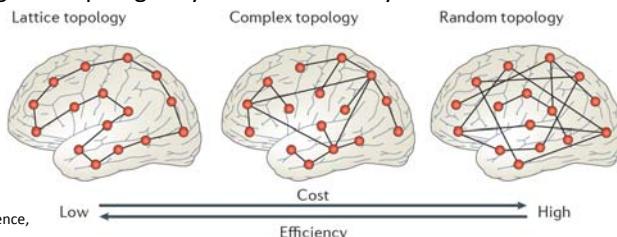
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## Human brain networks

Wiring costs ⇔ efficiency



- Clusters of lattice-like short-distance connections between spatially neighboring nodes
- Topologically direct interconnections between spatially remote brain regions → increase efficiency of information processing
- Nodes aggregated topologically and anatomically as modules → minimize wiring cost



Nature Reviews Neuroscience, 13: 336-349, 2012.

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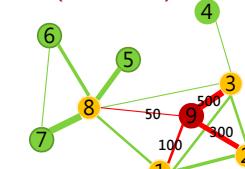
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### A small-world architecture

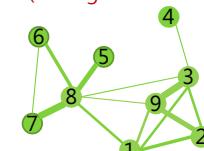


## Graph theory: topological properties

Local (each node)



Global (average over all nodes)



- **degree** (the number of neighbors)  
e.g. degree of node 9 = 4
- **strength** (the connected fiber number\*FA)  
e.g. strength of node 9  
 $= (50+100+300+500)/4 = 237.5$
- **clustering coefficient**  
(the connection between neighbors, [0~1])  
e.g. clustering coefficient of node 9  
 $= 5/6 = 0.83$
- **path length (separation)**  
(the minimal steps for connection)  
e.g. path length from node 9 to node 6  
 $= 2 \text{ steps } (9 \rightarrow 8 \rightarrow 6)$

Philos Trans R Soc Lond B Biol Sci, 360, 937-946, 2005

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## Available Softwares

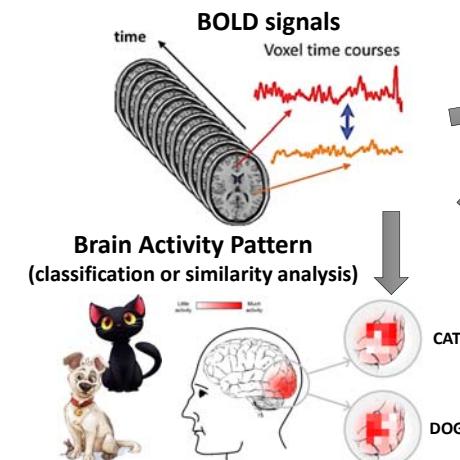


- Group ICA of fMRI Toolbox, GIFT
  - <http://mialab.mrn.org/software/gift/>
- FSL-MELODIC
  - <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/MELODIC>
- Statistical Parametric Mapping, SPM
  - <https://www.fil.ion.ucl.ac.uk/spm/>
- Data Processing & Analysis for Brain Imaging (DPABI)
  - <http://rfmri.org/dpabi>
- GRETA Toolbox
  - <https://www.nitrc.org/projects/gretna/>
- Brain Connectivity Toolbox
  - <https://sites.google.com/site/bctnet/>

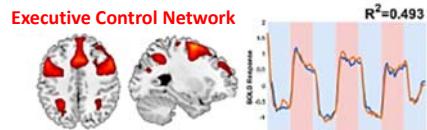
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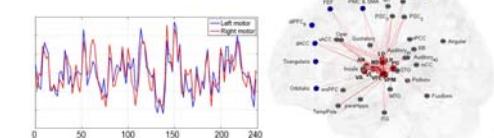
## fMRI Analysis



**Brain Activation**  
(similarity between local BOLD signal and task design)



**Brain Connectivity**  
(dependencies of BOLD signals between brain regions)





# THE END

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**fMRI Teaching Materials:**

[http://www.ym.edu.tw/~cflu/CFLu\\_course\\_fMRIana.html](http://www.ym.edu.tw/~cflu/CFLu_course_fMRIana.html)