

Extracting fMRI features

PRoNTo course
May 2018

Overview

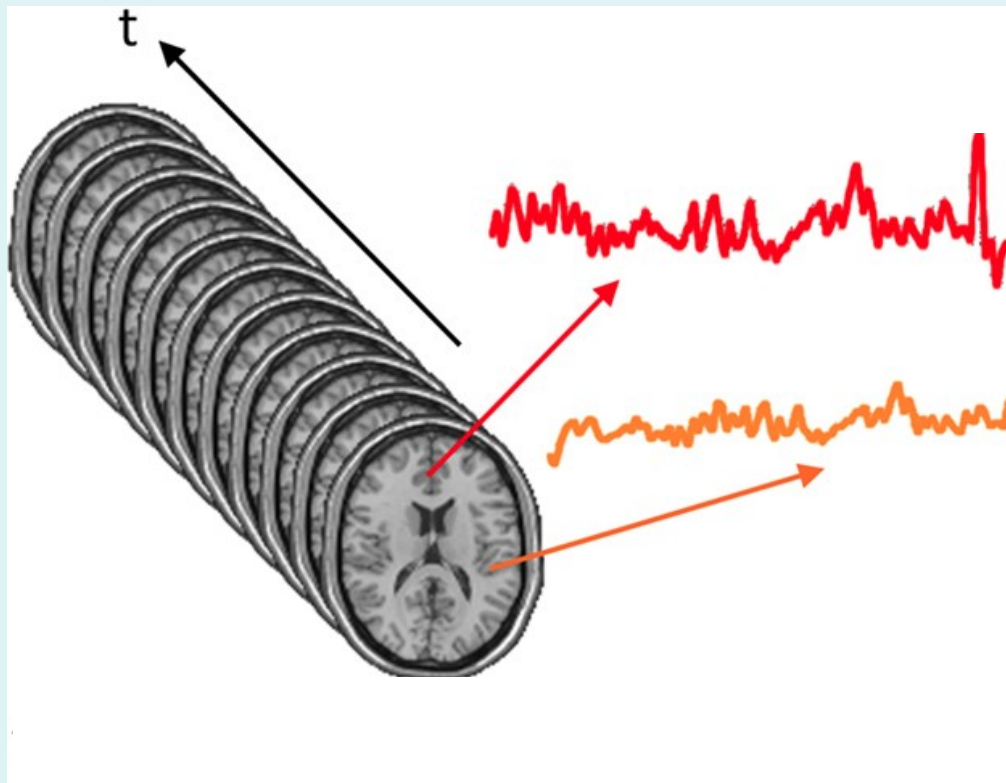
- Introduction
- “Brain decoding” problem
- “Subject prediction” problem
- Conclusion

Overview

- Introduction
 - fMRI data & processing
 - GLM & BOLD response
 - classical univariate approach
 - levels of inference
- “Brain decoding” problem
- “Subject prediction” problem
- Conclusion

Data

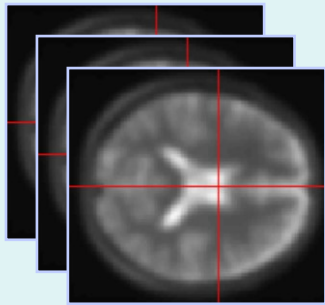
fMRI time series = 4D image
= time series of 3D fMRI's
= 3D array of time series.



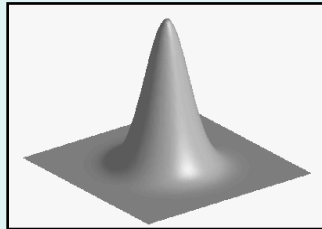
About the same for a series of structural MRI's...

Spatial pre-processing & SPM

Image time-series



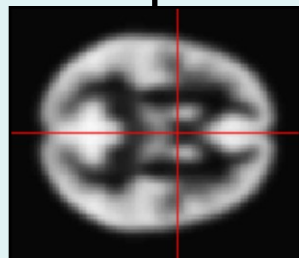
Spatial filter



Realignment

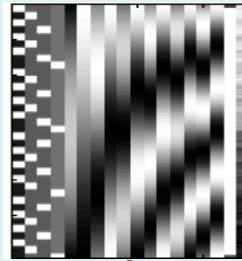
Smoothing

Normalisation

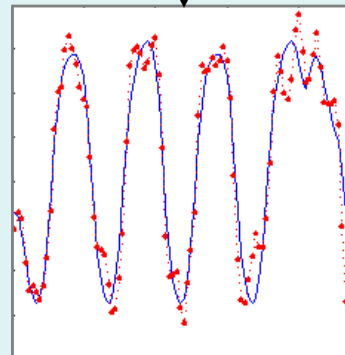


Anatomical reference

Design matrix

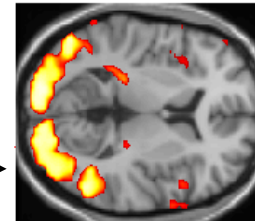
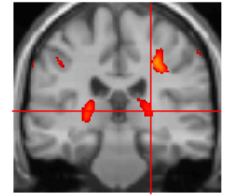
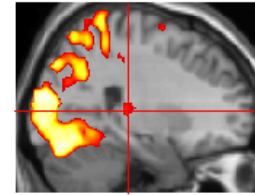


General Linear Model



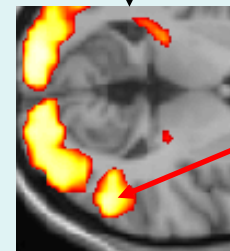
Parameter estimates

Statistical Parametric Map



Statistical Inference

RFT



$p < 0.05$

GLM *univariate* approach

$$Y = X\beta + \varepsilon$$

N : # images
 p : # regressors

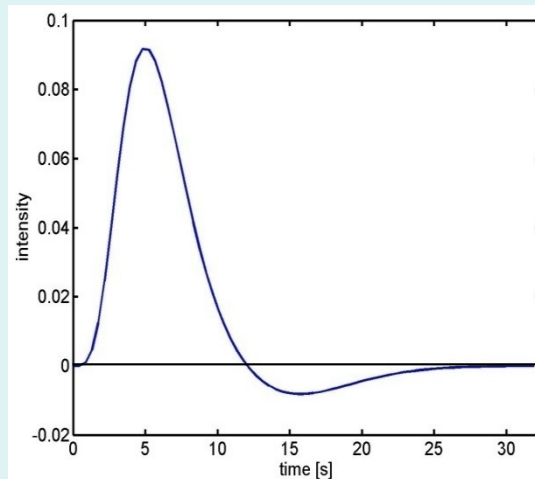
GLM defined by:

- design matrix X
- error term ε distribution, e.g.

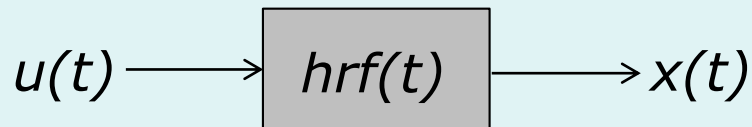
$$\varepsilon \sim N(0, \sigma V)$$

BOLD response

Hemodynamic response function (HRF):



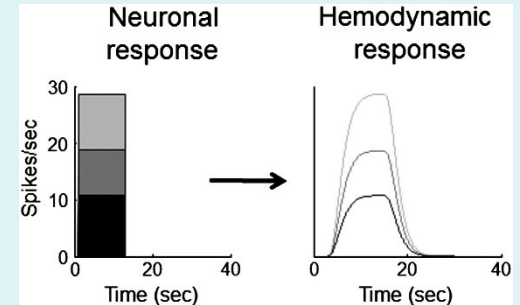
Linear time-invariant (LTI) system:



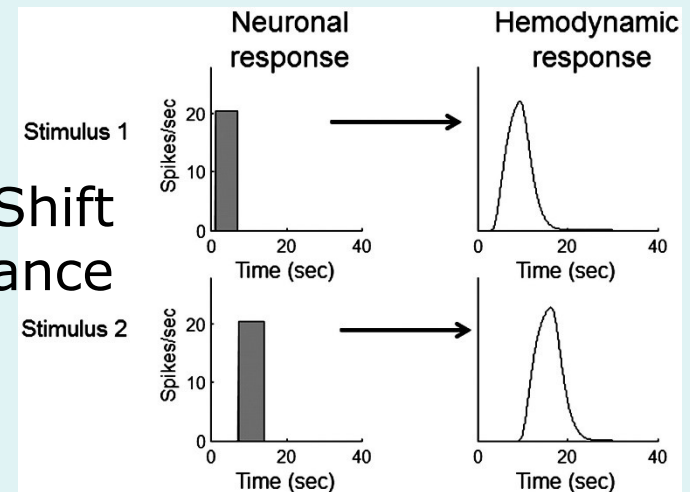
Convolution operator:

$$\begin{aligned} x(t) &= u(t) * hrf(t) \\ &= \int_0^t u(\tau) hrf(t - \tau) d\tau \end{aligned}$$

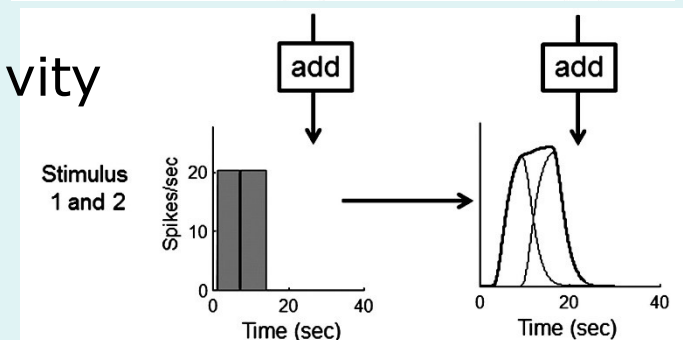
Scaling



Shift invariance



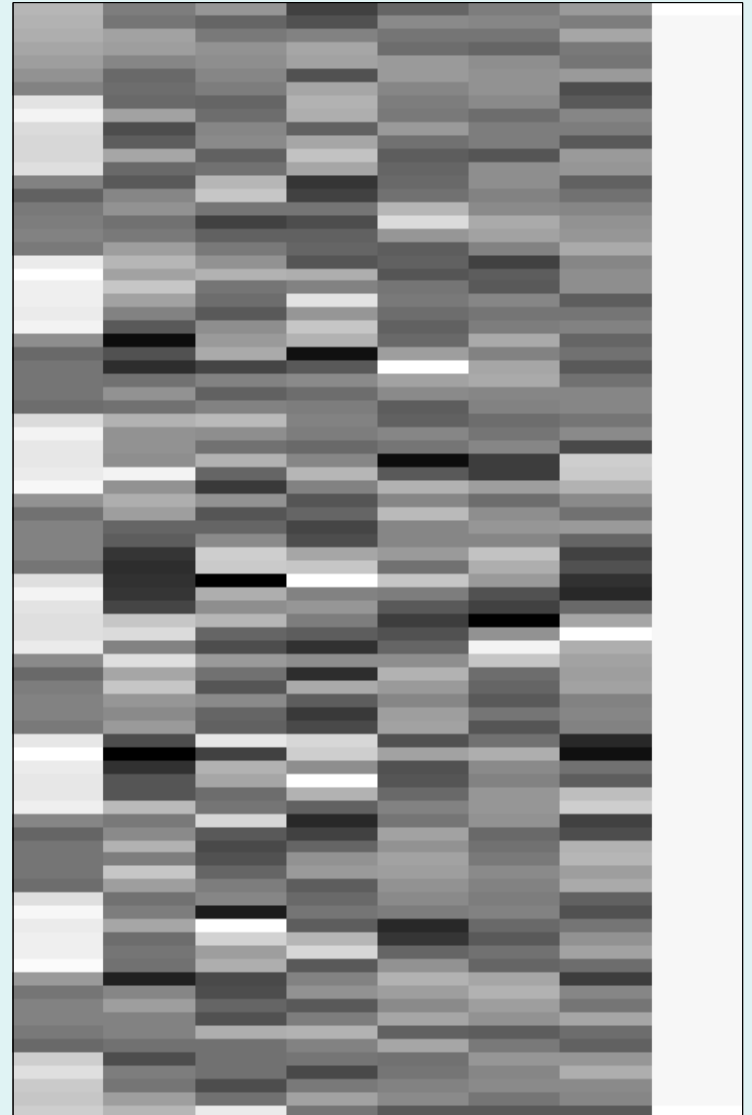
Additivity



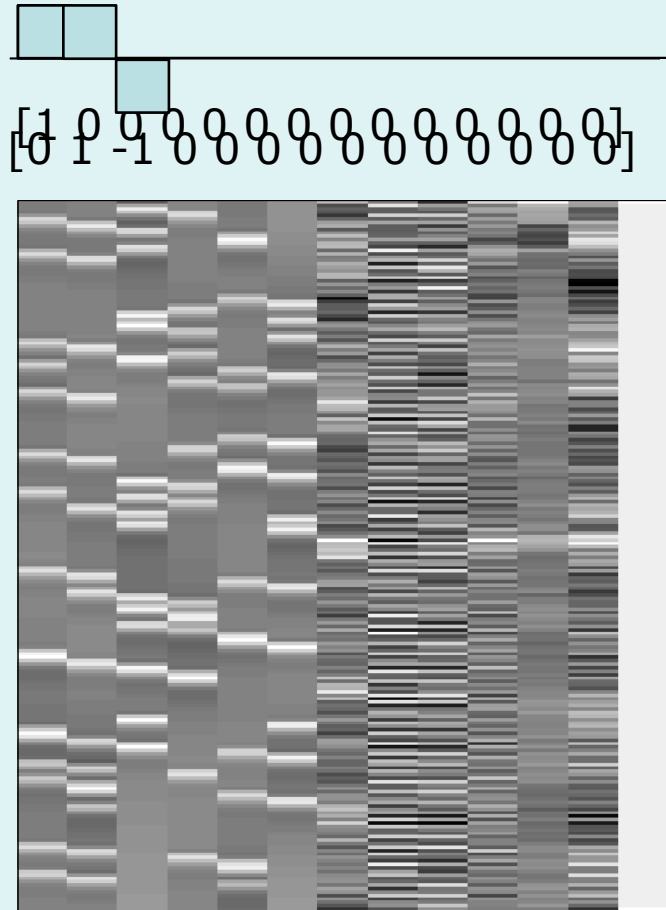
Confounds modelling

Model

- condition(s) of interest, i.e. “activation”
- “confounds”:
 - BOLD response
 - estimated movement parameters



Contrast & inference



A contrast selects a **specific effect of interest**.

⇒ A contrast c is a vector of length p .

$\Rightarrow c^T \beta$ is a linear combination of regression coefficients β .

$$c = [1 \ 0 \ 0 \ 0 \ \dots]^T$$

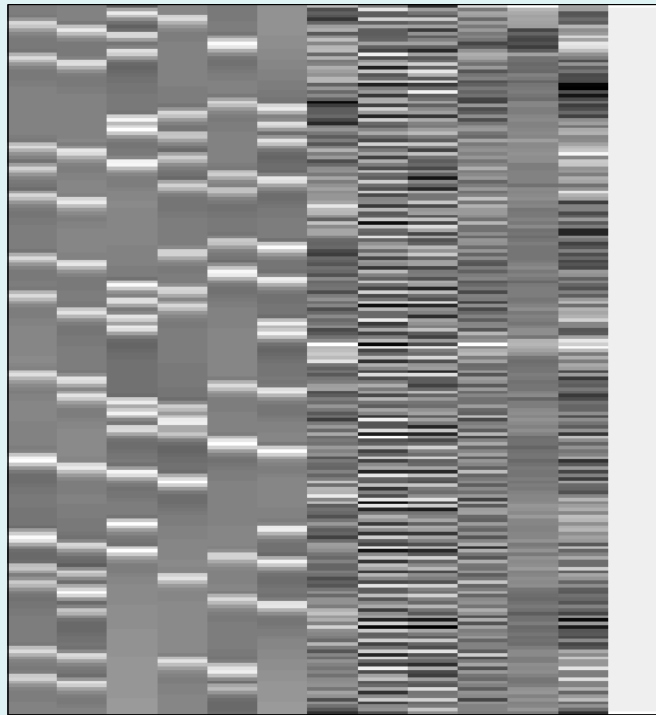
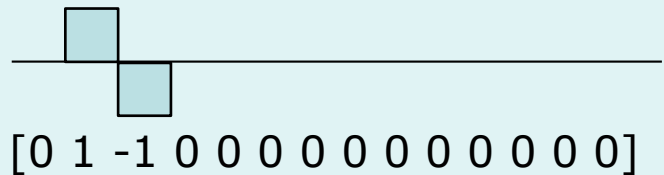
$$c^T \beta = \mathbf{1} \times \beta_1 + \mathbf{0} \times \beta_2 + \mathbf{0} \times \beta_3 + \mathbf{0} \times \beta_4 + \dots$$
$$= \beta_1$$

$$c = [0 \ 1 \ -1 \ 0 \ \dots]^T$$

$$\begin{aligned} c^T \beta &= \mathbf{0} \times \beta_1 + \mathbf{1} \times \beta_2 + \mathbf{-1} \times \beta_3 + \mathbf{0} \times \beta_4 + \dots \\ &= \beta_2 - \beta_3 \end{aligned}$$

$$c^T \hat{\beta} \sim N(c^T \beta, \sigma^2 c^T (X^T X)^{-1} c)$$

Contrast & inference



Question: β_2 larger than β_3 ?
=
 $\beta_2 - \beta_3 = c^T \beta > 0$?

Null hypothesis: $H_0: c^T \beta = 0$

Test statistic: $T = \frac{\text{contrast of estimated parameters}}{\sqrt{\text{variance estimate}}}$

$$T = \frac{c^T \hat{\beta}}{\sqrt{\text{var}(c^T \hat{\beta})}} = \frac{c^T \hat{\beta}}{\sqrt{\hat{\sigma}^2 c^T (X^T X)^{-1} c}} \sim t_{N-p}$$

Classical inference

The Null Hypothesis H_0

= what we want to disprove (no effect).

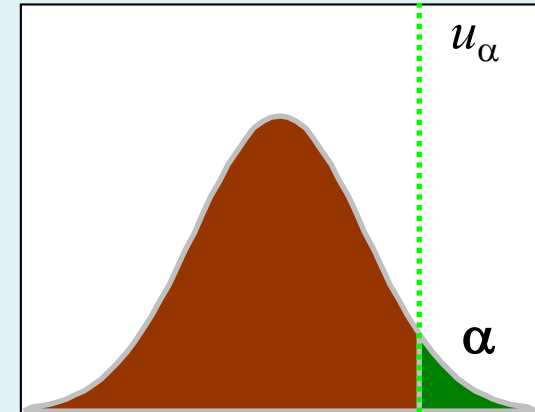
⇒ The Alternative Hypothesis H_A
= outcome of interest.

Significance level α :

Acceptable *false positive rate* α .

⇒ threshold u_α

$$\alpha = p(T > u_\alpha \mid H_0)$$



Null Distribution of T

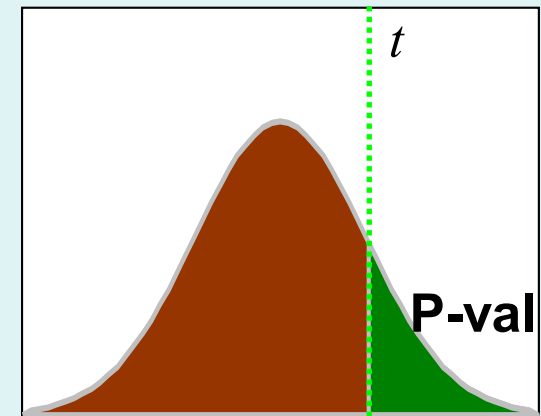
Observation of test statistic t

= a realisation of T

⇒ Reject H_0 in favour of H_A if $t > u_\alpha$

p -value = evidence against H_0

$$p(T > t \mid H_0)$$

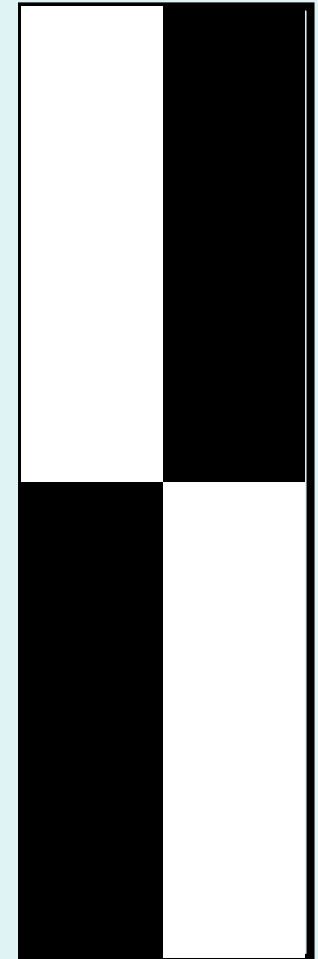
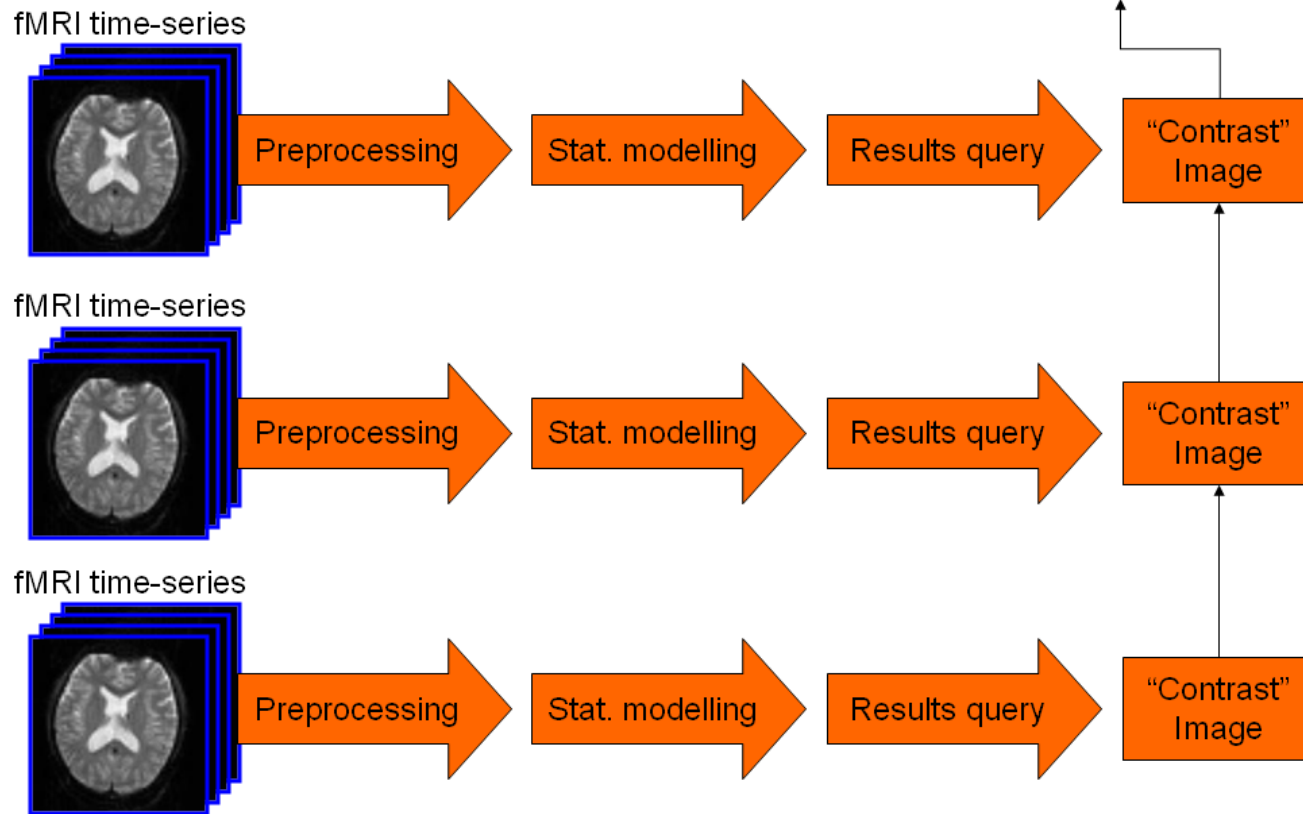


Null Distribution of T

Group-level analysis

Group A vs. Group B design

SPM for group fMRI



"Summary statistics" approach.

Overview

- Introduction
- “Brain decoding” problem
 - BOLD signal & HRF
 - Raw signal
 - Beta images
- “Subject prediction” problem
- Conclusion

Brain decoding

Data:

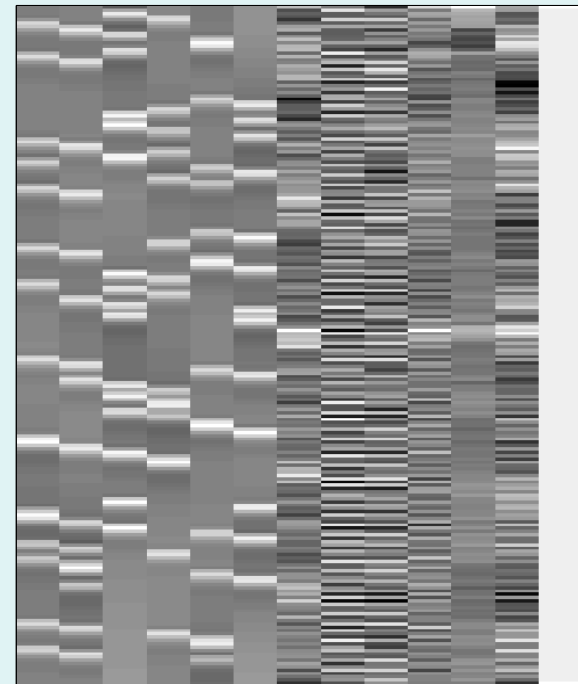
fMRI time series from 1 (or a few) subject(s)

Goal:

Find temporary mental state, from fixed set, of a subject based on pattern of brain activity.

→ decode BOLD signal over 1 or few images

Similar to FFX analysis

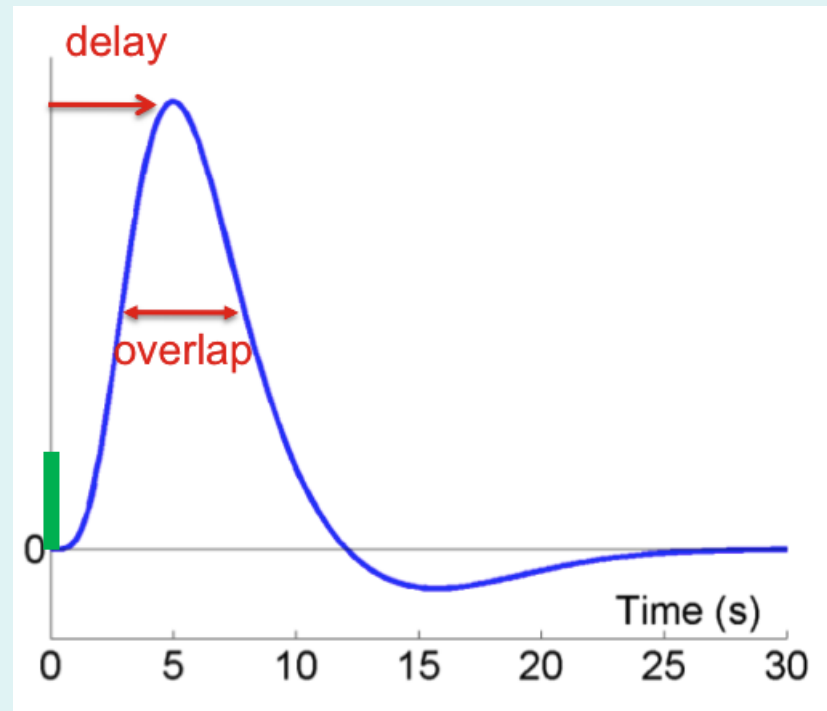


Brain decoding: signal

Use the raw BOLD signal but

- Block or event-related design?
- How to account for haemodynamic function?

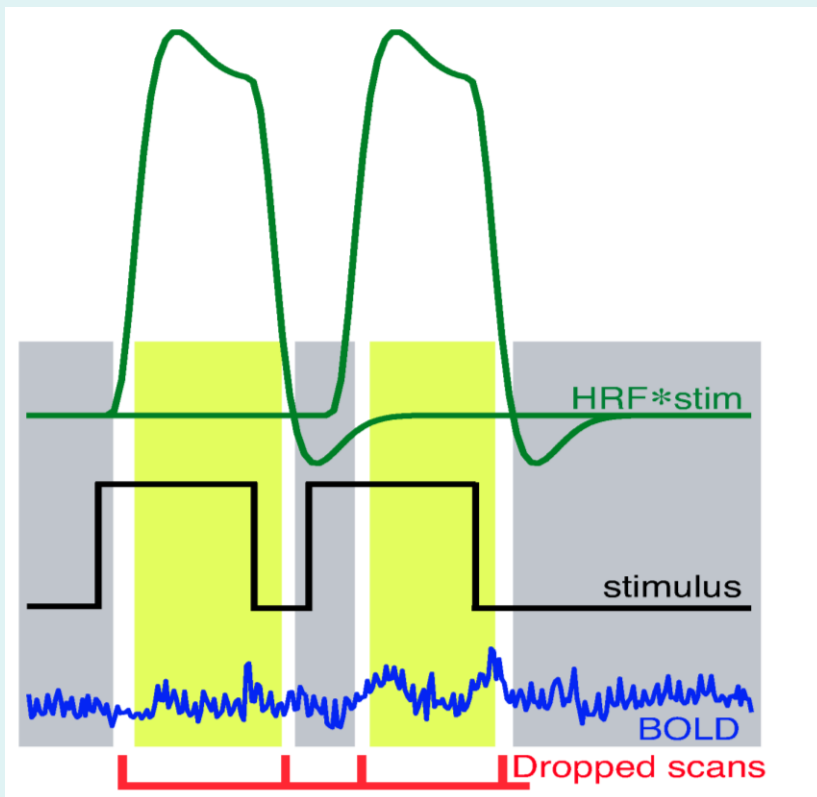
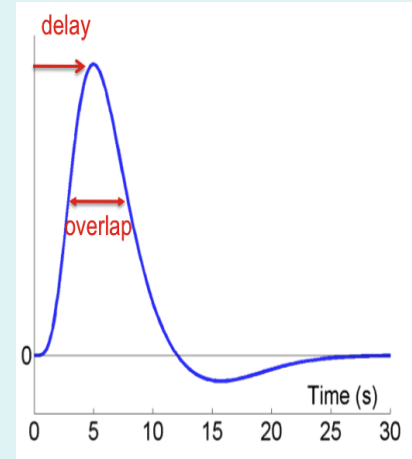
Standard impulse
response function



Brain decoding: raw BOLD

Design:

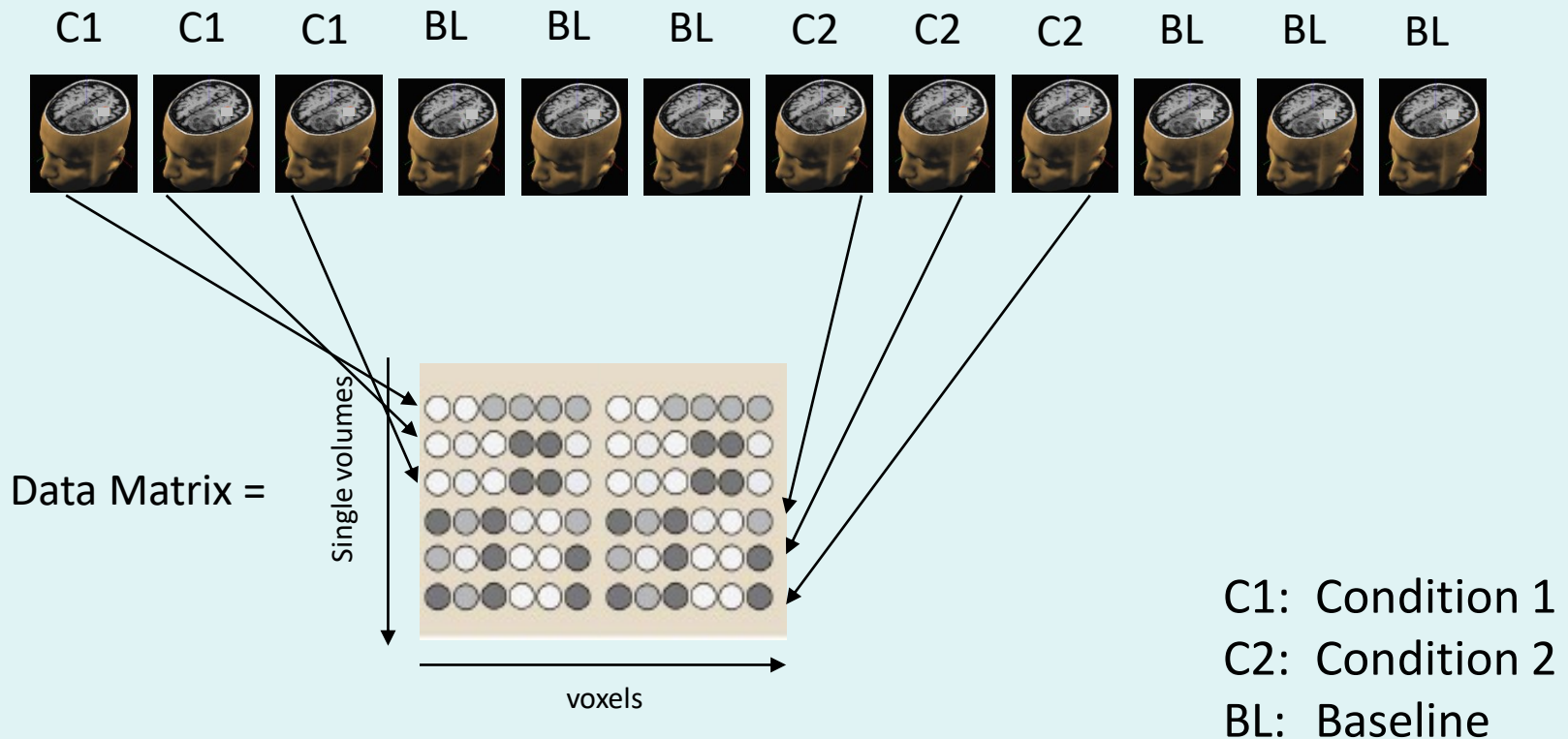
- Block or event-related design
- Accounting for haemodynamic function, with HRF 'delay' & 'overlap'



Brain decoding: raw BOLD

Design:

- Block design
- Use single scans

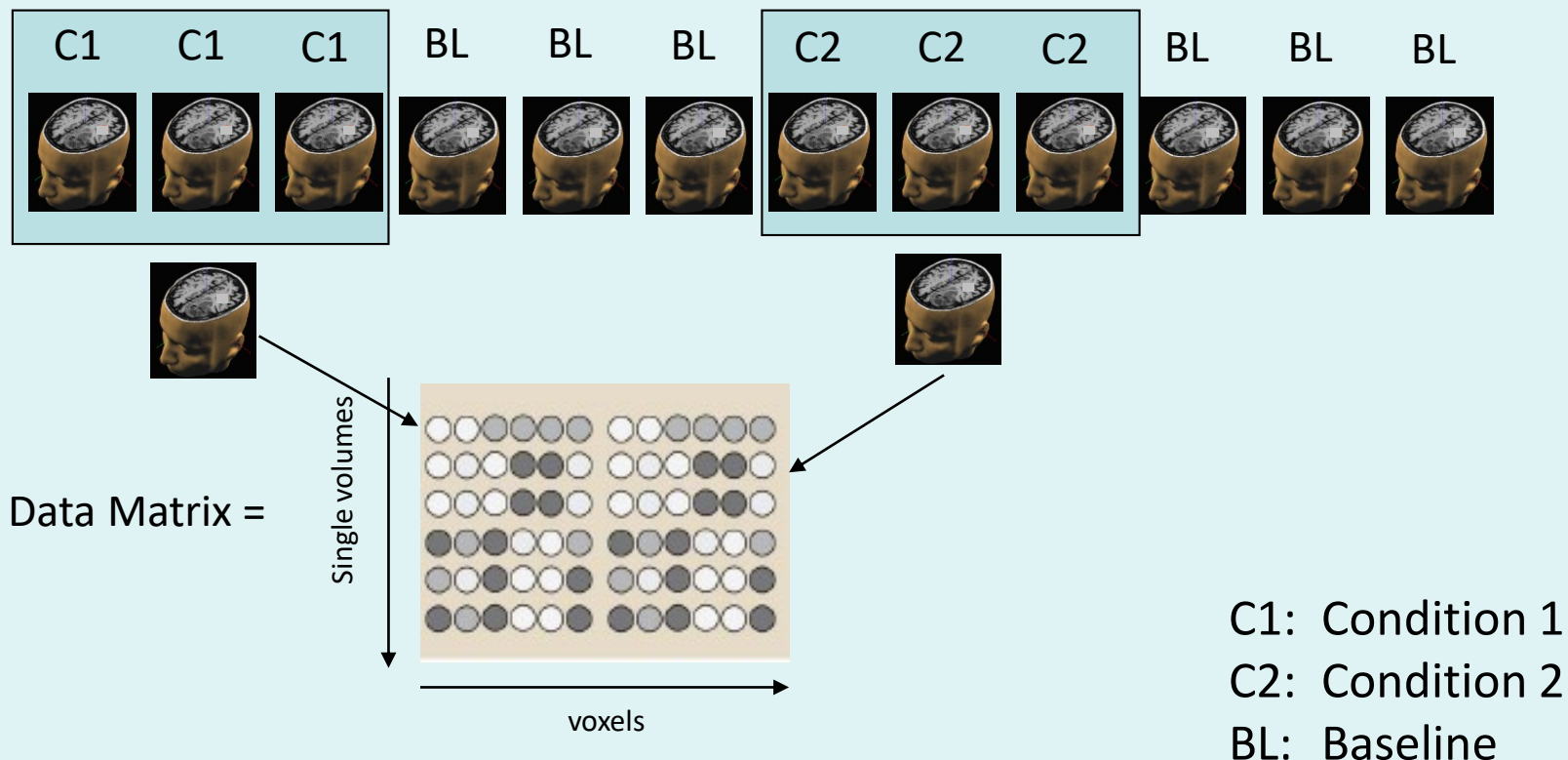


Brain decoding: raw BOLD

Design:

- Block design
- Use single scans

➔ **Average scans**
over blocks/events

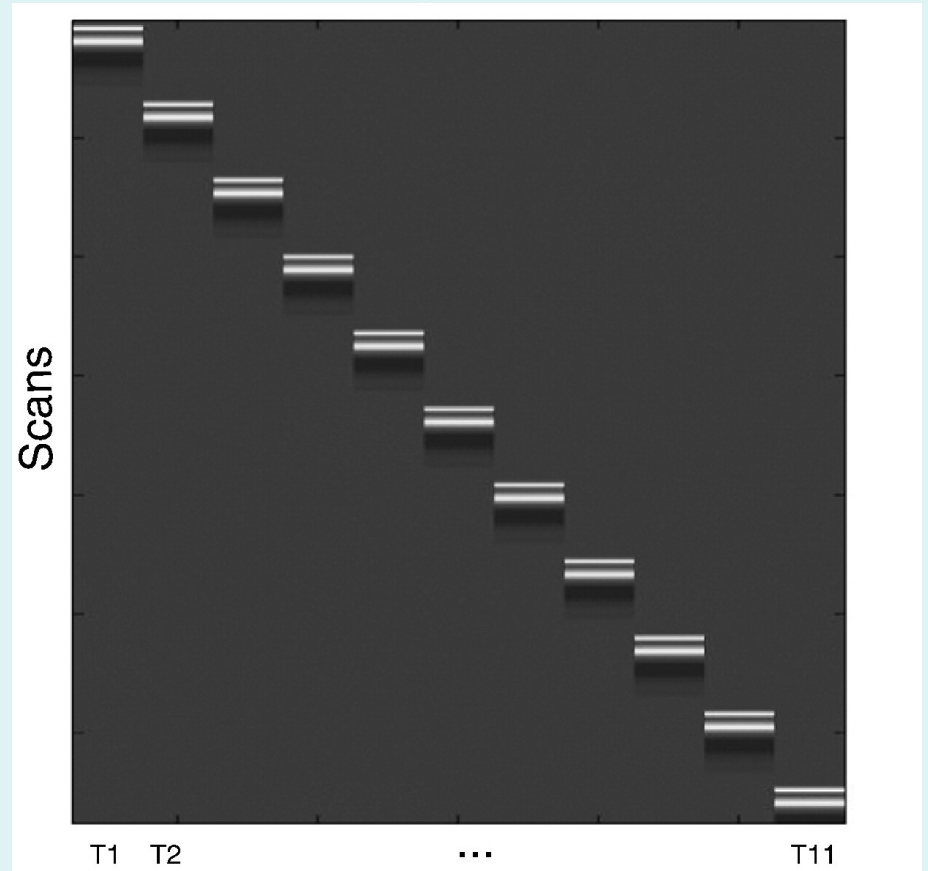
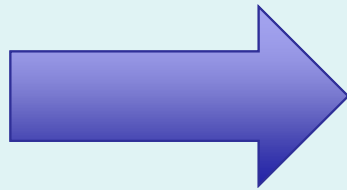


Brain decoding: beta image

1 GLM and 1 beta per event/block



LSA

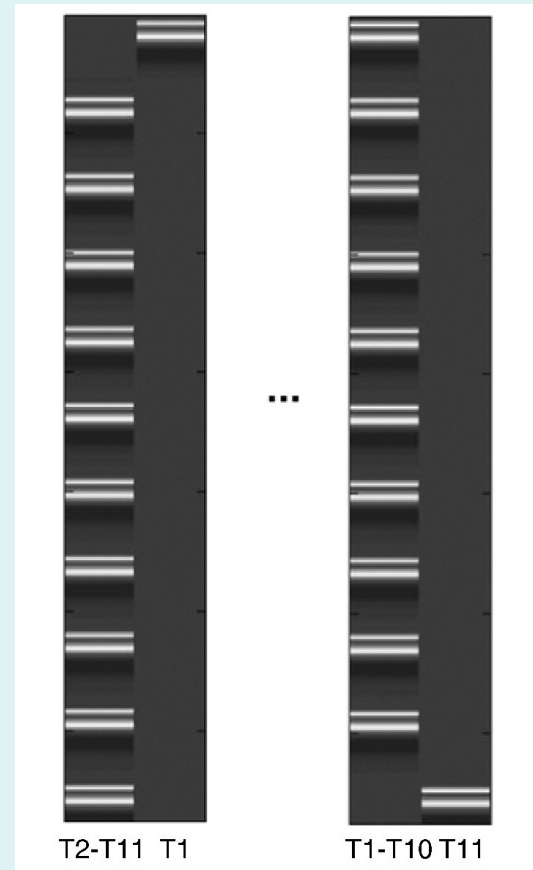
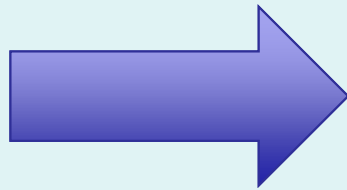


LSU

"Least Squares All" (LSA) and
"Least Squares Unitary" (LSU)

Brain decoding: beta image

N GLM's and 1 beta per event/block



LSA

LSS

"Least Squares All" (LSA) and
"Least Squares Separate" (LSS)

Overview

- Introduction
- “Brain decoding” problem
- “Subject prediction” problem
 - Definition & summary approach
 - Event/block design
 - Resting state design
- Conclusion

Subject prediction

Data:

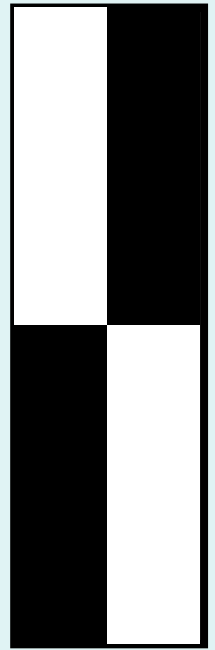
N subjects 1 (or a few) images
per subject(s)

Goal:

Find target value (class or score) of a subject based
on pattern from many subjects.

→ decode “summary” image(s) per subject

Similar to RFX analysis



Subject prediction: summary image

fMRI time series

→ summary image

Event/block design

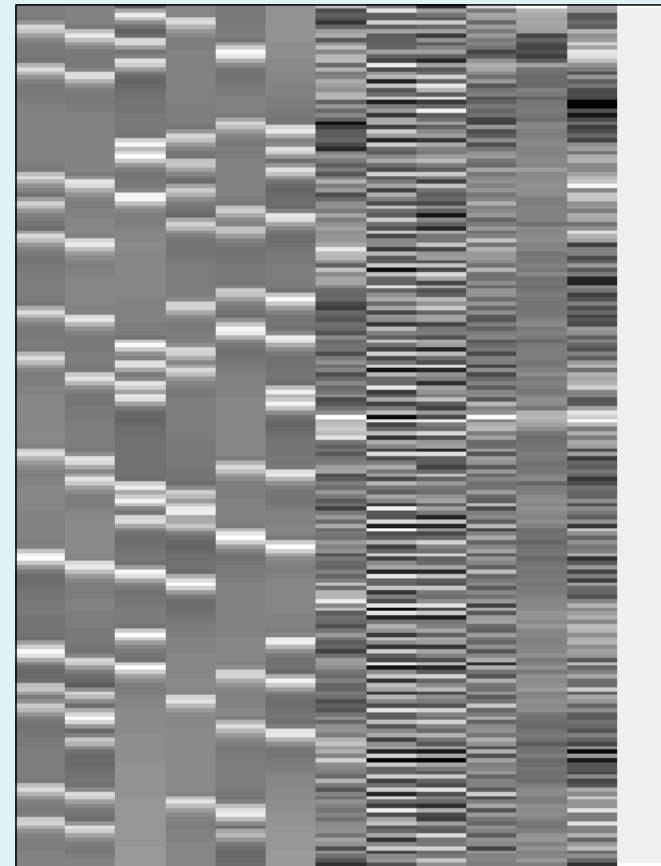
→ build contrast image(s)

e.g.

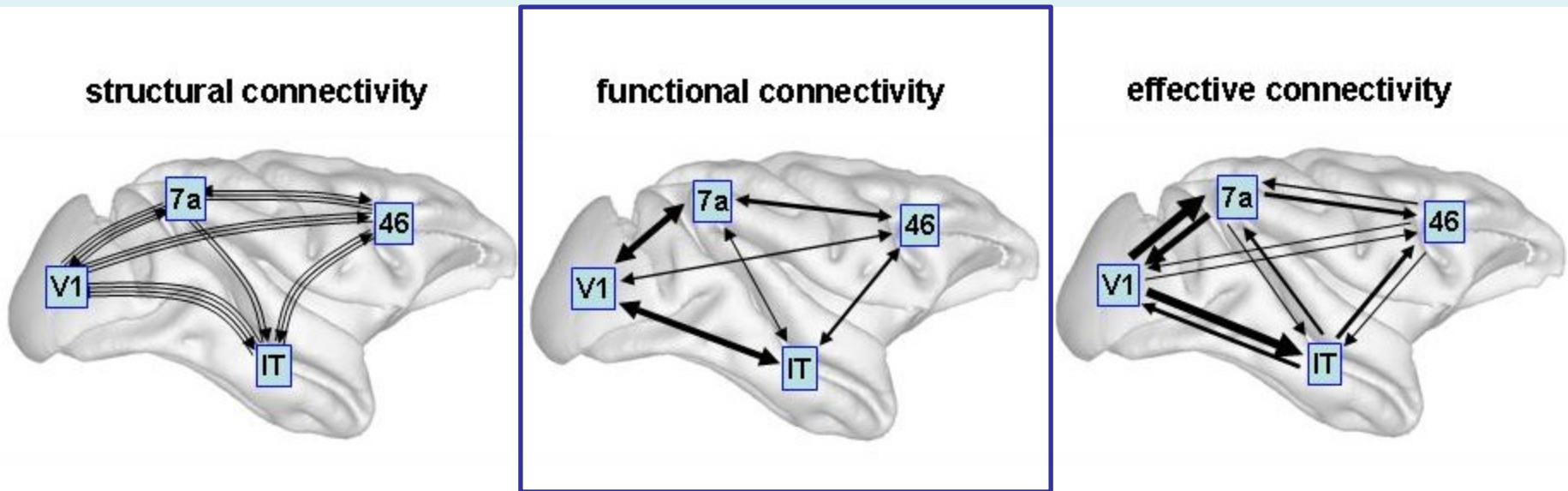
- main/average effect,
 $[1 \ 0 \ 0 \dots]$ or $[0 \ 1 \ 1 \ 0 \dots] / 2$
- difference between 2 conditions
 $[1 \ 0 \ -1 \ 0 \dots]$
- ...

$[1 \ 0 \ -1 \ 0 \dots]$

$[0 \ 0 \ 0 \dots] / 2$



Brain connectivity



Functional connectivity = statistical concept

Statistical dependence estimated by measuring *correlation* or *covariance*

Resting state fMRI

Resting state functional MRI [...] is a [...] method for evaluating regional interactions that occur when a subject is not performing an explicit task.

Paradigm shift:

- Activation → functional segregation
- Spontaneous → functional integration

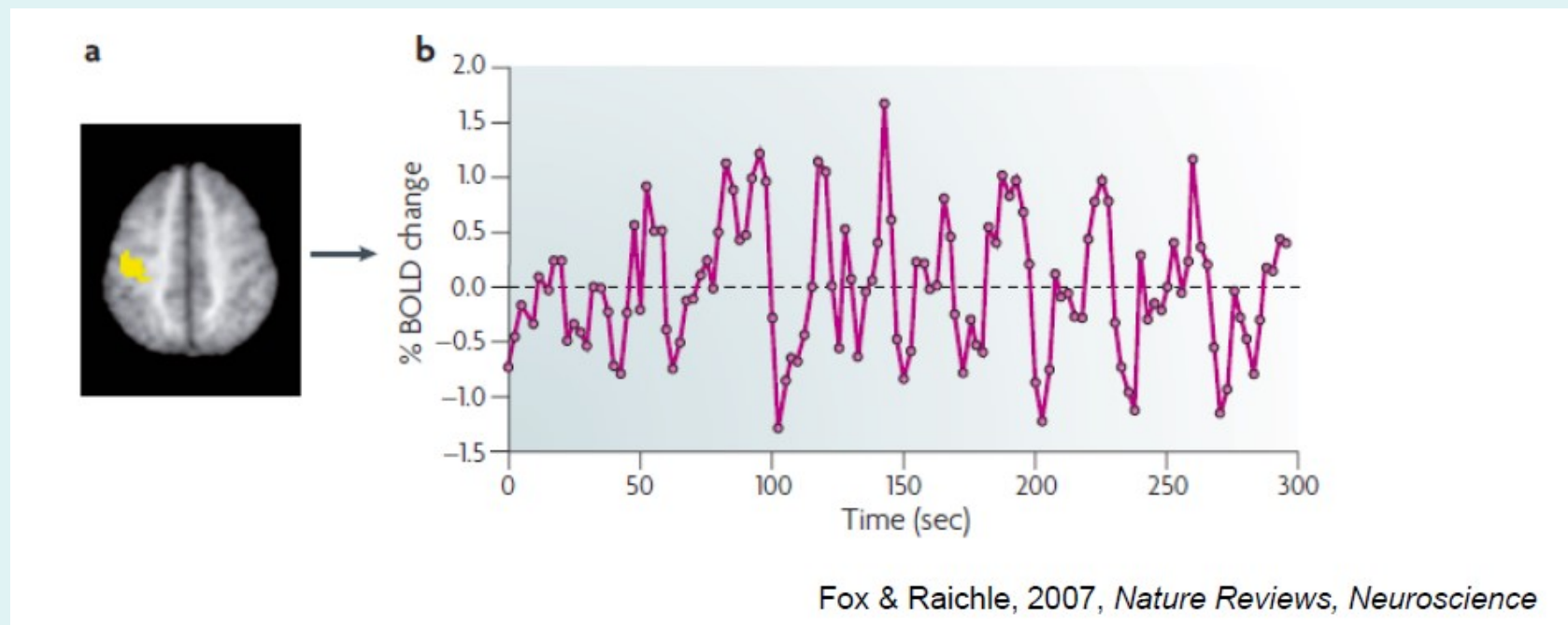
Goal:

Summarize rs-fMRI time series into 1 (few) map(s).

Rs-fMRI: model based approach

Pick one (few) region(s) of interest:

- Extract BOLD signal time-series

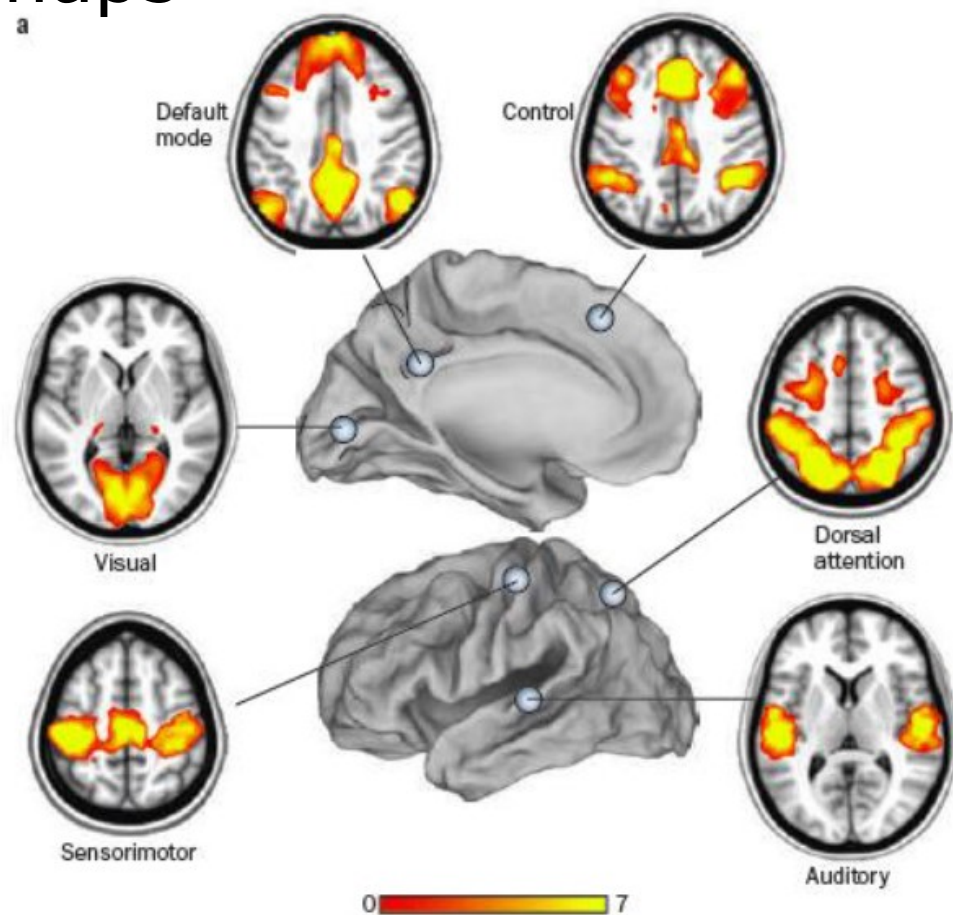


- Enter time series as regressor in a GLM & find correlation map

Rs-fMRI: model based approach

Multiple/different region of interest

→ multiple/different maps



Rs-fMRI: model free approach

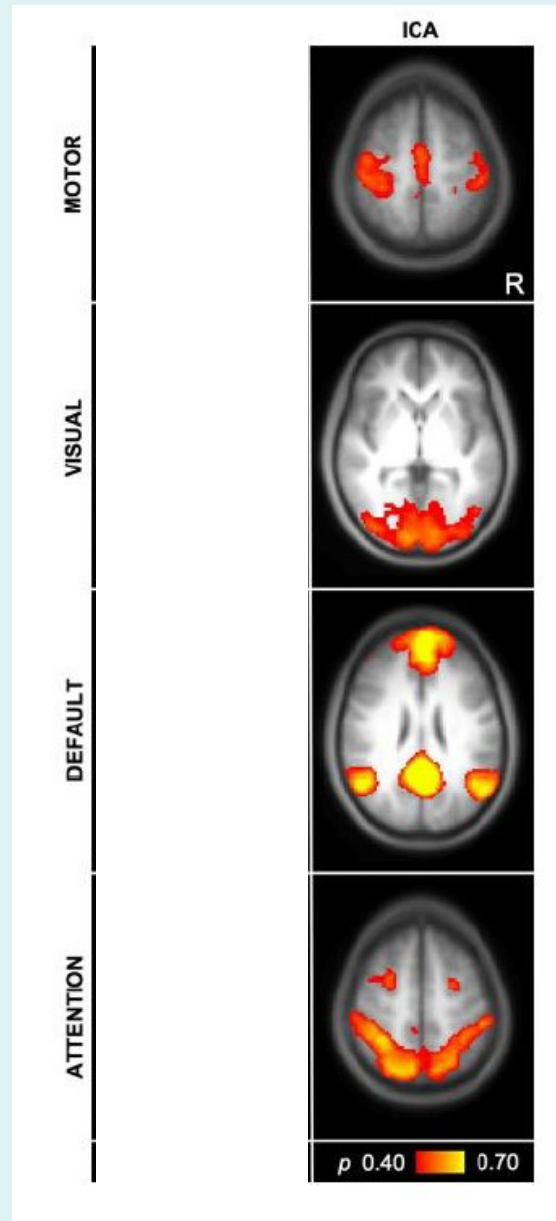
Decompose original fMRI time series into linear combination of

- basis vectors, PCA
- independent components, ICA

i.e. data driven approach.

➔ A few basis/component maps per subject

Rs-fMRI: model free approach



Overview

- Introduction
- “Brain decoding” problem
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- Conclusion

Conclusion

1 sample = 1 image

- What is your question of interest?
- At what level of inference ?
- What is the experimental design?
- How much data is/will be available?

Thank you for your attention!

Any question?

“Univariate vs. multivariate” concepts

Univariate

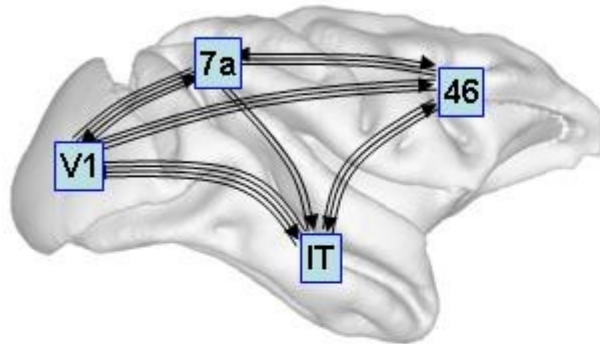
- 1 voxel
- target \rightarrow data
- look for difference or correlation
- General Linear Model
- GLM inversion
- calculate contrast of interest

Multivariate

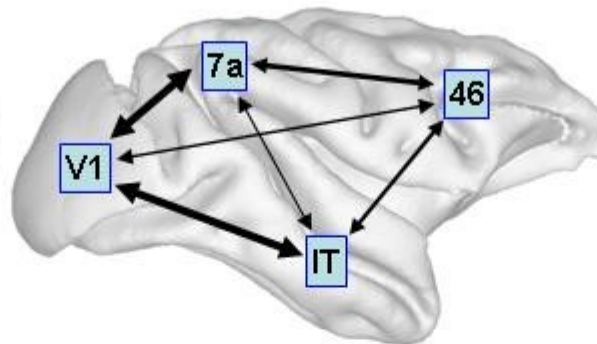
- 1 volume
- data \rightarrow target
- look for similarity or score
- Specific machine (SVM, GP,...)
- training & testing cross-validation
- estimate accuracy of prediction

Brain connectivity concepts

structural connectivity



functional connectivity



effective connectivity

