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Turku PET Centre

First level fMRI data analysis

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Outline

- We learned the general linear model (GLM), preprocessing, and experimental design

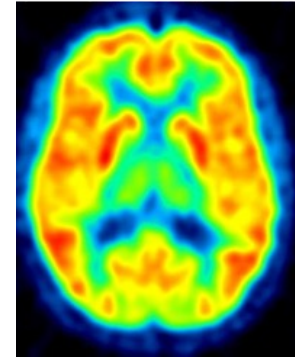
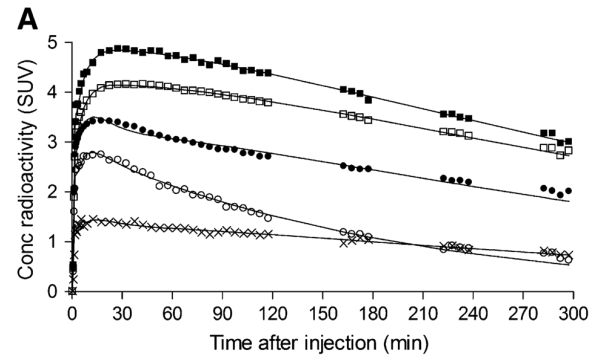
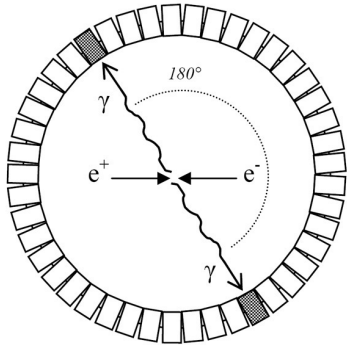
- What is next:

Apply the GLMs to the preprocessed data

Note: 1st level analysis = within subject analysis

Within-subject modelling

Positron emission tomography



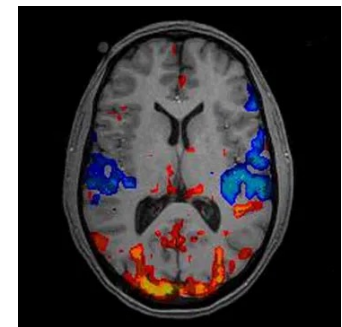
Density

fMRI

Time



Task

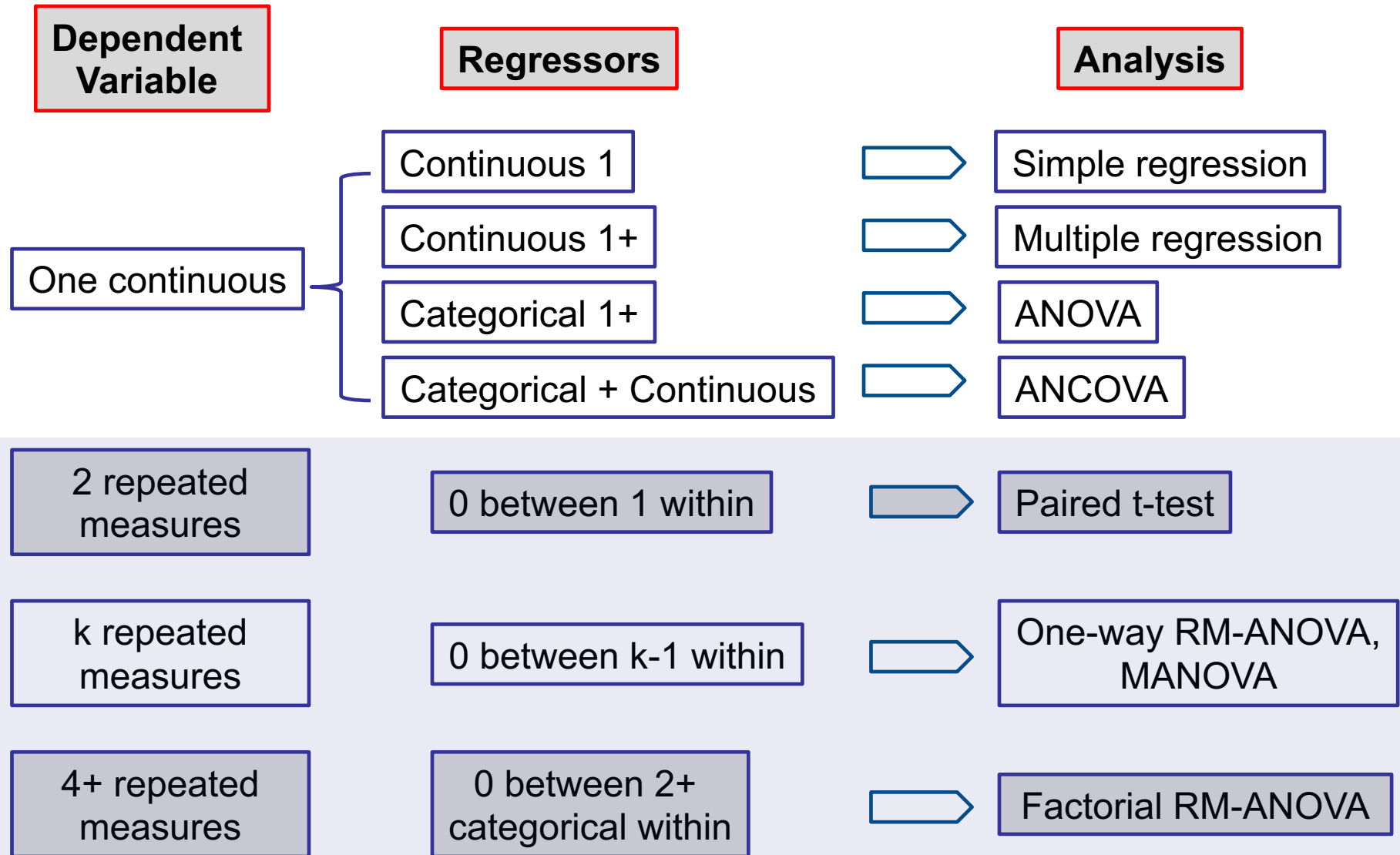


Statistical map

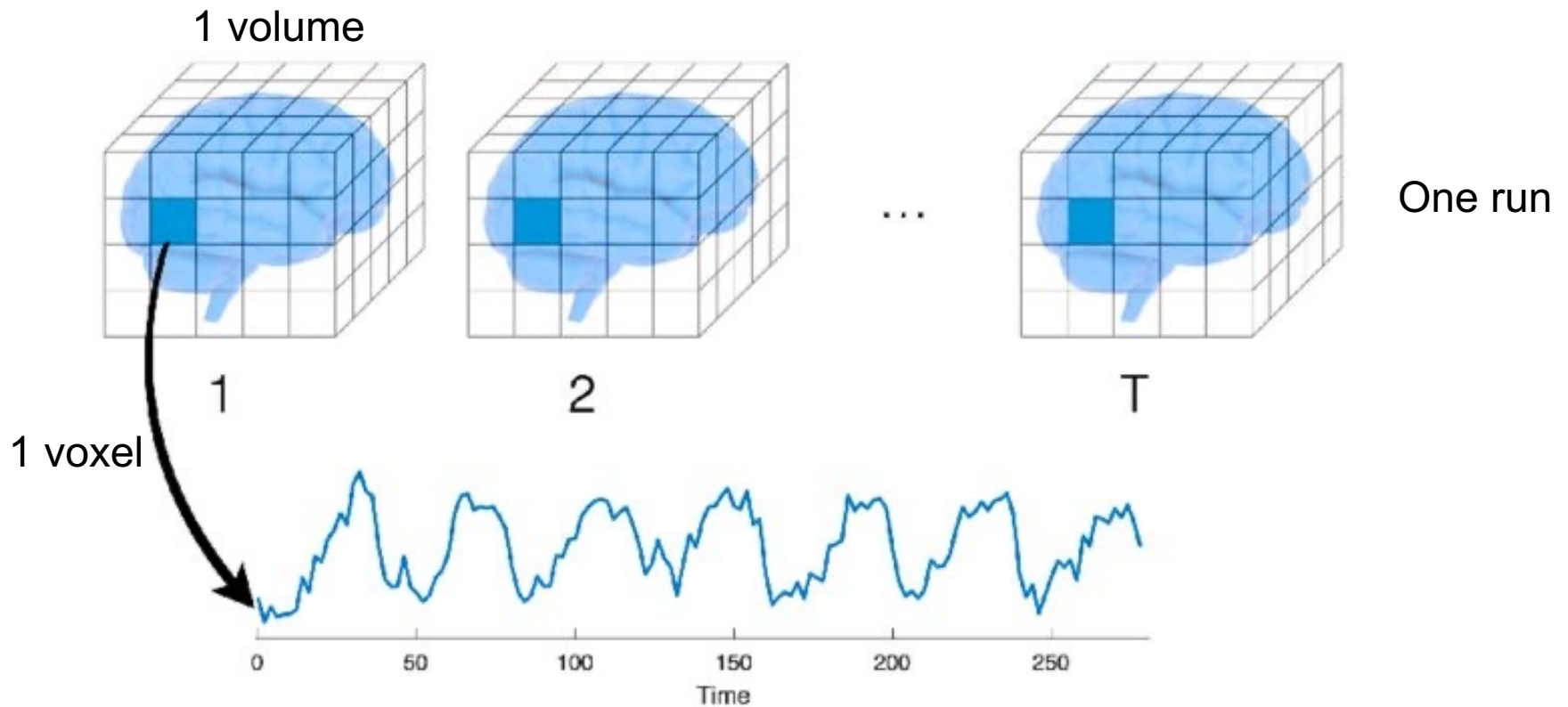
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- Statistical models according to experiment design
- Contrasts
- Operation in SPM12
 - To set the parameters
 - To make contrasts: T contrasts

The general linear model (GLM) family

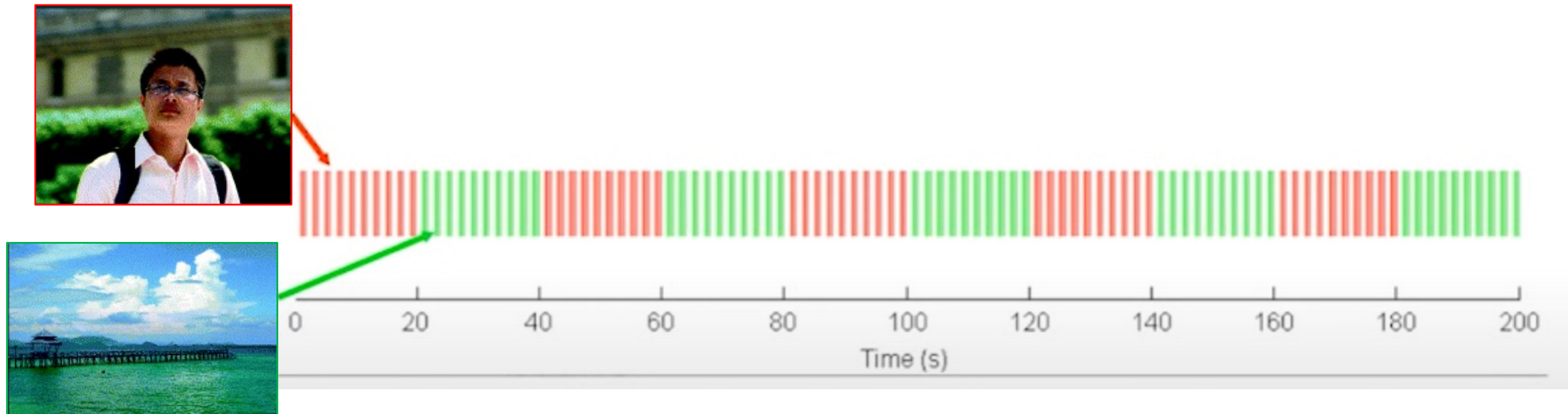


Mass univariate approach

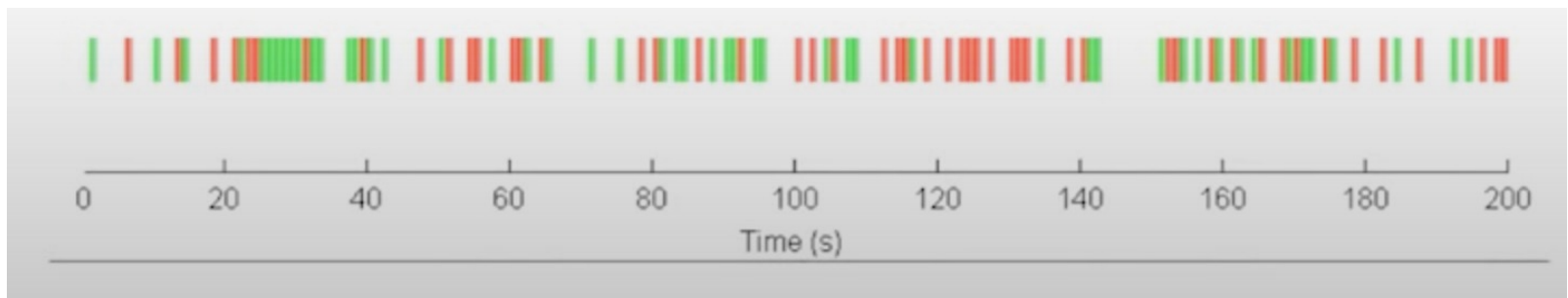


1st level GLM: single-voxel, signal subject

1. Block design



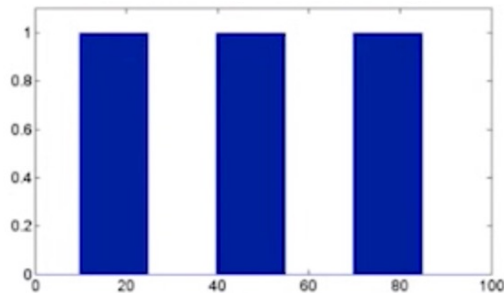
2. Event-related design



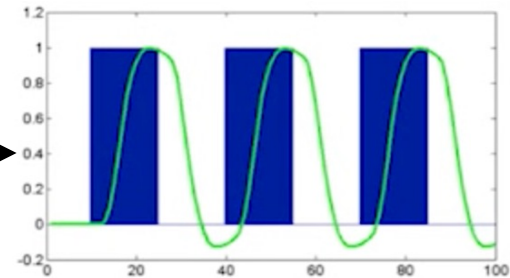
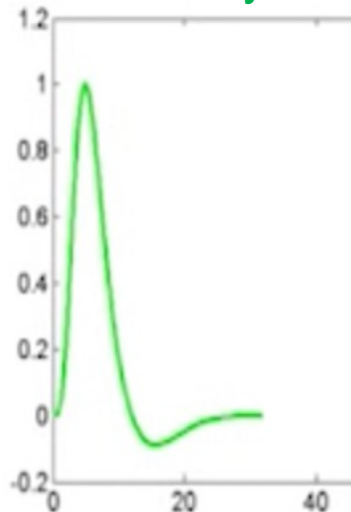
How to build a regressor?

- BOLD responses are delayed: **peak at 4-6 s** and **baseline 20-30 s**.
- Convolved with the hemodynamic response function (**HRF**)
- The linear time-invariant (**LTI**) system

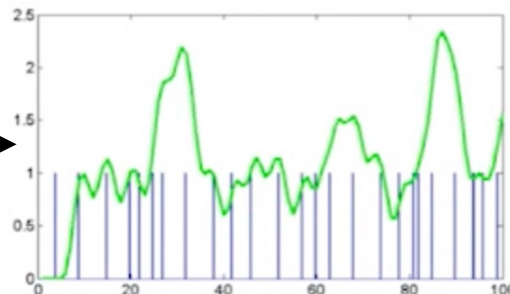
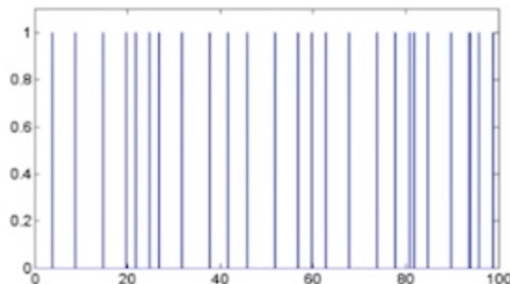
Block design



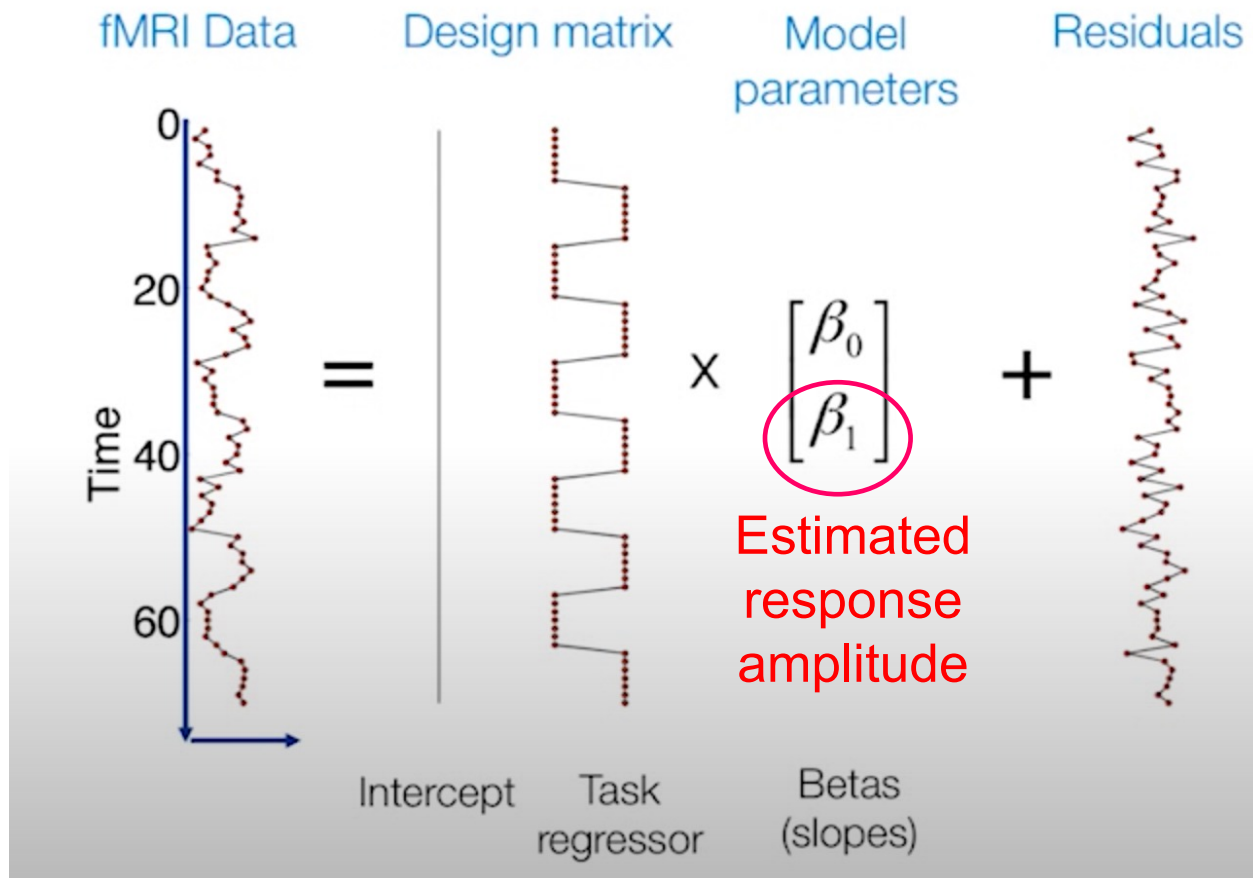
Hemodynamic
delay



Event-related design

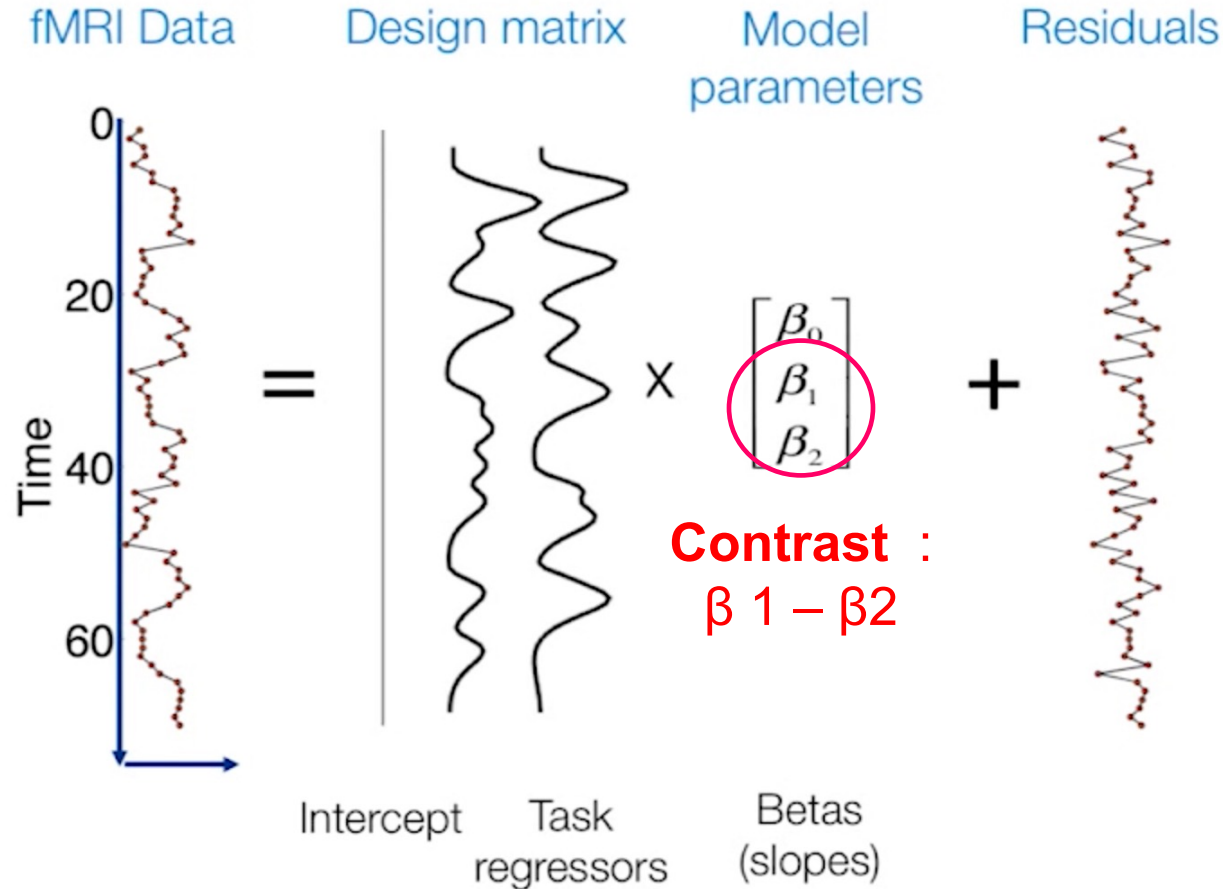


Block design: one regressor of interest (e.g. people vs. landscape)





Event-related design: two continuous regressors (people vs. landscape)



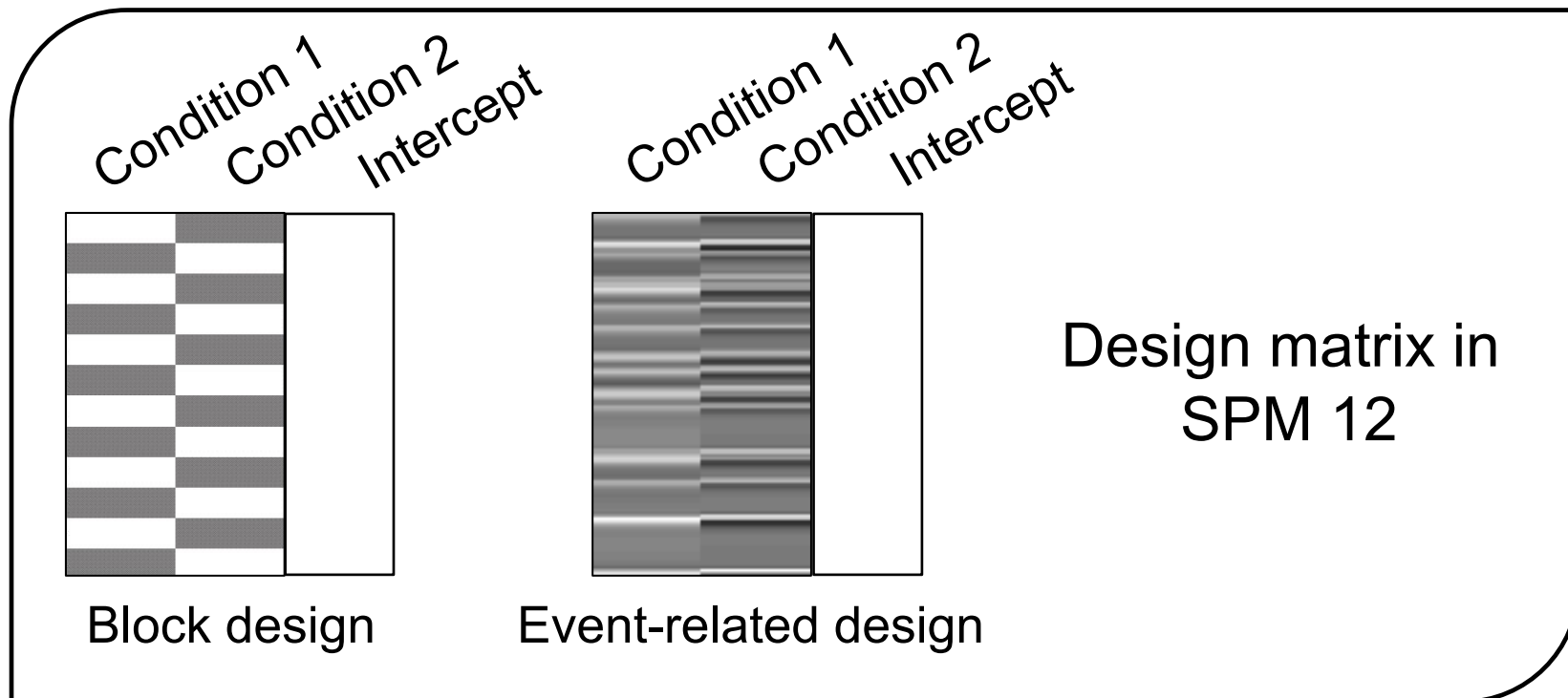
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Contrasts

Experimental conditions: people vs. landscape

- Your interest is often the difference between the two conditions, which is “contrast”
 - You can calculate the difference, sum or separately each conditions, which are calculated by different linear contrasts.
-
- We only focus on T contrast in this lecture ! !



Difference between conditions

$[1 \ -1]$ = “Con1 > Con2”

$[-1 \ 1]$ = “Con1 < Con2”

Separately

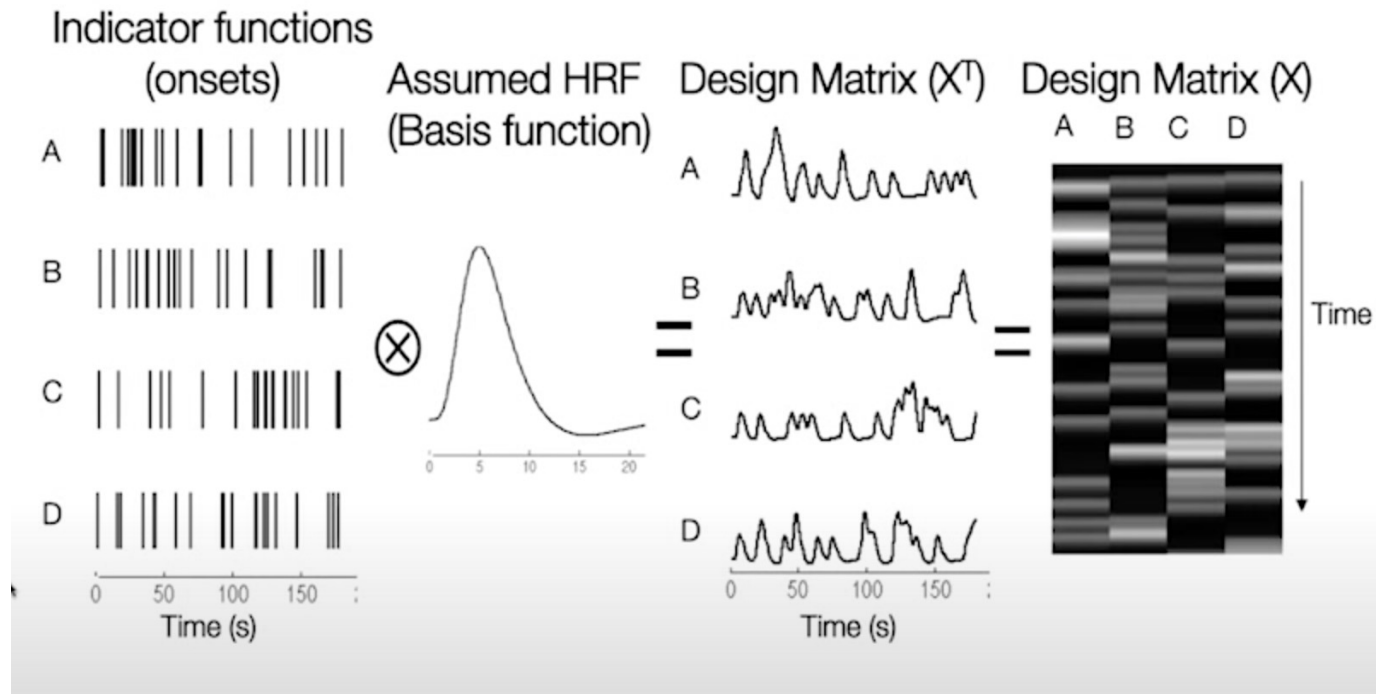
$[1 \ 0]$ or $[-1 \ 0]$ = “main effect Con1”

$[0 \ 1]$ or $[0 \ -1]$ = “main effect Con2”

Sum

$[1 \ 1]$ or $[-1 \ -1]$

More regressors



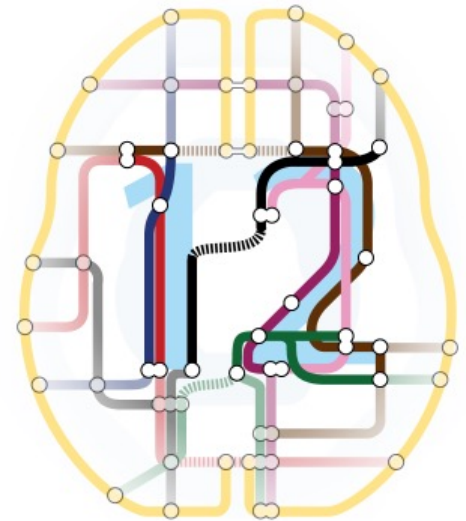
- $[1 \ 1 \ -1 \ -1]$: $(A+B) > (C+D)$
- $[1 \ -1 \ 1 \ -1]$: $(A+C) > (B+D)$
- $[1 \ 0 \ 0 \ 0]$: main effect of A
- $[1 \ 1 \ 0 \ 0]$: Sum of $(A+B)$ vs the mean of the signal
-

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SPM12

- SPM - theoretical concepts of Statistical Parametric Mapping in a complete analysis package.
- Run in matlab
- See more information:
<https://www.fil.ion.ucl.ac.uk/spm/software/spm12/>



Simple or multiple regression, t-tests, ANOVA, ANCOVA

$$\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$$

The diagram illustrates the linear regression equation $\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$ with the following components and labels:

- Observed Data:** The vector $\begin{bmatrix} Y_1 \\ Y_2 \\ \vdots \\ Y_n \end{bmatrix}$ is labeled "Observed Data".
- Design matrix:** The matrix $\begin{bmatrix} 1 & X_{11} & \cdots & X_{1p} \\ 1 & X_{21} & \cdots & X_{2p} \\ \vdots & \vdots & & \vdots \\ 1 & X_{np} & \cdots & X_{np} \end{bmatrix}$ is labeled "Design matrix".
- Model parameters:** The vector $\begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}$ is labeled "Model parameters".
- Residuals:** The vector $\begin{bmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \vdots \\ \varepsilon_n \end{bmatrix}$ is labeled "Residuals".

The equation is shown as $\begin{bmatrix} Y_1 \\ Y_2 \\ \vdots \\ Y_n \end{bmatrix} = \begin{bmatrix} 1 & X_{11} & \cdots & X_{1p} \\ 1 & X_{21} & \cdots & X_{2p} \\ \vdots & \vdots & & \vdots \\ 1 & X_{np} & \cdots & X_{np} \end{bmatrix} \times \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix} + \begin{bmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \vdots \\ \varepsilon_n \end{bmatrix}$.

The laughter experiment

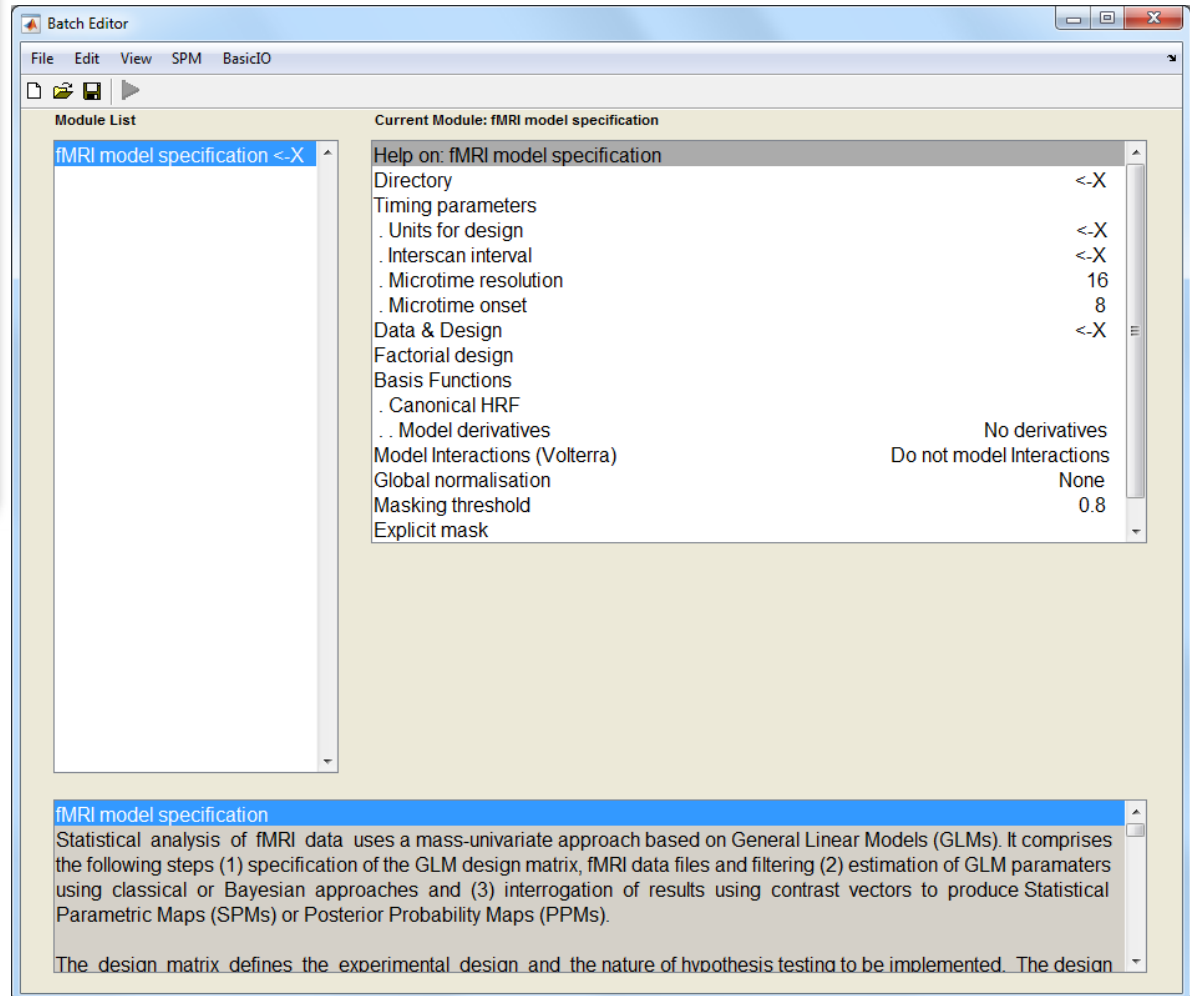
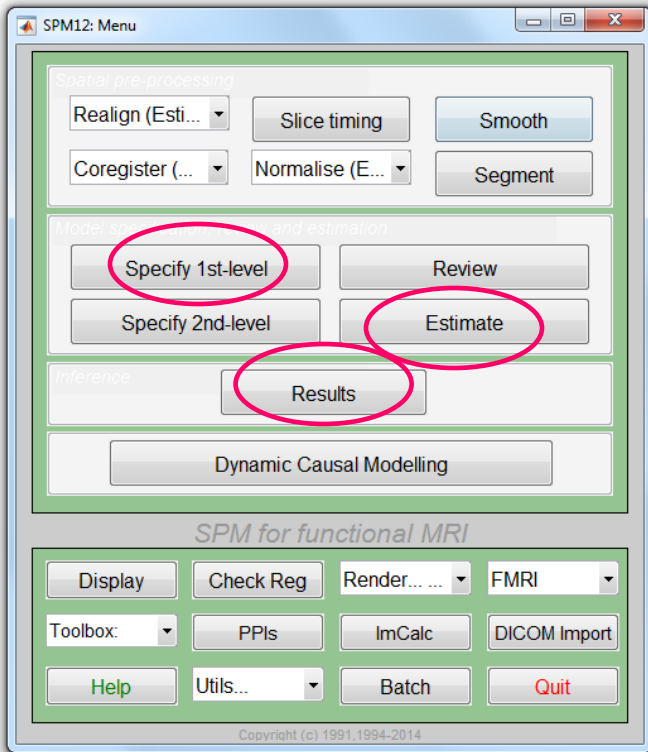
Four stimuli types:

Laughter / Crying vocalization /
Scrambled laughter / Scrambled Crying

Block design (event-related design?)

Statistical model (ANOVA, or multiple regression)

Operation in SPM12



SPM12: parameter setting

Current Module: fmri model specification

Help on: fmri model specification

Directory

Timing parameters

- . Units for design
- . Interscan interval
- . Microtime resolution
- . Microtime onset

Data & Design

Factorial design

Basis Functions

- . Canonical HRF
- . . Model derivatives

Model Interactions (Volterra)

Global normalisation

Masking threshold

Explicit mask

Result to store

seconds
2.6

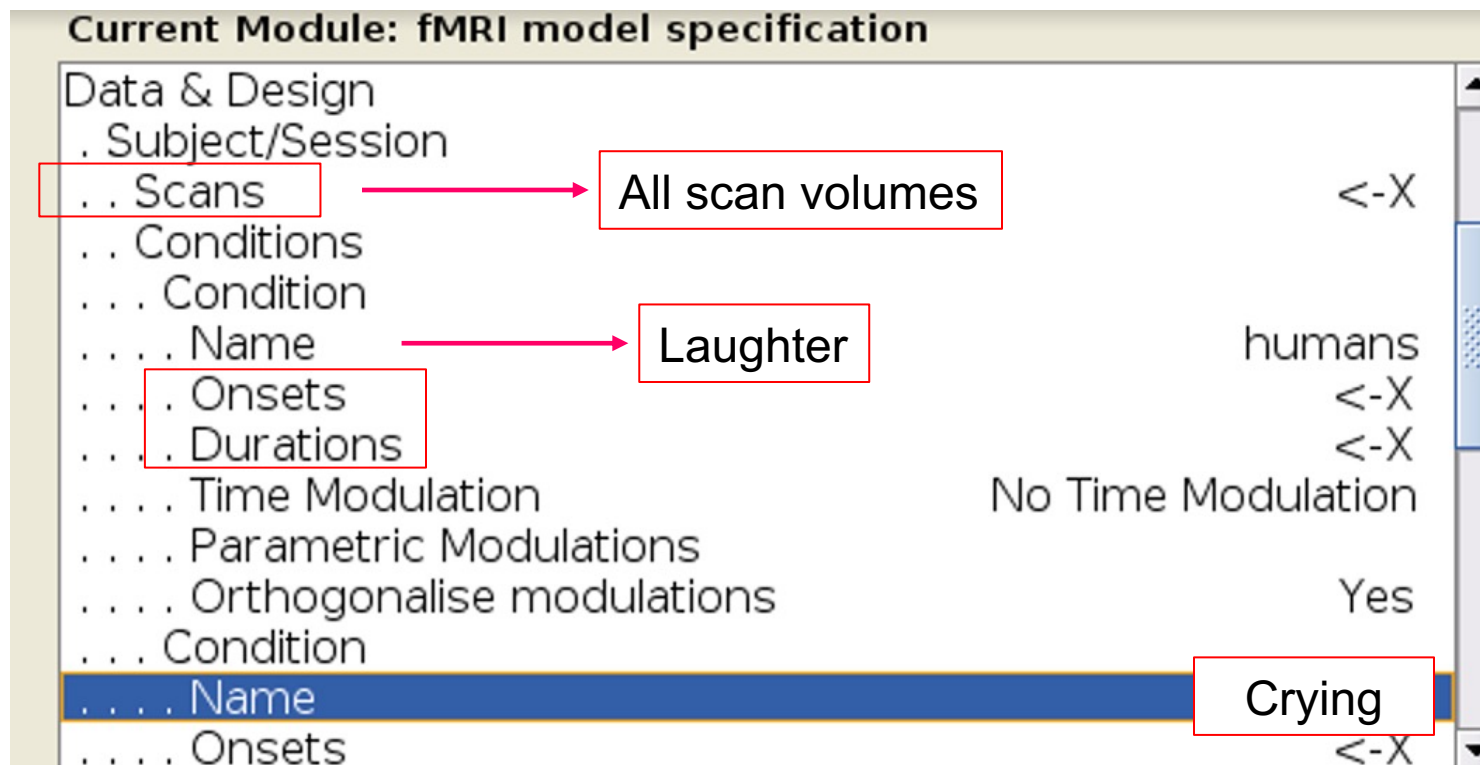
Dependent variables

Statistical model

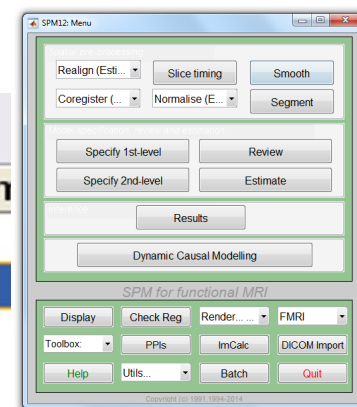
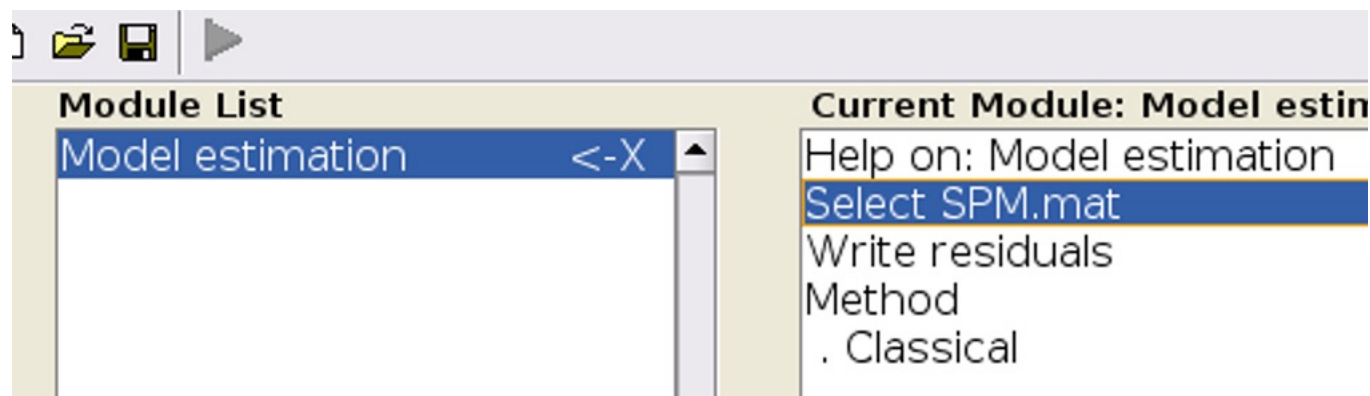
Regressor building

Leave as default,
or specify if necessary

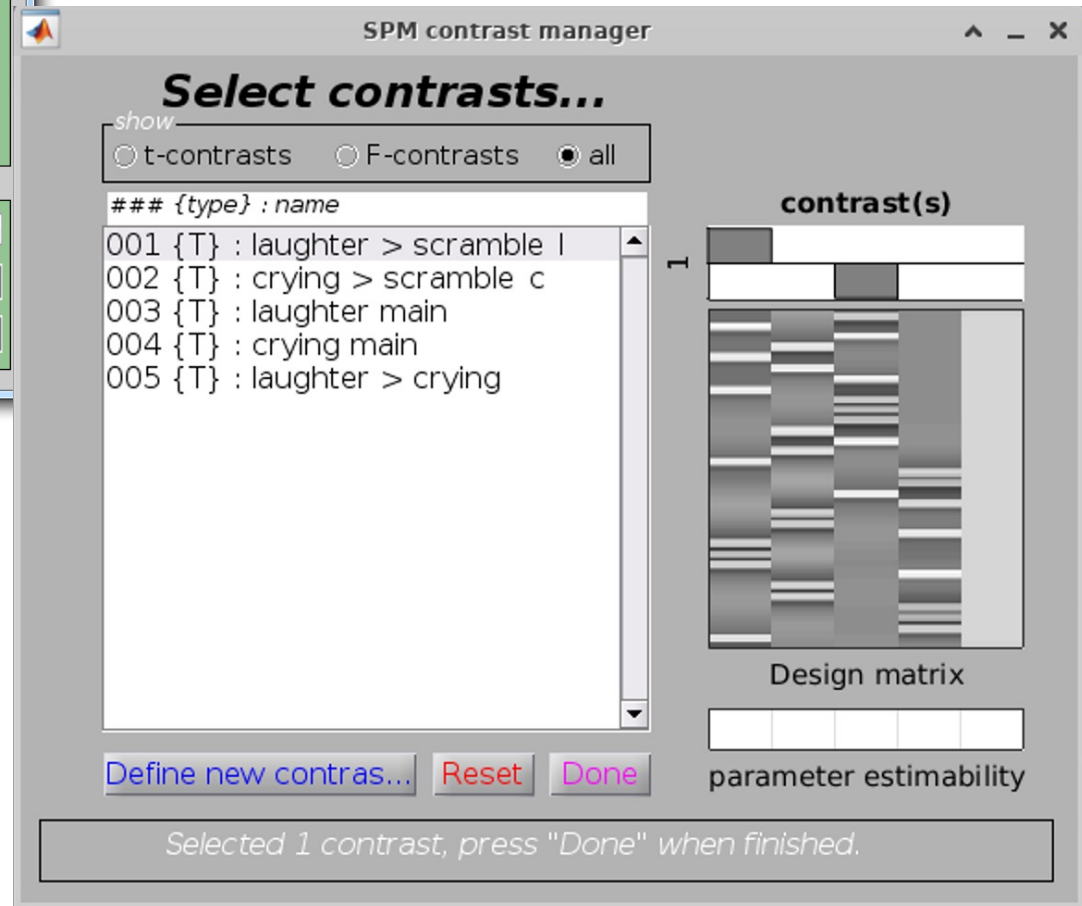
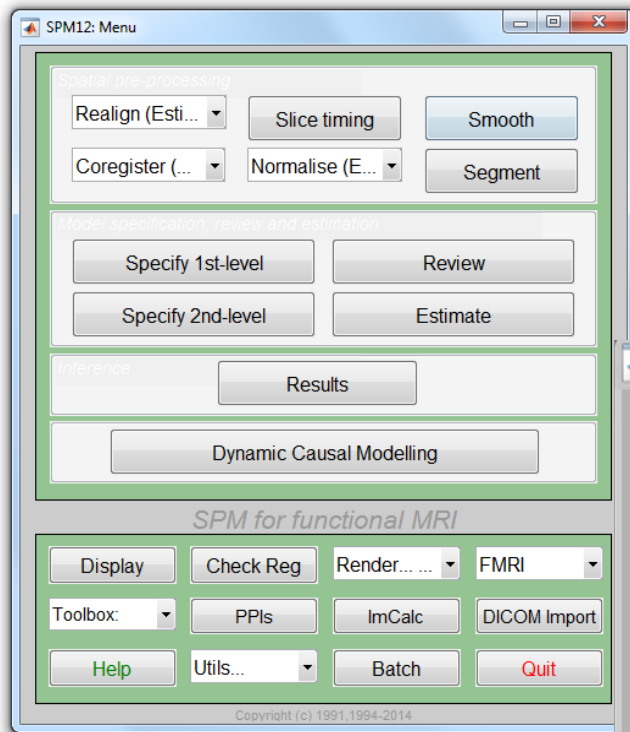
No masks



Data estimation

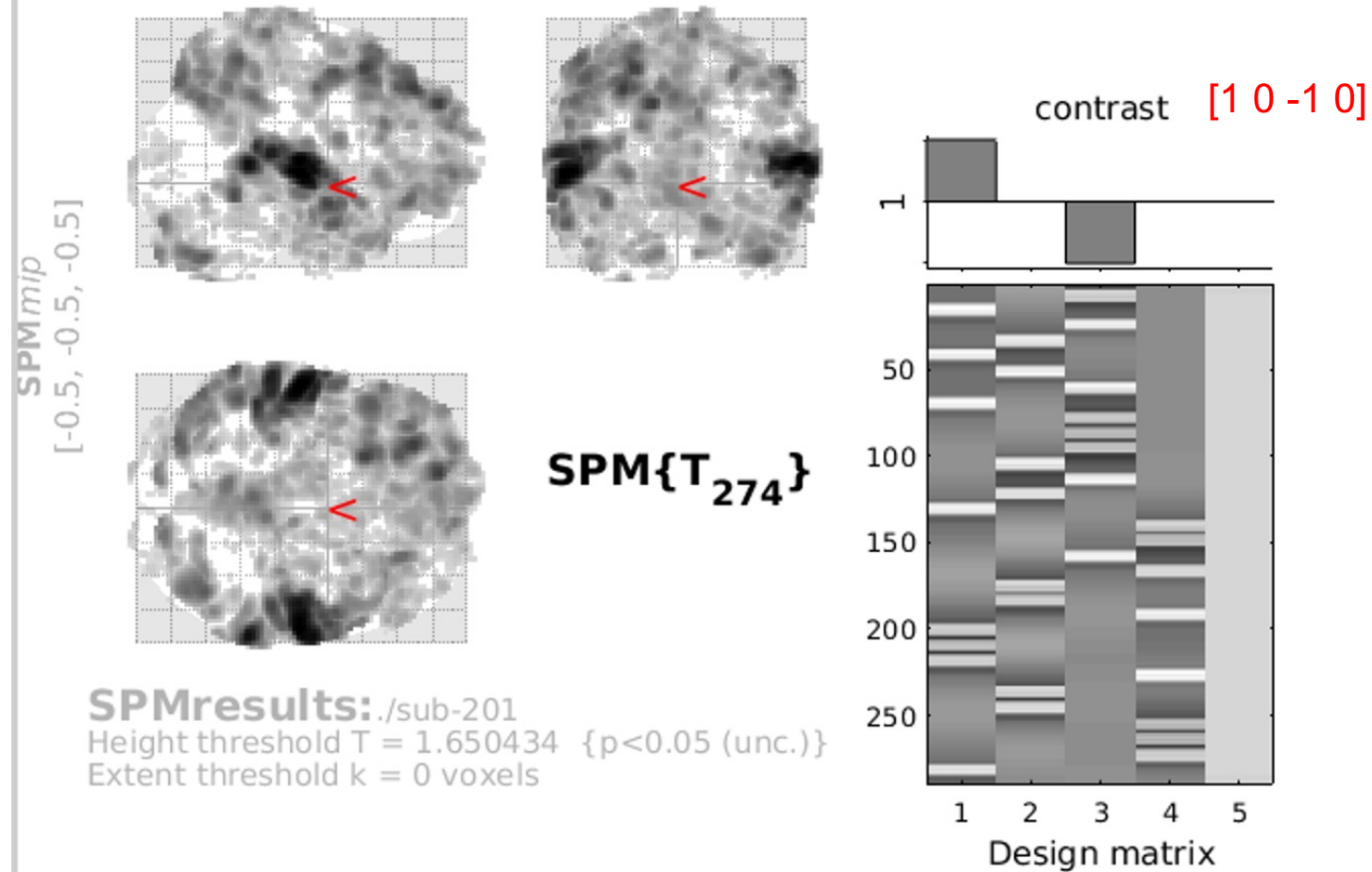


SPM12: setting contrasts

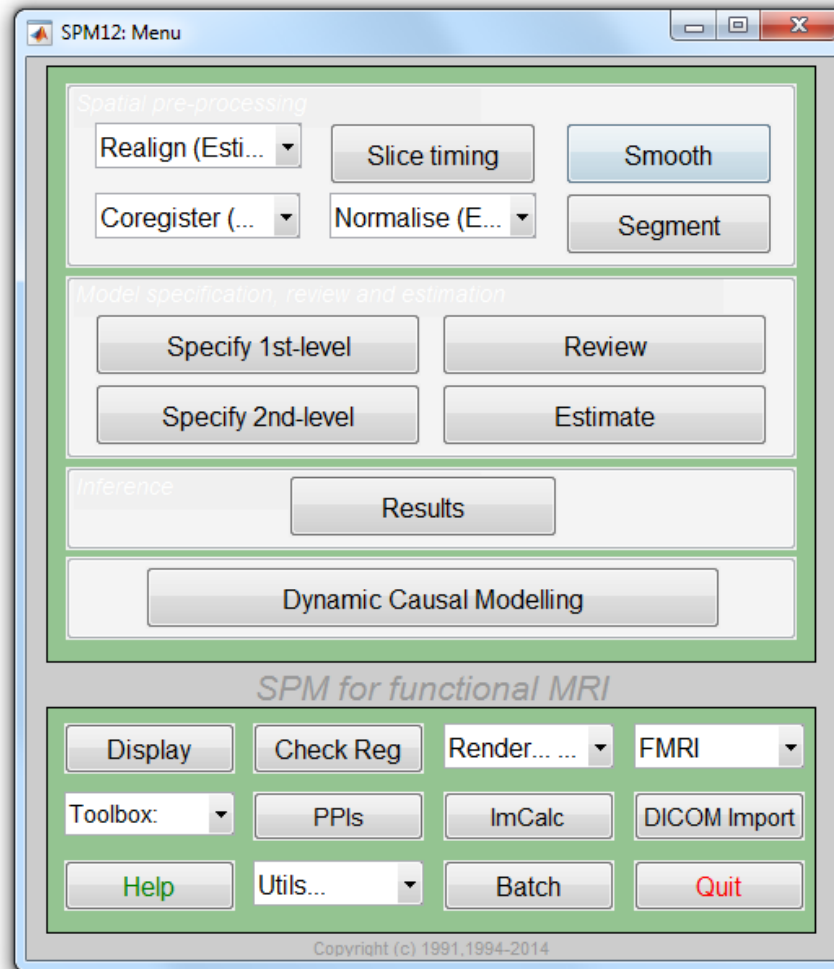


SPM12: result

laughter > scramble_I



Ready for 2nd-level analysis



Resources

- Principles of fMRI: <https://youtu.be/OyLKMb9FNhg>
- Spm12 manual:
https://www.fil.ion.ucl.ac.uk/spm/doc/spm12_manual.pdf
- Behav Res (2014) 46:596–610