

First level fMRI data analysis

Turku PET Centre Brain Imaging Course 2024

Lihua Sun, Turku PET Centre



Outline

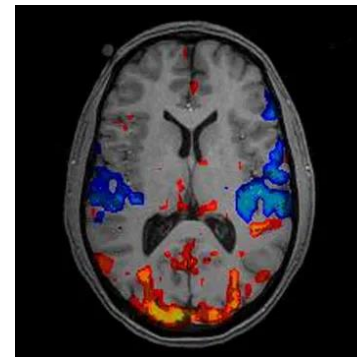
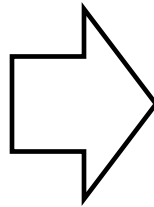
- What is 1st level analysis?
- Statistical models
- Contrasts
- Examples for illustration
 - Operation in SPM12
 - Reading the results, e.g., the contrasts

What is 1st level analysis?

- fMRI data analysis has two levels: 1st (within-subject) and 2nd (group-level).



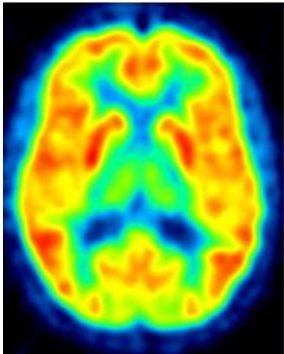
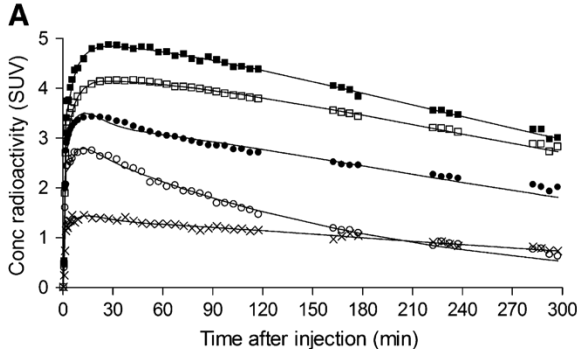
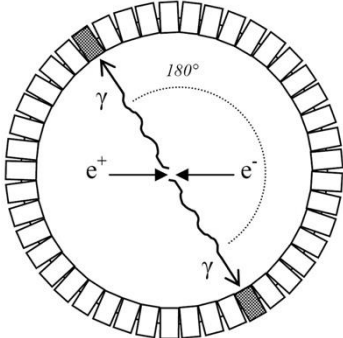
One subject's raw T2* data



The subject's statistical map

Within-subject: comparison between PET and fMRI data

1. Traditional PET



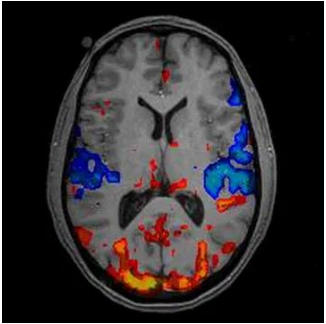
Density
e.g., glucose uptake

2. fMRI

Time



Task

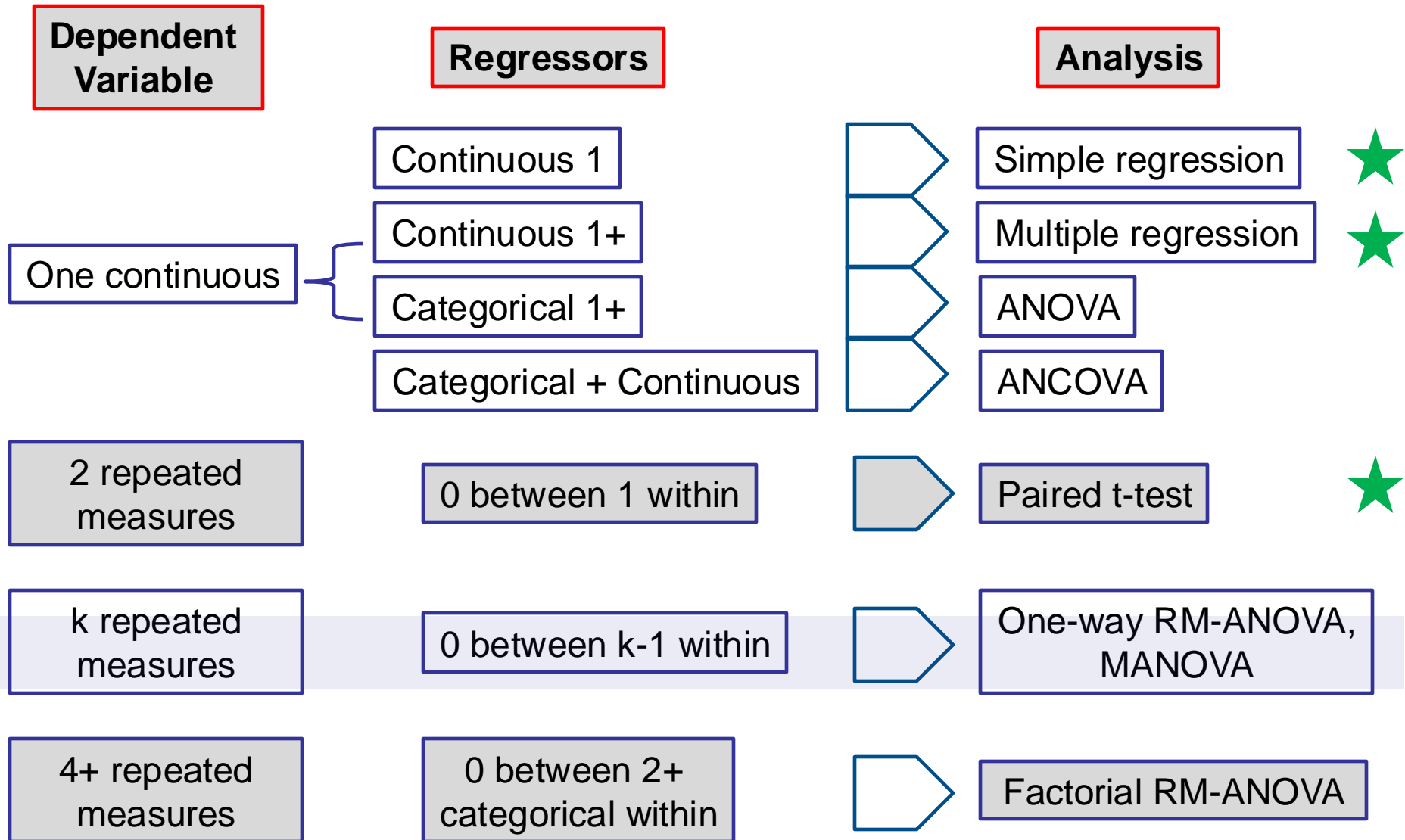


Statistical map

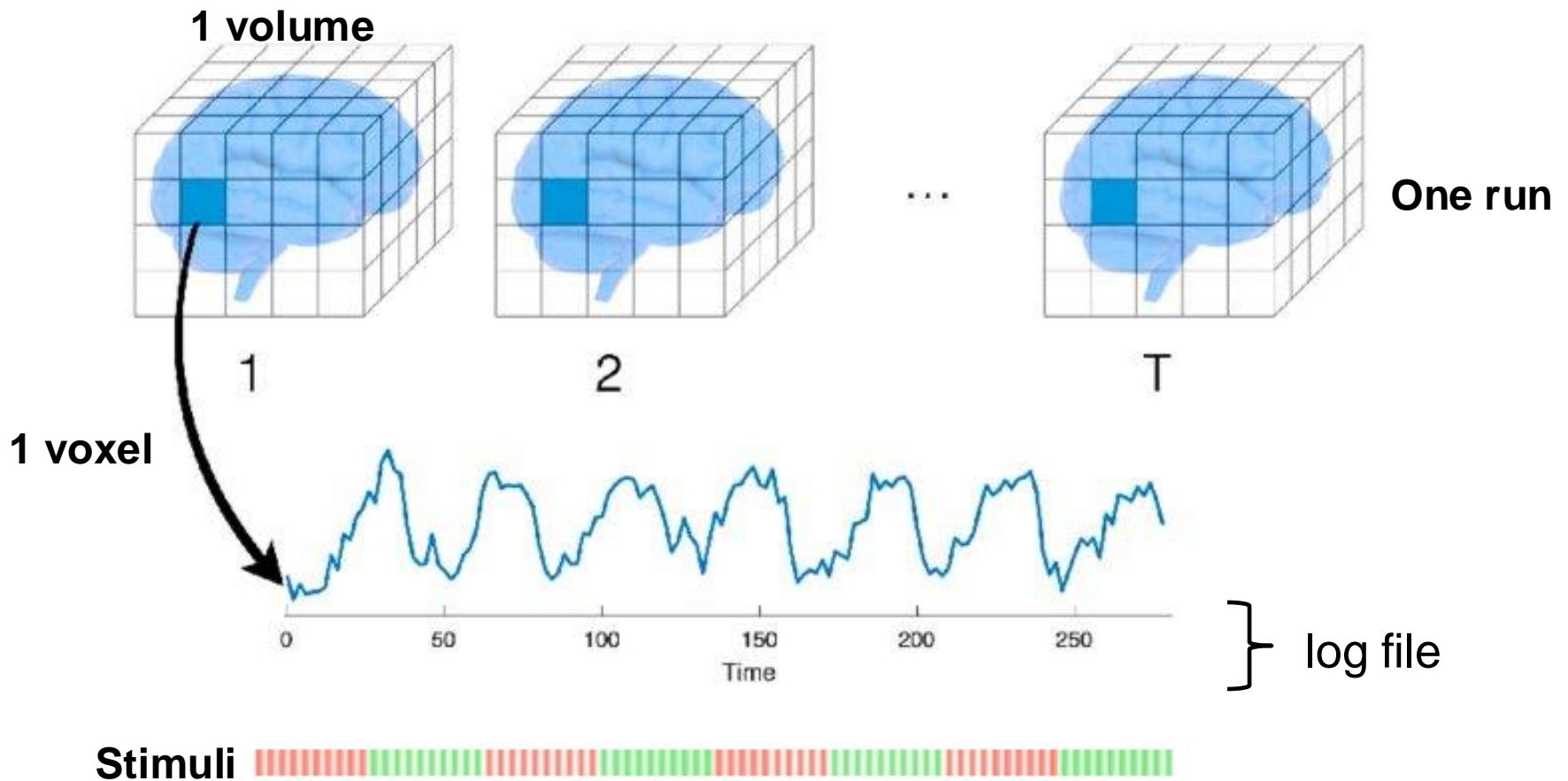
Why is 1st level analysis?

- BOLD signals are dominated by noise:
physiological factors (breathing, heart rate), head movement, scanner instabilities, magnetic susceptibility artifacts, and neuronal variability, leading to low signal-to-noise ratio
>> preprocessing NOT enough!
- Multiple repeated exposure to stimuli to increase signal-to-noise ratio.
- Statistical analysis to extract the signal-associated brain responses **>> statistical map**

The general linear model (GLM) family

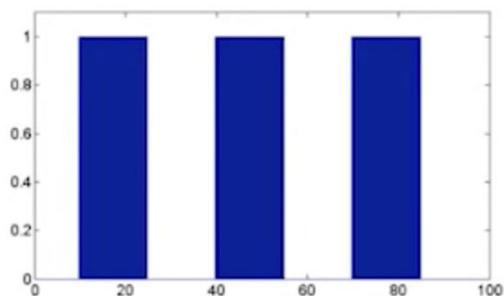


Voxel-level data modelling

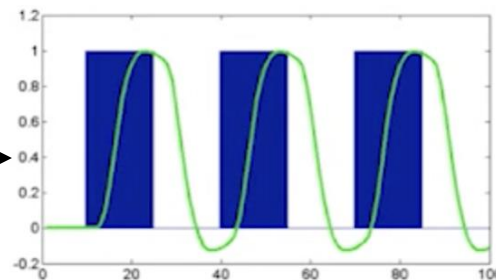
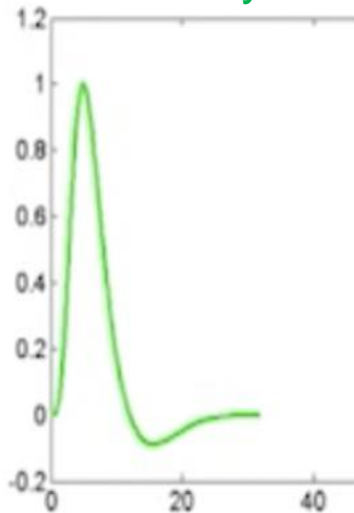


- BOLD responses are delayed: **peak at 4-6 s** and **baseline 20-30 s**.
- Convolved with the hemodynamic response function (HRF)
- The linear time-invariant (LTI) system

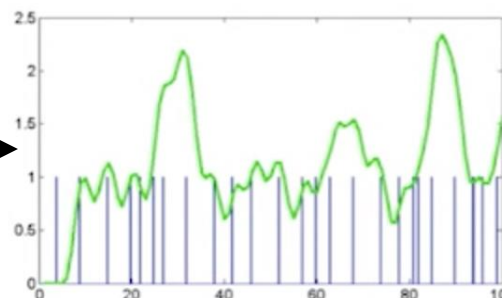
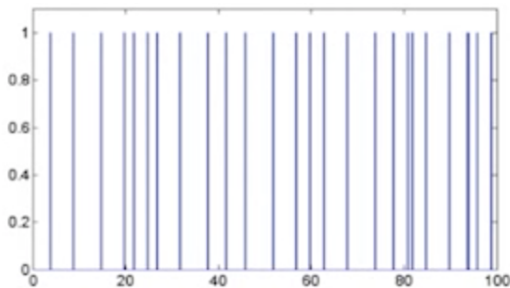
Block design



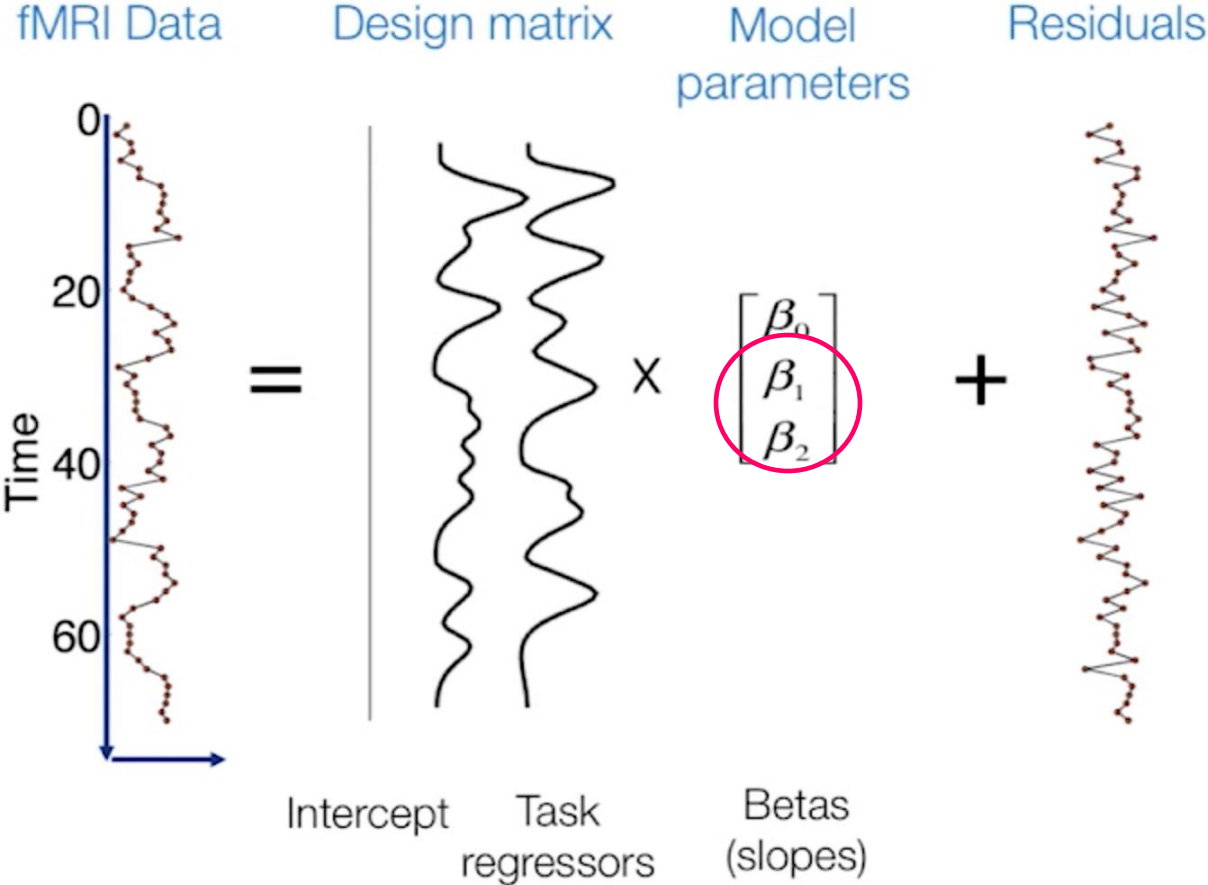
Hemodynamic delay



Event-related design



Event-related design: two continuous regressors (laughter vs. scrambled laughter regressors)



Four example studies

1. Social laughter experiment
2. Naturalistic movie stimuli
3. Repeated measure food-reward responses
4. Resting states

The laughter experiment

Laughter is a contagious behavioural stimulus that is commonly used to study social brain functions. We have studied the social brain functions of participants with high psychopathy or autism traits.

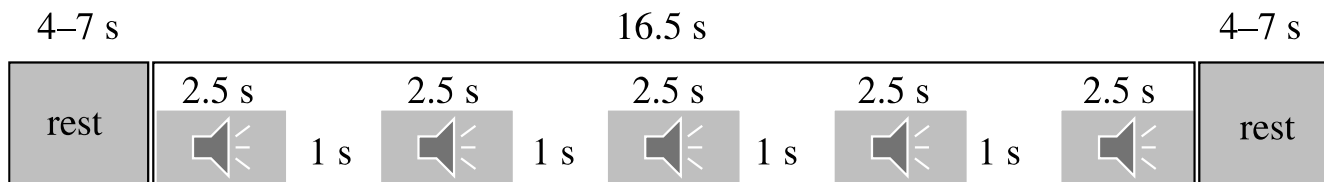
>> Sun L., et al., *Cerebral Cortex*, Volume 33, Issue 2, 15 January 2023

Four stimuli types:

Laughter / Crying vocalization /

Scrambled laughter / Scrambled Crying

Block (also event-related) design

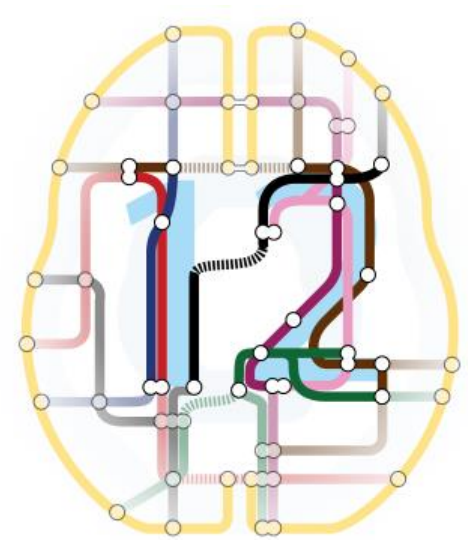


↑
1 event

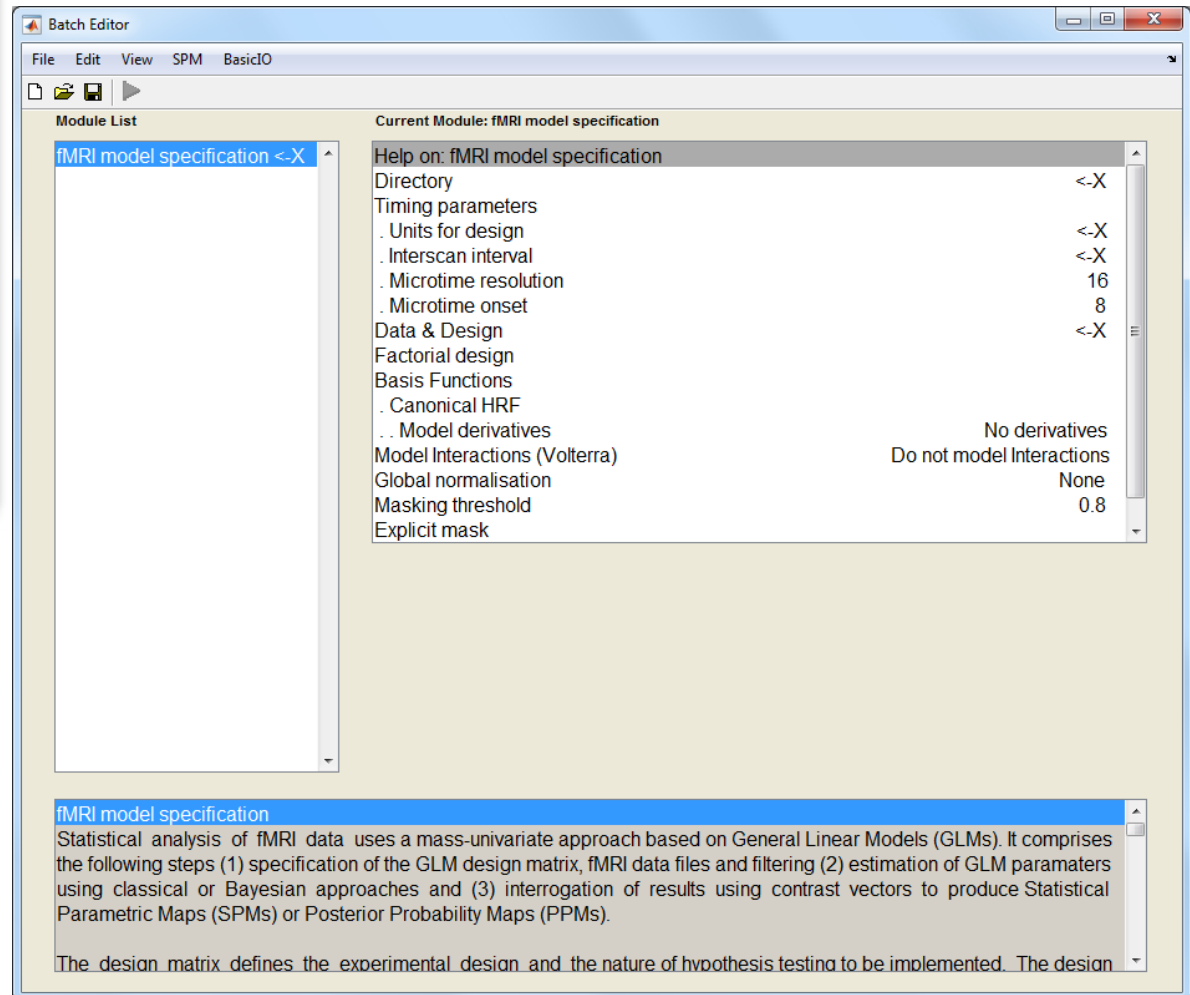
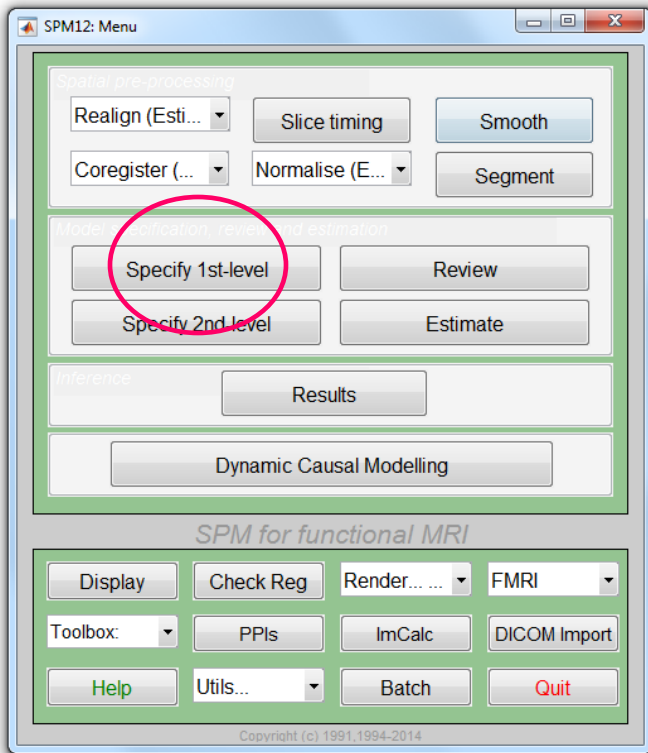
The 16.5 s block contains 5 Laughter, crying, or scrambled sound clips.

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/>



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

} Basic settings

} Statistical model

Regressor building

} Leave as default,
or specify if necessary

No masks

Current Module: fMRI model specification

Data & Design
 . Subject/Session
 . . Scans → All scan volumes <-X
 . . Conditions
 . . . Condition
 Name → Laughter humans
 Onsets <-X
 Durations <-X
 Time Modulation No Time Modulation
 Parametric Modulations
 Orthogonalise modulations Yes
 . . . Condition
 Name → Crying <-X
 Onsets <-X



Data estimation

Module List
Model estimation <-X

Current Module: Model estimation
Help on: Model estimation
Select SPM.mat
Write residuals
Method
. Classical

SPM12: Menu

Realign (Esti... Slice timing Smooth
Coregister (... Normalise (E... Segment

Specify 1st-level Review
Specify 2nd-level Estimate

Results

Dynamic Causal Modelling

SPM for functional MRI

Display Check Reg Render... FMR
Toolbox: PPIs ImCalc DICOM Import
Help Utis... Batch Quit

Batch Editor

File Edit **View** SPM BasicIO

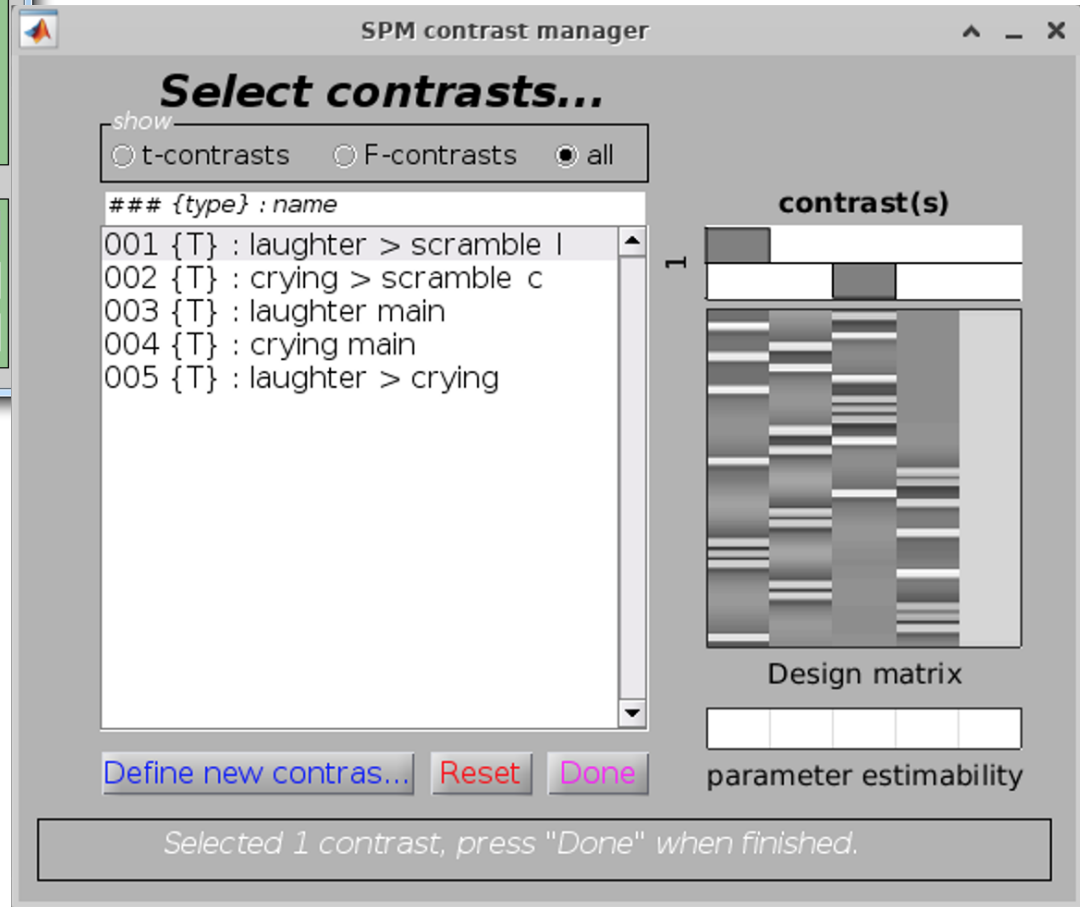
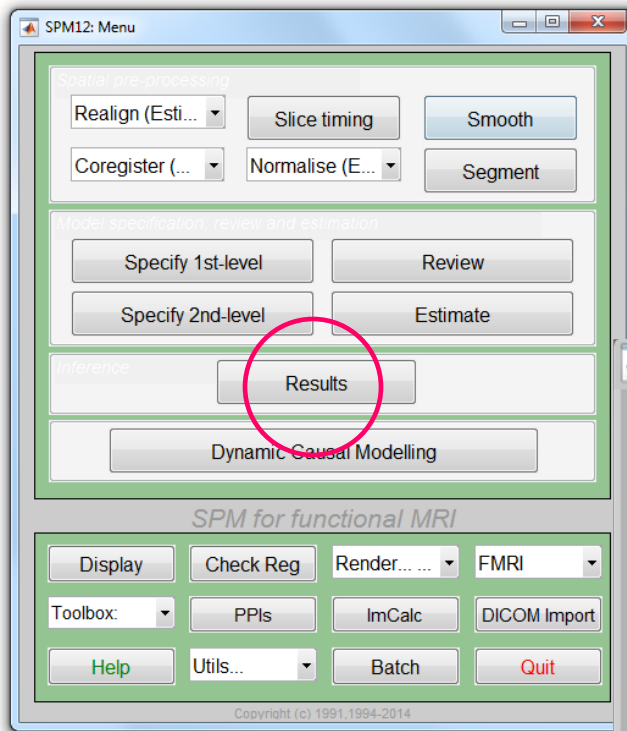
Update View Ctrl+U
Change Font/Fontsize Ctrl+F
Expert Edit Ctrl+E
Show .m Code Ctrl+S

Module: fMRI mode
fMRI model specification

Timing parameters

```
matlabbatch{1}.spm.stats.fmri_spec.dir = {output_directory};  
matlabbatch{1}.spm.stats.fmri_spec.timing.units = 'secs';  
matlabbatch{1}.spm.stats.fmri_spec.timing.RT = 2.6;  
matlabbatch{1}.spm.stats.fmri_spec.timing.fmri_t = 45;  
matlabbatch{1}.spm.stats.fmri_spec.timing.fmri_t0 = 23;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.scans = epi_files;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(1).name = 'laughter';  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(1).onset = onset_laughter;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(1).duration = duration_laughter;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(1).tmod = 0;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(1).pmod = struct('name', {}, 'param', {}, 'poly', {});  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(1).orth = 1;  
  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(2).name = 'crying';  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(2).onset = onset_crying;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(2).duration = duration_crying;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(2).tmod = 0;  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(2).pmod = struct('name', {}, 'param', {}, 'poly', {});  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(2).orth = 1;  
  
matlabbatch{1}.spm.stats.fmri_spec.ssess.cond(3).name = 'scramble';
```


SPM12: setting contrasts

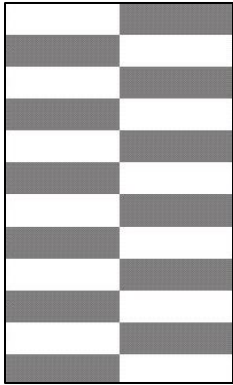


Contrasts

Experimental conditions: stimuli or interest vs. control stimuli

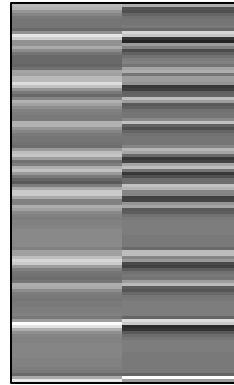
- Your interest is often the difference between the two conditions, which is “contrast”
 - You can calculate the difference, sum or separately each conditions, which are calculated by different linear contrasts.
-
- We only introduce T contrast in this lecture !!

Condition 1
Condition 2
Intercept



Block design

Condition 1
Condition 2
Intercept



Event-related design

Design matrix in
SPM 12

Difference between
conditions

$$[1 \ -1] = \text{“Con1 > Con2”}$$

$$[-1 \ 1] = \text{“Con1 < Con2”}$$

Separately

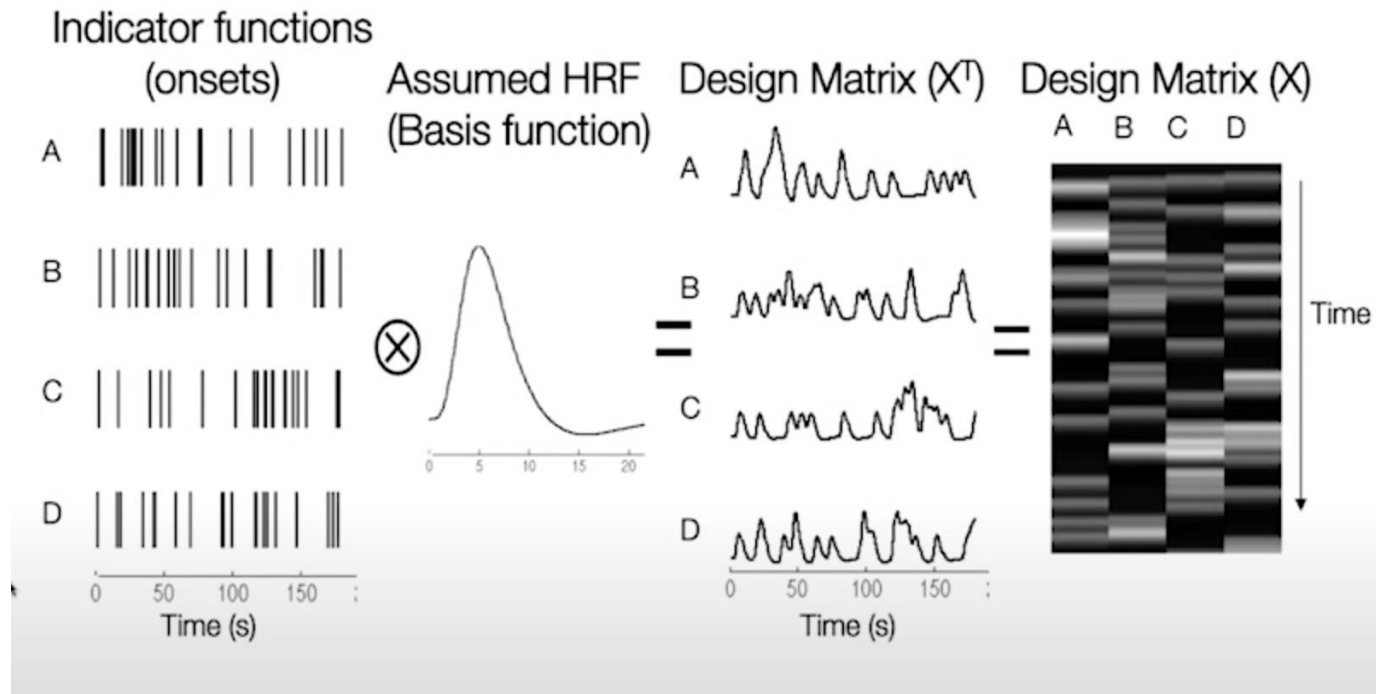
$$[1 \ 0] \text{ or } [-1 \ 0] = \text{“main effect Con1”}$$

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

Sum

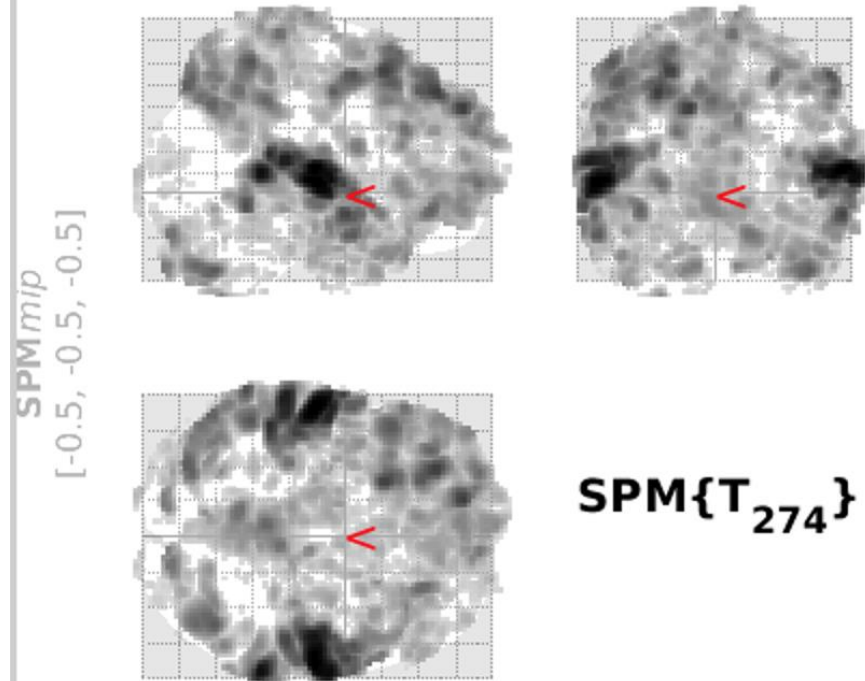
$$[1 \ 1] \text{ or } [-1 \ -1]$$

More regressors

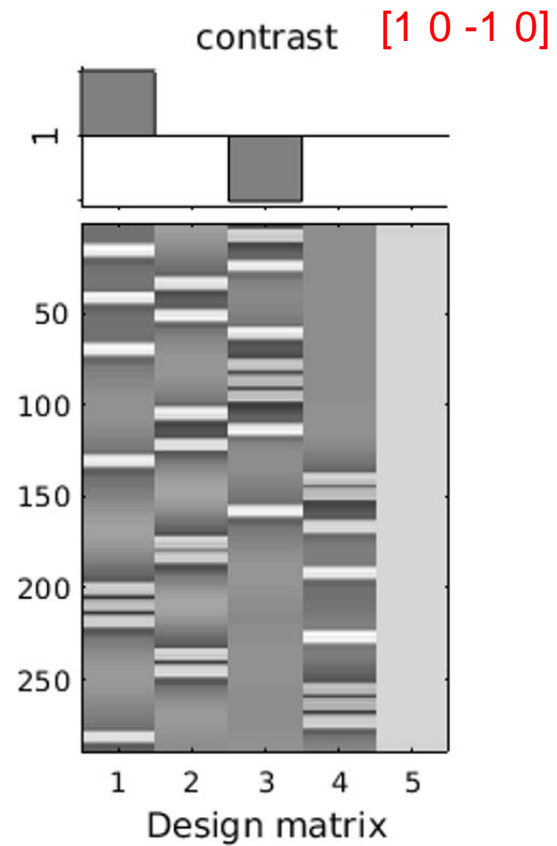


- $[1 \ 1 \ -1 \ -1]$: $(A+B) > (C+D)$
- $[1 \ -1 \ 1 \ -1]$: $(A+C) > (B+D)$
- $[1 \ 0 \ 0 \ 0]$: main effect of A
- $[1 \ 1 \ 0 \ 0]$: Sum of $(A+B)$ vs the mean of the signal
-

laughter > scramble_I



SPMresults: ./sub-201
 Height threshold $T = 1.650434$ { $p < 0.05$ (unc.)}
 Extent threshold $k = 0$ voxels



Example 2: Naturalistic stimuli

Naturalistic fMRI offers **ecological validity**, engages **complex brain functions**, **richer data**, **better participant engagement**, etc.

>> *Santavirta S. et al., NeuroImage, Volume 272, 15 May 2023, 120025*

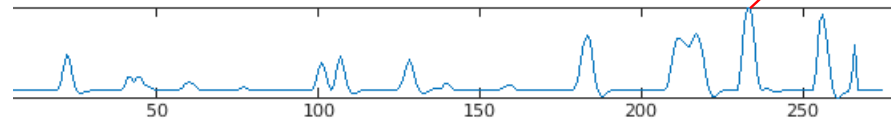
>> *Nummenmaa L., et al., Cerebral Cortex, Volume 31, Issue 9, September 2021, Pages 4104–4114*

Movie-based fMRI

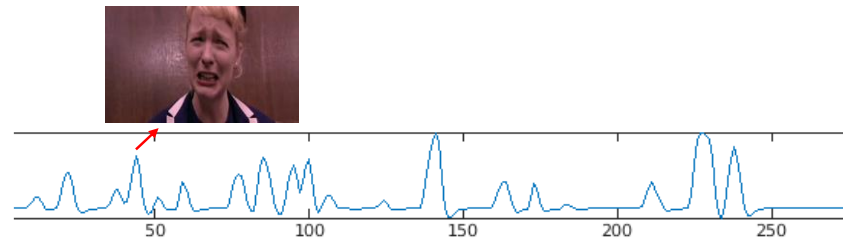
- ✓ Ratings of different dimensions (social, emotional, neutral, objective...)
- ✓ Each regressor should contain certain number of stimuli. CAN NOT be too small number!

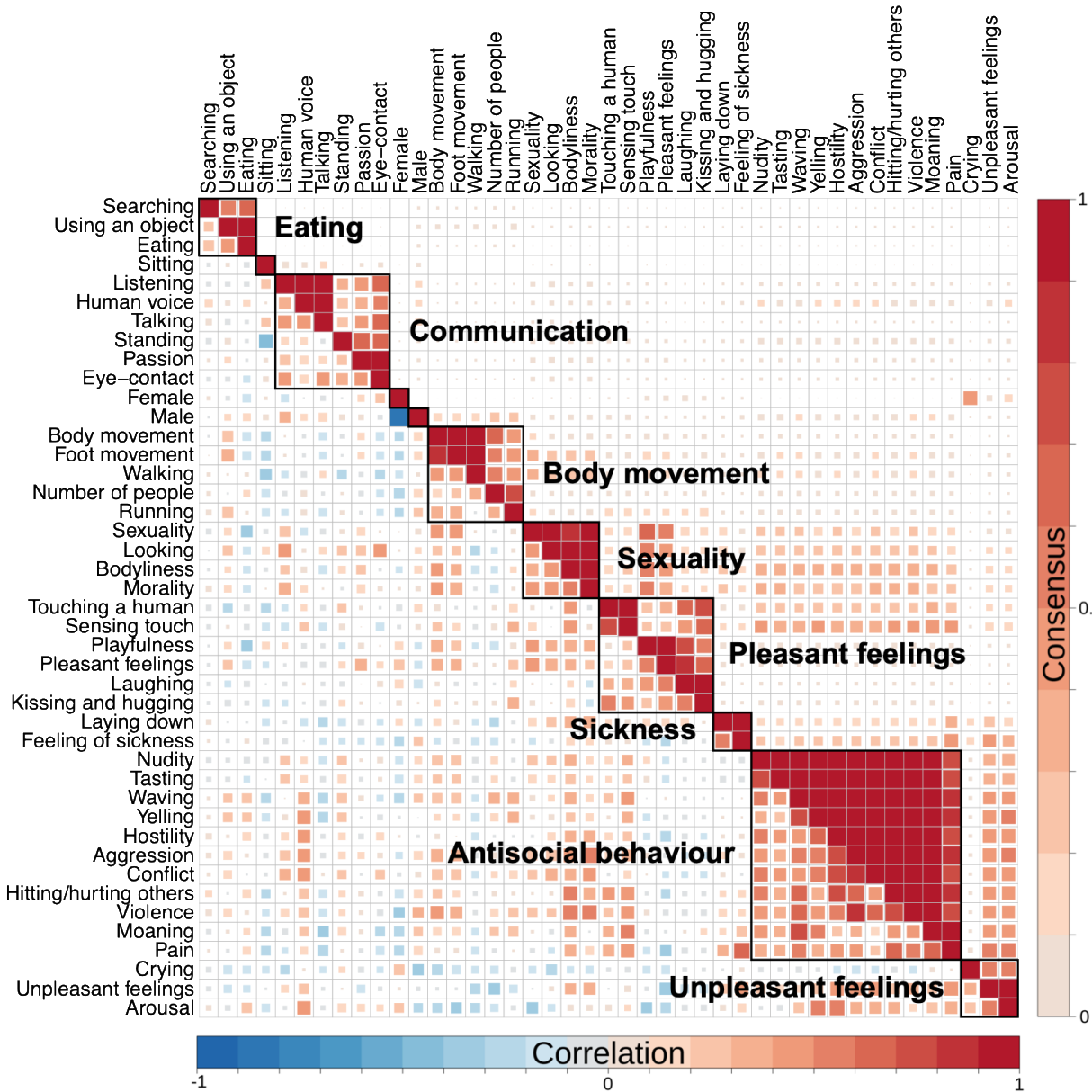


Self-control



Wanting





PCA analysis

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 (M)

Global normalisation

Masking threshold

Explicit mask

Result to store

scan

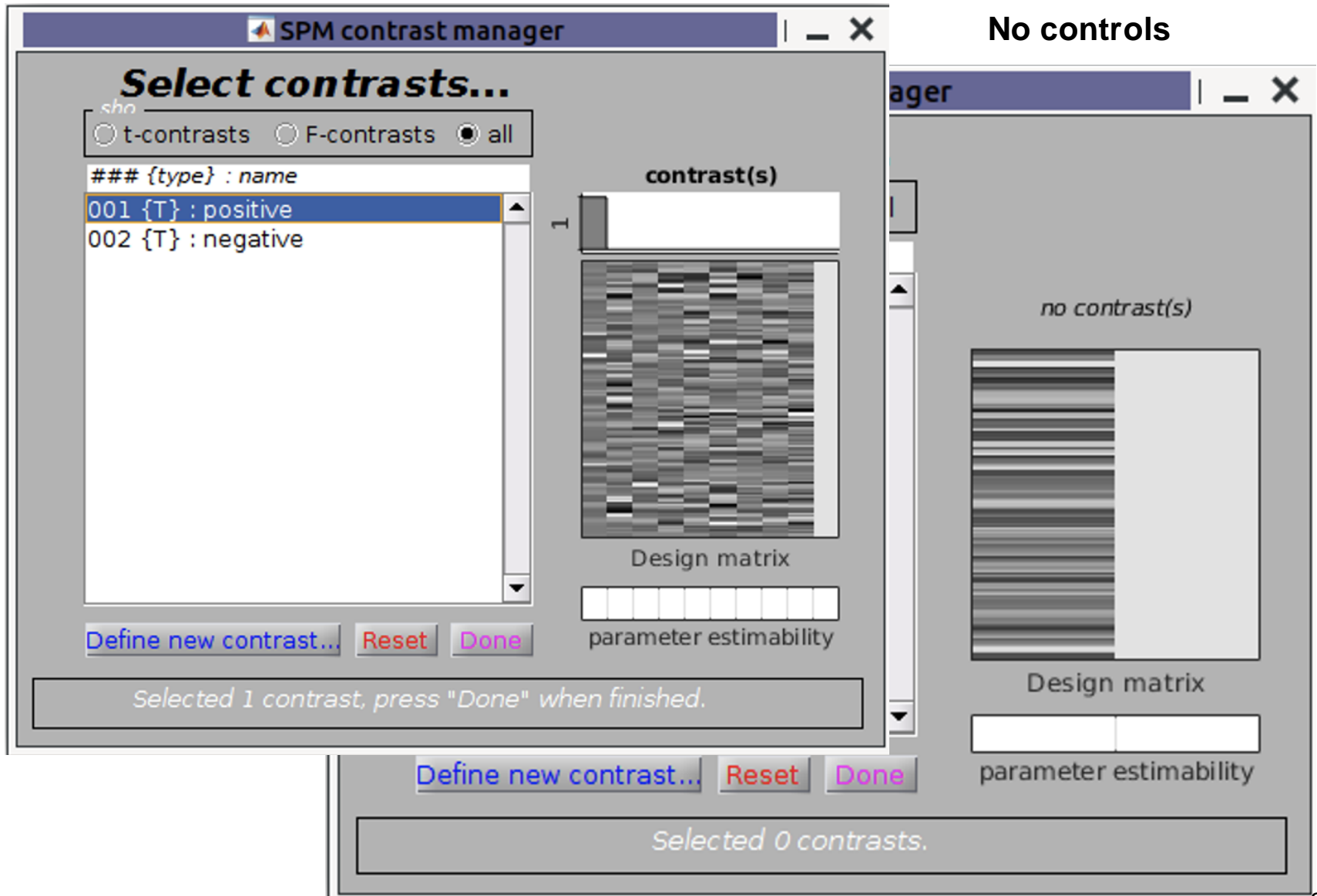
Dependent variables

Current Module: fMRI model specification

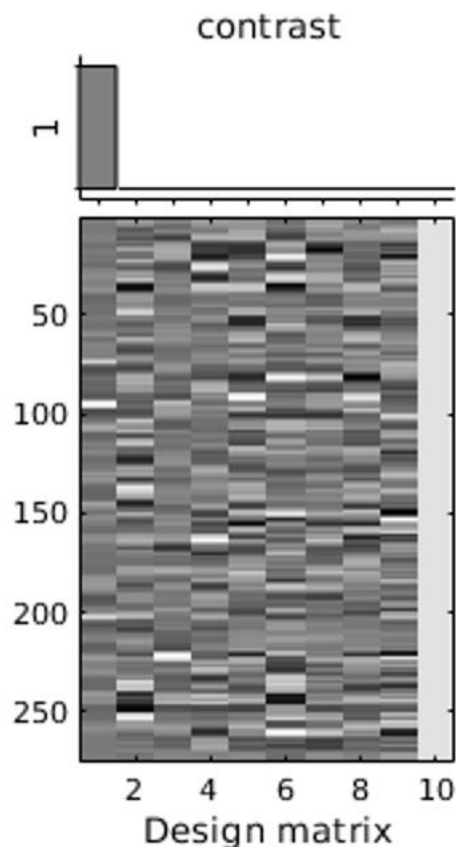
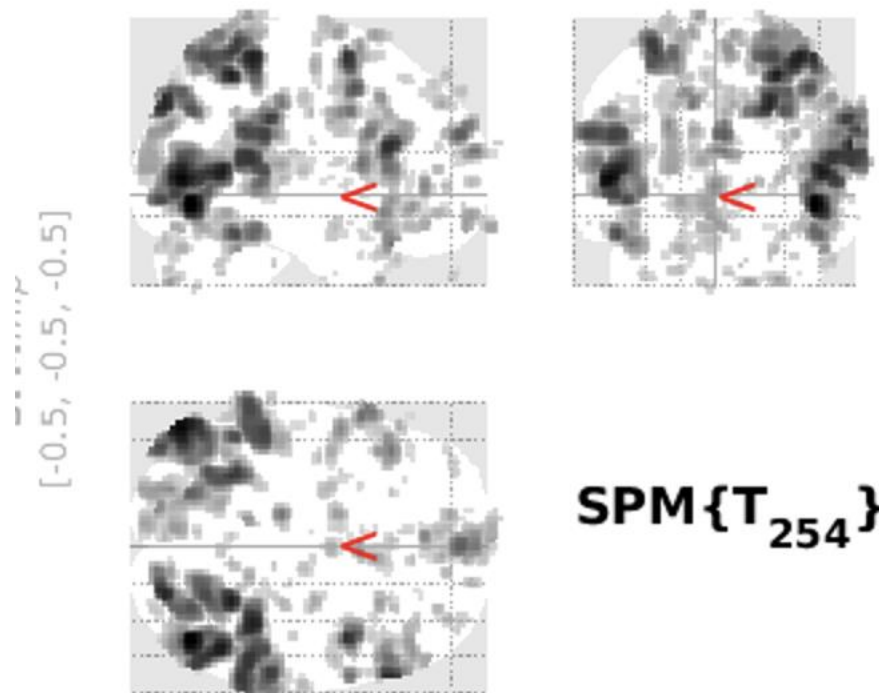
Data & Design

- . Subject/Session
- .. Scans → All scan volumes <-X
- .. Conditions
- ... Condition
- Name → e.g. Self control humans
- Onsets <-X
- Durations → The same as volume No. <-X
- Time Modulation No Time Modulation
- Parametric Modulations
- Orthogonalise modulations Yes

Controlling for low-level regressors



positive



SPMresults: ./Pleasant_feelings/sub-101
 Height threshold $T = 1.650875$ { $p < 0.05$ (unc.)}
 Extent threshold $k = 0$ voxels

Statistics: *p-values adjusted for search volume*

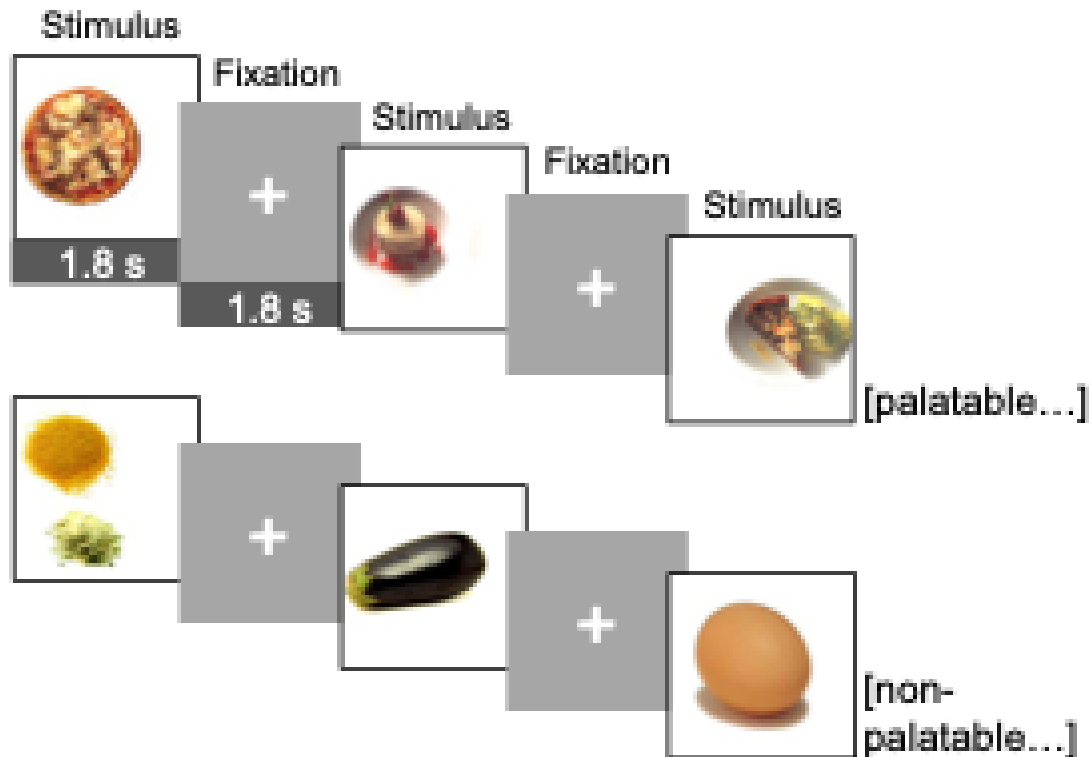
set-level		cluster-level				peak-level					mm mm mm		
ρ	c	$\rho_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	k_E	ρ_{uncorr}	$\rho_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	T	$(Z_{\text{=}})$	ρ_{uncorr}			
1.000	100	0.009	0.002	1576	0.000	0.026	0.025	5.21	5.07	0.000	46	-72	-6
						0.211	0.055	4.67	4.57	0.000	48	-56	8
						0.701	0.100	4.23	4.15	0.000	54	-66	6
		0.034	0.005	1272	0.000	0.070	0.034	4.97	4.85	0.000	-52	-76	6

Example 3: Repeated measure food-reward

Food reward experiment has been used in decoding the brain conceptualization of satiation as induced by secretin hormone.

>> *Lauri S., Sun L., et al., Nature Metabolism, volume 3, pages 798–809 (2021)*

Twelve 16.2s blocks for each food category
Blocks in pseudo-randomized order
6 food stimuli intermixed with 3 fixations in one block



- Control condition vs. condition with infusion of secretin.
- Each subject **scanned twice** with fMRI: one in control condition, the other under intervention (secretin infusion).

Current Module: fMRI model specification

Data & Design

- . Subject/Session
 - .. Scans
 - .. Conditions
 - ... Condition
 - Name
 - Onsets
 - Durations
 - Time Modulation
 - Parametric Modulations
 - Orthogonalise modulations
 - .. Multiple conditions
 - .. Regressors
 - .. Multiple regressors
 - .. High-pass filter
- . Subject/Session
 - .. Scans
 - .. Conditions

Current Item: Data & Design

- New: Subject/Session
- Replicate: Subject/Session (1)
- Replicate: Subject/Session (2)
- Delete: Subject/Session (1)
- Delete: Subject/Session (2)

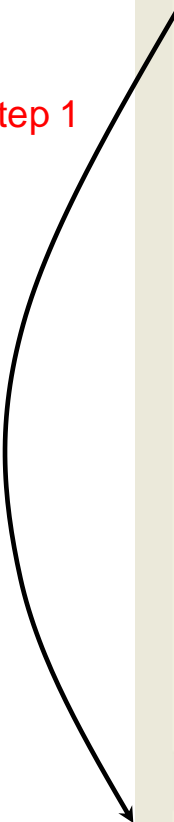
Current Module: fMRI model specification

- .. Scans
- .. Conditions**
 - ... Condition
 - Name
 - Onsets
 - Durations
 - Time Modulation
 - Parametric Modulations
 - Orthogonalise modulations
 - .. Condition
 - Name
 - Onsets
 - Durations
 - Time Modulation
 - Parametric Modulations
 - Orthogonalise modulations
 - .. Multiple conditions
 - .. Regressors
 - .. Multiple regressors

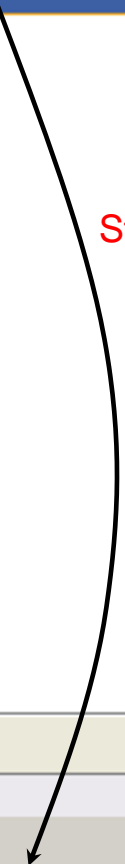
Current Item: Conditions

- New: Condition
- Replicate: Condition (1)
- Replicate: Condition (2)
- Delete: Condition (1)
- Delete: Condition (2)

Step 1



Step 2

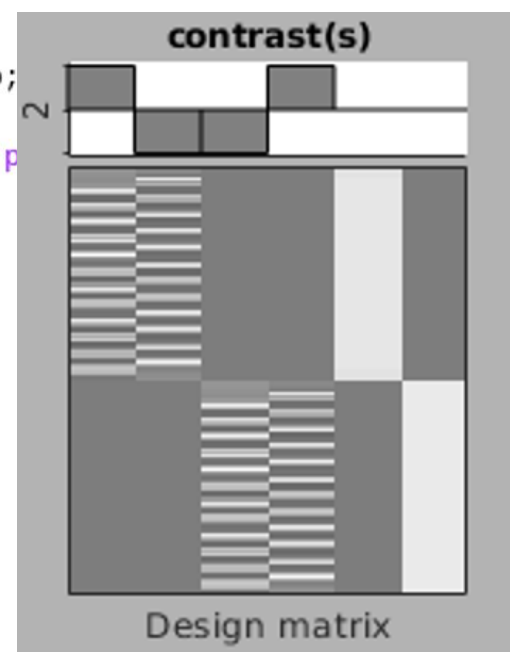


```

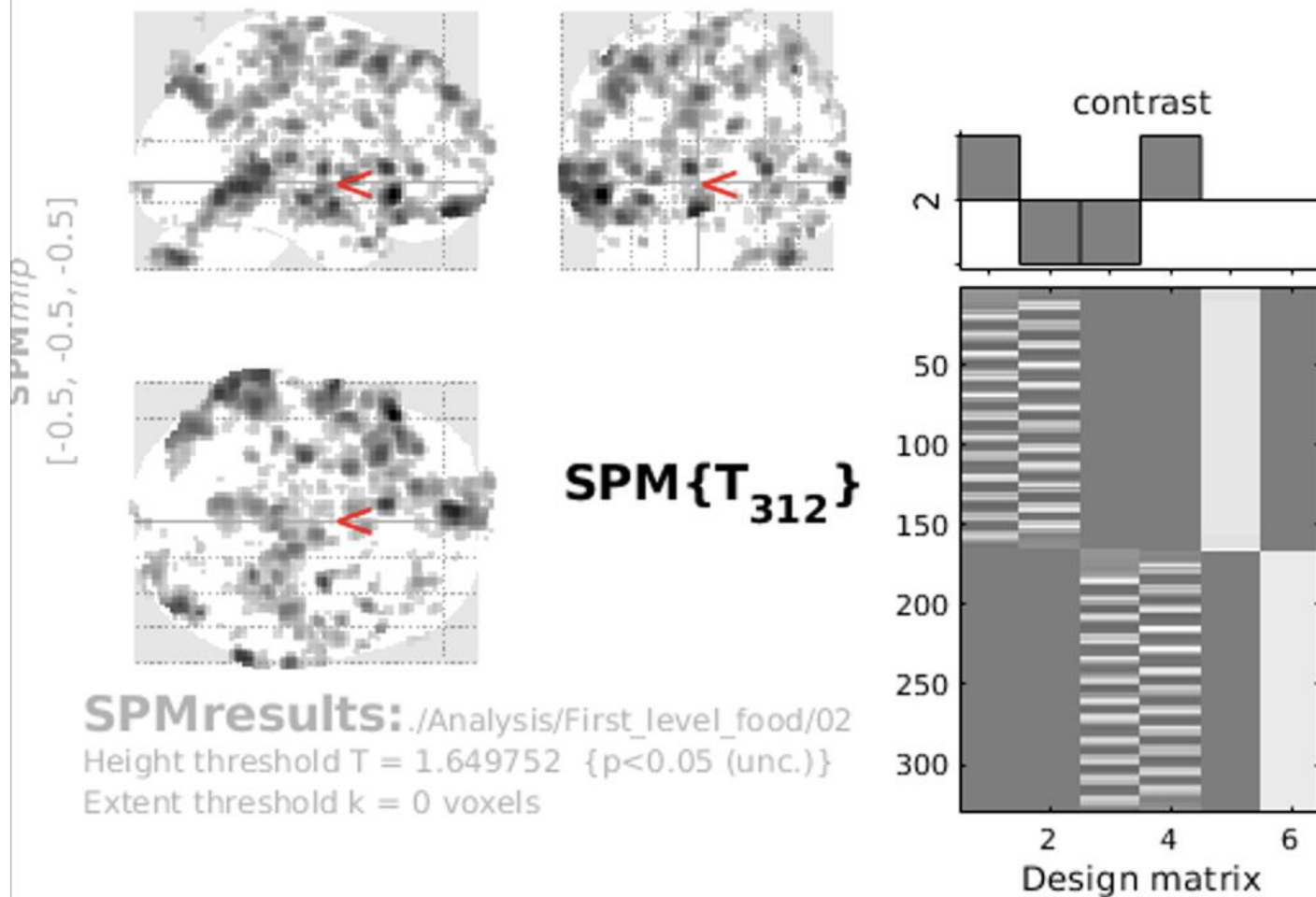
matlabbatch{1}.spm.stats.fmri_spec.sess(1).scans = epi_files1;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(1).name = 'appetite';
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(1).onset = onset_apep;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(1).duration = duration_apep;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(1).tmod = 0;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(1).pmod = struct('name', {}, 'param', {}, 'poly', {});
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(1).orth = 1;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(2).name = 'bland';
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(2).onset = onset_bland;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(2).duration = duration_bland;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(2).tmod = 0;
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(2).pmod = struct('name', {}, 'param', {}, 'poly', {});
matlabbatch{1}.spm.stats.fmri_spec.sess(1).cond(2).orth = 1;

matlabbatch{1}.spm.stats.fmri_spec.sess(2).scans = epi_files2;
matlabbatch{1}.spm.stats.fmri_spec.sess(2).cond(1).name = 'appetite';
matlabbatch{1}.spm.stats.fmri_spec.sess(2).cond(1).onset = s2_onset_apep;
matlabbatch{1}.spm.stats.fmri_spec.sess(2).cond(1).duration = s2_duration_apep;
matlabbatch{1}.spm.stats.fmri_spec.sess(2).cond(1).tmod = 0;
matlabbatch{1}.spm.stats.fmri_spec.sess(2).cond(1).pmod = struct('name', {}, 'p

```



interaction



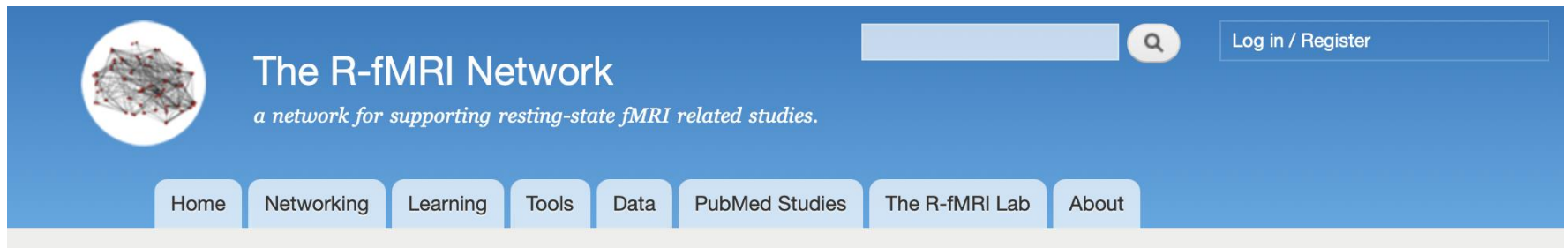
Example 4: resting state fMRI

- ReHo = regional homogeneity: larger value indicates a higher regional synchronization.
- ALFF = amplitude of low-frequency fluctuation: indicate the magnitude of neural activity
- **FC = functional connectivity (between ROIs): inter-regional correlations**



Toolbox for rs-fMRI analysis

- Matlab based
- Download DPARSF from: <http://rfmri.org/DPARSF>



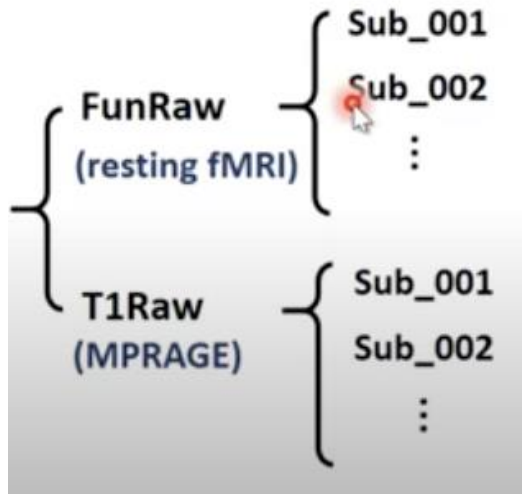
The R-fMRI Network
a network for supporting resting-state fMRI related studies.

Home Networking Learning Tools Data PubMed Studies The R-fMRI Lab About

Log in / Register

The screenshot shows the header of the website 'The R-fMRI Network'. It features a blue background with a circular logo on the left containing a network diagram. To the right of the logo is the site title and a tagline. Further right is a search bar and a 'Log in / Register' button. Below this is a horizontal navigation menu with buttons for 'Home', 'Networking', 'Learning', 'Tools', 'Data', 'PubMed Studies', 'The R-fMRI Lab', and 'About'.

Data folder



Data Processing Assistant for Resting-State fMRI

Advanced Edition DPARSF A

Working Directory: ...

Participants:

Time Points:
TR (s):

Template P... EPI DICOM to NIFTI T1 DICOM to NIFTI BIDS to DPARSF

Apply Mats Remove First Time Points Slice Timing Slice Number: Slice Order:
Reference Slice: FieldMap Correction Realign Voxel-Specific Head Motion

Reorient Fun* AutoMask Crop T1 Reorient T1* Bet T1 Coreg to Fun

Segment New Segment + DARTEL East Asian European

Nuisance Covariates Regression Head Motion model: Rigid-body 6 Derivative 12
 Friston 24 Voxel-specific 12 Head motion scrubbing regressors

Other covariates Add mean back Filter (Hz): ~

Normalize
 Normalize by using EPI templates Normalize by using T1 image unified segmentation Normalize by DARTEL

Smooth Smooth by DARTEL

Default mask No mask User-defined mask ... Warp Masks into Individual Space

Detrend Nuisance Covariates Regression ALFF+fALFF ~ Filter

Scrubbing ReHo Smooth ReHo Degree Centrality

Functional Connectivity Extract ROI time courses Define ROI Interactively* CWAS

Normalize to Symmetric Template Smooth VMHC Normalize Derivatives Smooth Derivatives

Parallel Workers #: Functional Sessions #: Starting Directory Name:

Summary

- The first level analysis is a **within-subject analysis, necessary due to the low signal-noise ratios.**
- **Experimental design decides the statistical model.**
- We have showed 4 example studies on how to conduct the first level analysis.
- Using **Contrasts** to view the results
- Contrast images are ready for second level analysis.

References

- Sun, Lihua, et al. "Aberrant motor contagion of emotions in psychopathy and high-functioning autism." *Cerebral Cortex* 33.2 (2023): 374-384.
- Santavirta, Severi, et al. "Functional organization of social perception networks in the human brain." *NeuroImage* 272 (2023): 120025.
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Thanks!

Correspondence:

lihua.sun@utu.fi; sunlihua@fudan.edu.cn