



fMRI analysis

GitHub https://github.com/dcdace/fMRI-COGNESTIC-23

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DataOrganise & Manage



Pre-process

Analyse

BOLD signal = X * b + errors

explained variation unexplained variation
task-related activity changes noise (other changes)

What we know?

- BOLD signal: we collect this from the brain (functional data)
- X: the design matrix (each column is a predictor that we build ourselves)

What we want to find?

• b: vector of beta-weights (one weight for predictor in X) that give the best approximation of the BOLD signal

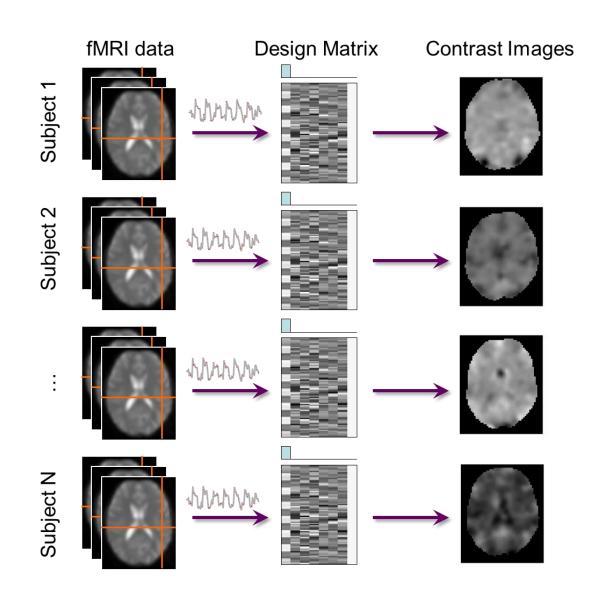
• How we find it?

 By minimising the sum of squared errors. In practice, the GLM has a formula, which guarantees to find these beta-weights

- 1. Extract the signal time-series from a given voxel
- 2. Run GLM (the signal and the design matrix are the inputs) to find beta-weights that best approximate the true signal
- 3. Define your **contrast** and test it
- 4. Repeat for all voxels
 - Produces an image file with contrast values for each voxel: contrast-maps

First-level analysis

• Run the GLM for each subject



First-level analysis

Let's see the nbo5_Subject-Level-Analysis.ipynb notebook

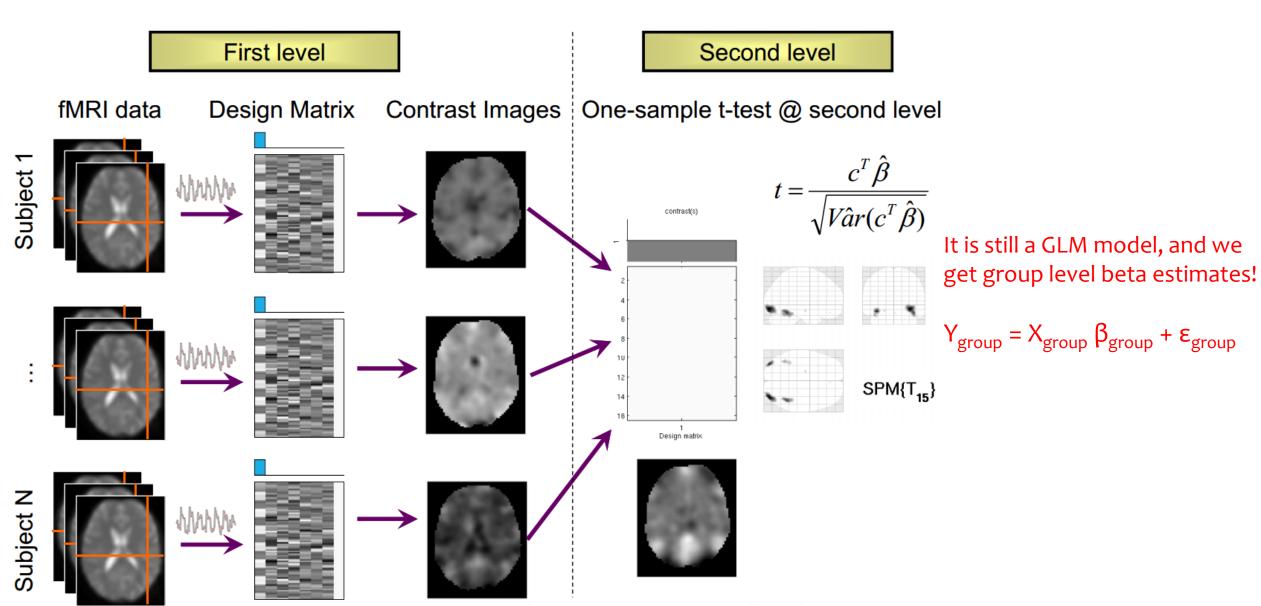


Group level (2nd level) analysis is across subjects

- Which voxels are showing significant activation differences between our conditions consistently within a group
 - Average contrast effect across the sample (e.g., one-sample ttest)

• Importantly, all subject brains need to be in a common space, e.g. MNI, to perform voxel-wise group analyses

Summary statistics, Random effects approach



Generalisability, Random Effects & Population Inference. Holmes & Friston, NeuroImage,1998.

Stats tests at the 2nd level

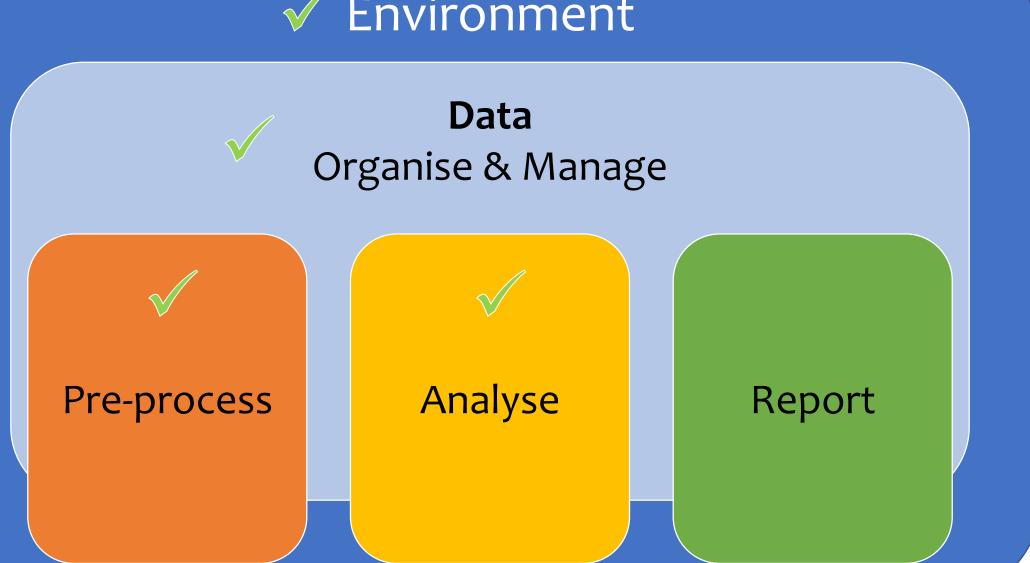
- Condense where possible
 - If a factor can be collapsed through a contrast at the 1st level, do so and use the simplest possible 2nd level model
 - T-tests at the 2nd level are preferred
 - Avoids need to estimate non-sphericity to account for within-subject correlations across repeated measures
 - Generally more accurate estimation of error

- However, if more than 2 factors or levels exist, a single tcontrast cannot capture the main effects and interactions
 - 2nd level ANOVA will be necessary

Let's see the nbook _Group-Level-Analysis.ipynb notebook







Sharing & Reporting



• Share your code and notebooks on GitHub



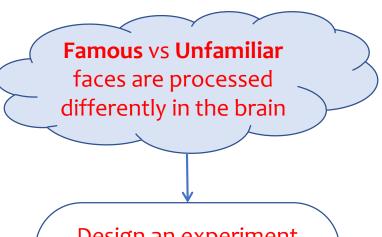
- Make it citable with Zendono
 - https://docs.github.com/en/repositories/archiving-a-github-repository/referencing-and-citing-content



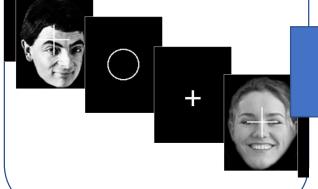
• If you have consent from participants, share the **BIDS data** on OpenNeuro



• Add your contrast maps to NeuroVault







Stimuli Timing

Collect the MRI data



Anatomical image Functional images Event details

