

FMRI Pre-Statistics

Brief introduction to Task FMRI experiments and analysis







FMRI Experiments



- Simple paradigm design:
 - stimulus vs baseline
 - constant stimulus "intensity"
 - constant block lengths
 - many repetitions: ABABA
- Need baseline (rest) condition to measure *change*

The Haemodynamic Response



- Field changes (perturbations) --> dephasing --> T_2^* effect
- BOLD-tuned MRI (T_2^* -weighted) is sensitive to this effect



Predicted Response

• The process can be modelled by convolving the activity curve with a "haemodynamic response function" or HRF





FMRI Experiments: Analysis

• Each voxel contains a time-varying signal (BOLD signal)





FMRI Experiments: Analysis

- Each voxel contains a time-varying signal (BOLD signal)
- Model the stimulus-induced change in BOLD signal (predicted response)



- Find which voxels have signals that match the model
- Good match implies activation related to stimulus





Standard GLM Analysis

- Correlate model at each voxel separately
- Measure residual noise variance
- *t*-statistic = model fit / noise amplitude









Standard GLM Analysis

- Correlate model at each voxel separately
- Measure residual noise variance
- t-statistic = model fit / noise amplitude
- Threshold *t*-stats and display map

Signals of no interest (e.g. artifacts) can affect both activation strength and residual noise variance

Use pre-processing to reduce/eliminate some of these effects





FMRI Pre-Statistics

Summary:

- Task experiment involves stimuli that is:
 - repeated many times
 - includes two or more conditions
- FMRI is based on BOLD effect
- HRF models the delay and spread of blood response
- GLM matches predicted response with data at each voxel to get amplitude
- Ratio with residual error gives statistic
- Thresholded values give map of significant voxels