

General Linear Model for fMRI: bases of statistical analyses

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Objectives

- Intuitive understanding of the GLM
- Get an idea how t-tests, ANOVA, regressions, etc .. are instantiation of the GLM
- Learn key concepts: linearity, model, design matrix, contrast, colinearity, orthogonalization

Overview

- What is linearity?
- Why do we speak of models?
- A simple fMRI model
- Contrasts
- Issues with regressors

What is linearity?

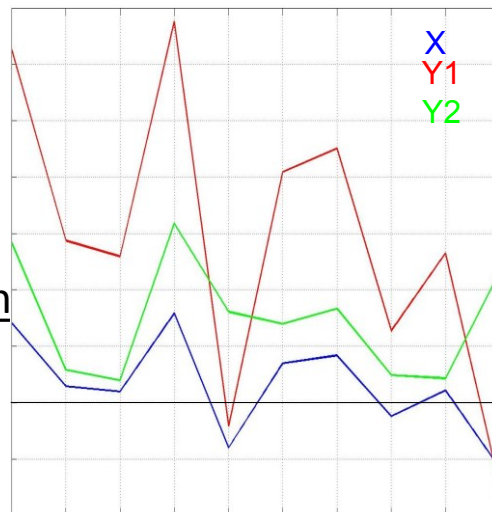
Linearity

- Means created by lines
- In maths it refers to equations or functions that satisfy 2 properties: additivity (also called superposition) and homogeneity of degree 1 (also called scaling)
- Additivity $\rightarrow y = x_1 + x_2$ (output is sum of inputs)
- Scaling $\rightarrow y = \beta x_1$ (output is proportional to input)

<http://en.wikipedia.org/wiki/Linear>

Examples of linearity / non linearity

- $X = \text{randn}(10,1)$
- Linear correlation
- $Y_1 = 3x + 2$
- Pearson $r = 1$
- Non linear correlation
- $Y_2 = \text{abs}(2x)$
- Pearson $r = 0.38$



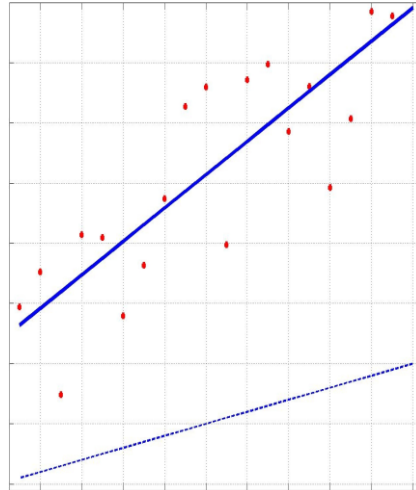
What is a linear model?

What is a linear model?

- An equation or a set of equations that models data and which corresponds geometrically to straight lines, plans, hyperplans and satisfy the properties of additivity and scaling.
- Simple regression: $y = \beta_1 x_1 + \beta_2 + \varepsilon$
- Multiple regression: $y = \beta_1 x_1 + \beta_2 x_2 + \beta_3 + \varepsilon$
- One way ANOVA: $y = \mu + \alpha_i + \varepsilon$
- Repeated measure ANOVA: $y = \mu + S_i + \alpha_i + \varepsilon$
- ...

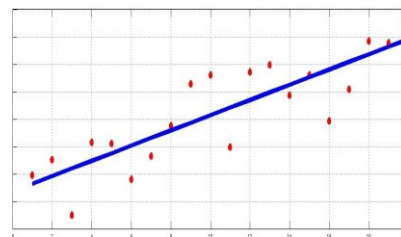
A regression is a linear model

- We have an experimental measure x (e.g. stimulus intensity from 0 to 20)
- We then do the expe and collect data y (e.g. RTs)
- Model: $y = \beta_1 x + \beta_2$
- Do some maths / run a software to find β_1 and β_2
 $\hat{y} = 2.7x + 23.6$

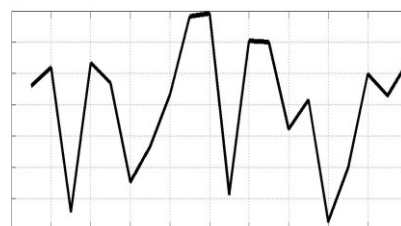


A regression is a linear model

- The error is the distance between the data and the model
- $F = (SS_{\text{effect}} / df) / (SS_{\text{error}} / df_{\text{error}})$
- $SS_{\text{effect}} = \text{norm}(\text{model} - \text{mean}(\text{model}))^2$;
- $SS_{\text{error}} = \text{norm}(\text{residuals})^2$;



Error = $Y - XB$



Summary

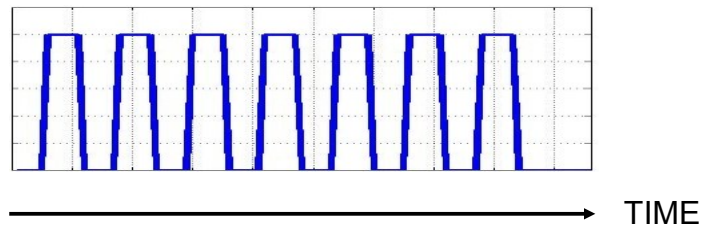
- Linear model: $y = \beta_1 x_1 + \beta_2 x_2$ (output = additivity and scaling of input)

A simple fMRI model

<http://www.fil.ion.ucl.ac.uk/spm/data/auditory/>

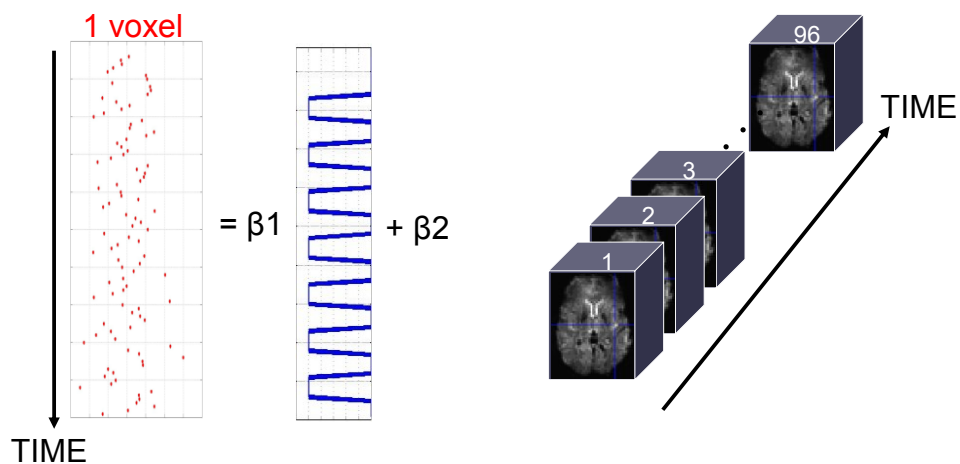
FMRI experiment

- *SPM data set*: which areas are activated by the presentation of bi-syllabic words presented binaurally (60 per minute)
- Experimental measure x : 7 blocks of 42 sec of stimulation

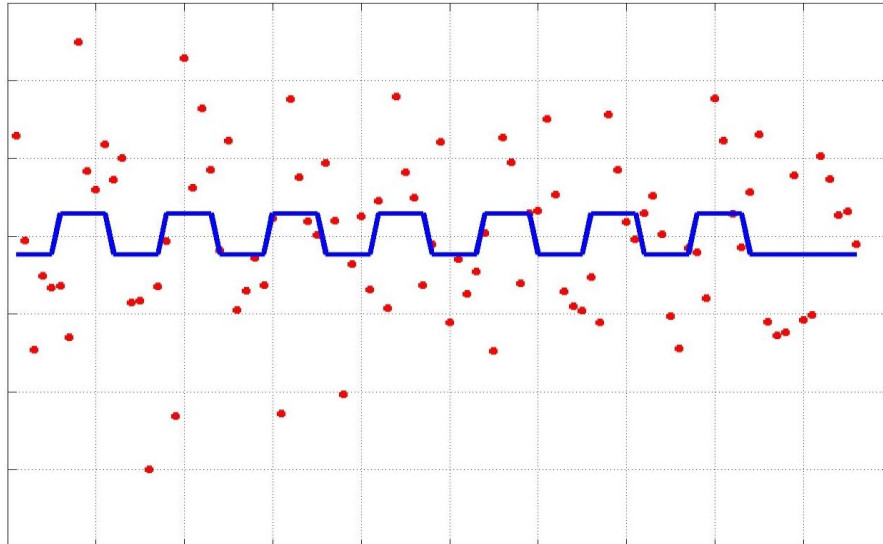


FMRI experiment

- Collect the data : 96 fMRI volumes (RT=7s)
- Model: $y = \beta_1 x + \beta_2$

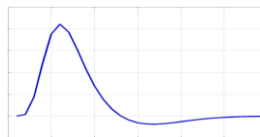


FMRI experiment

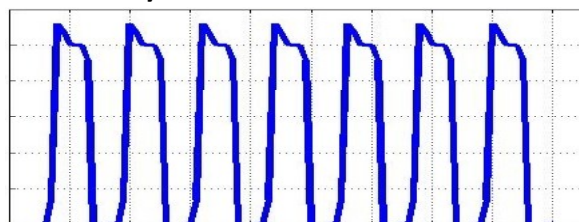


FMRI experiment

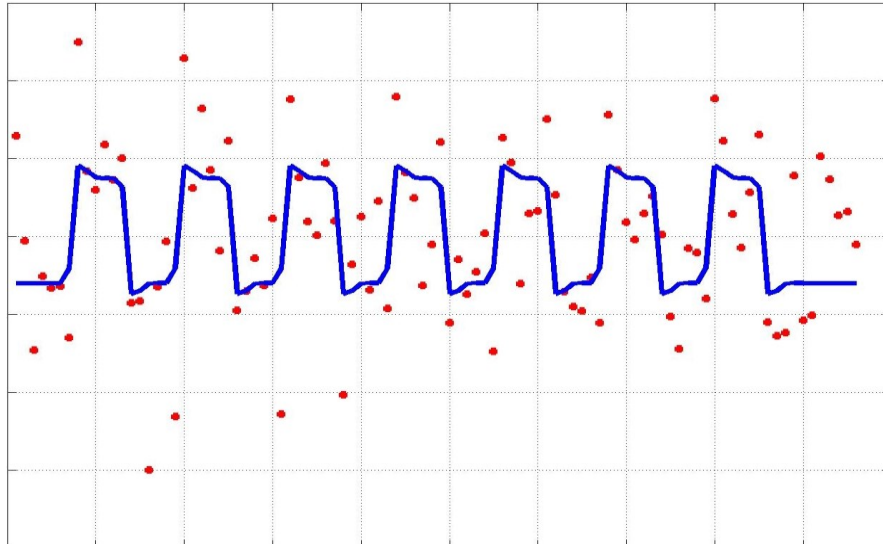
- A better model: we know the shape of the BOLD response



- Convolution by the hrf: $x \otimes \text{hrf}$

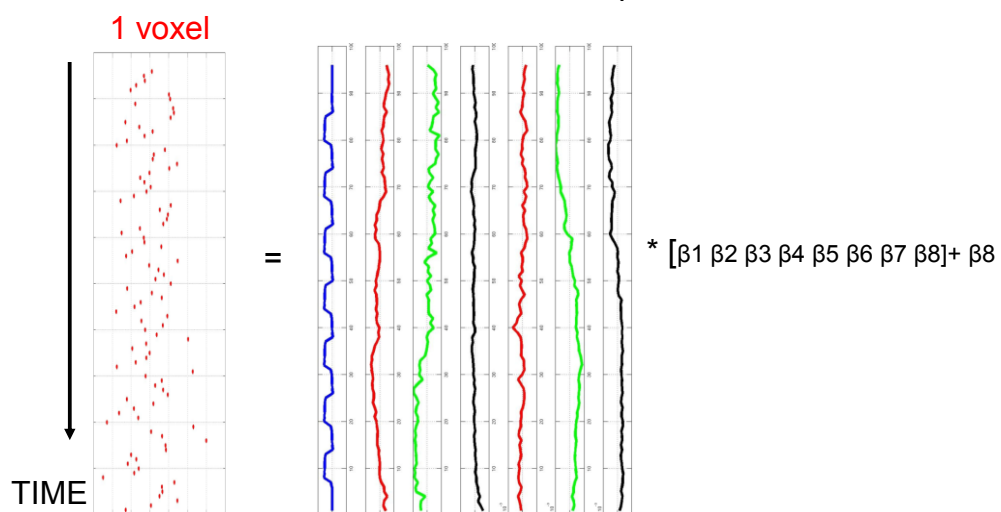


FMRI experiment

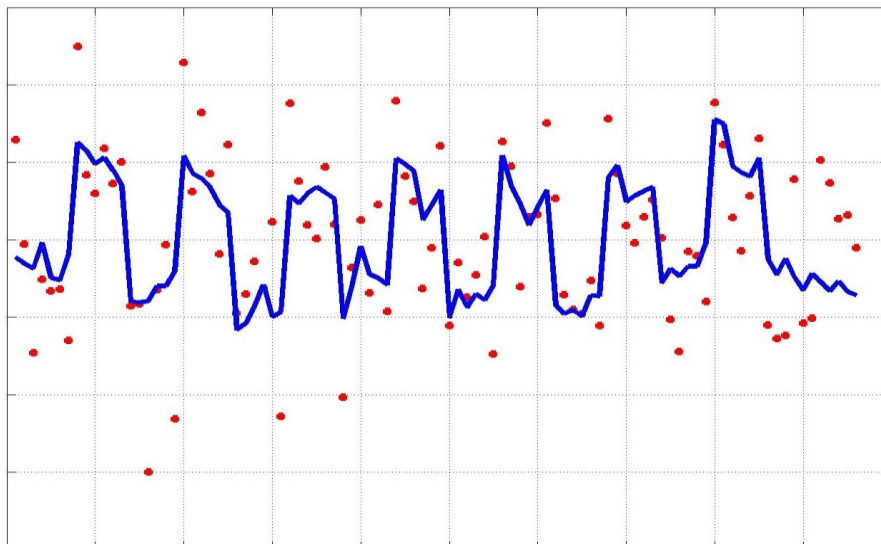


FMRI experiment

- An even better model: add motion parameters

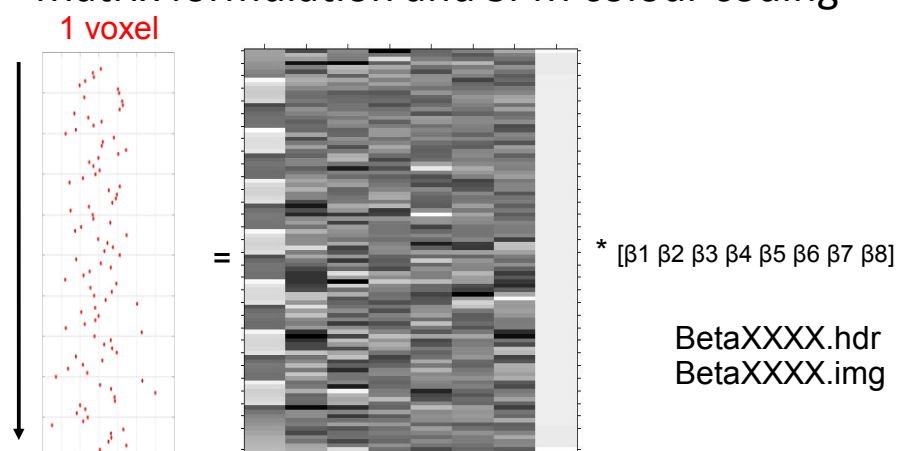


FMRI experiment



FMRI experiment

- Matrix formulation and SPM colour coding

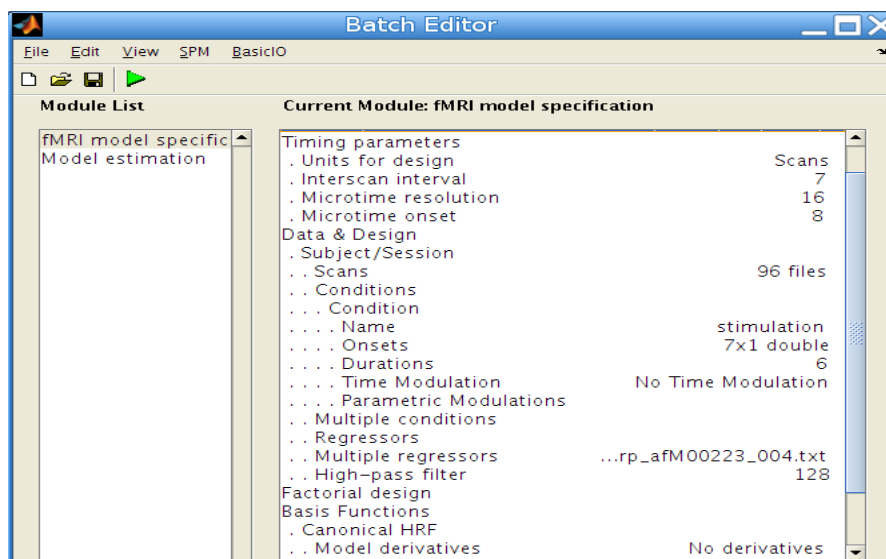


$$\text{FMRI data (Y)} = \text{Design matrix (X = SPM.mat)} * B + E (\text{ResMS})$$

Noise modelling

- GLM: $Y = XB + E$ with $E \sim N(0, V)$
- Because of scanner issue, data often have a drift, that we can incorporate into the model X
- V is the covariance matrix whose depends on your model – in SPM we use an AR(1) model + white noise to remove dependencies related to physiological artefacts (e.g. cardiac aliasing) and model residual noise

FMRI experiment



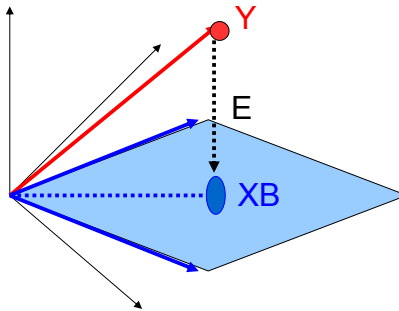
Summary

- Linear model: $y = \beta_1 x_1 + \beta_2 x_2$ (output = additivity and scaling of input)
- GLM: $Y = XB + E$ (matrix formulation, works for any statistics, express the data Y as a function of the design matrix X)

Contrasts

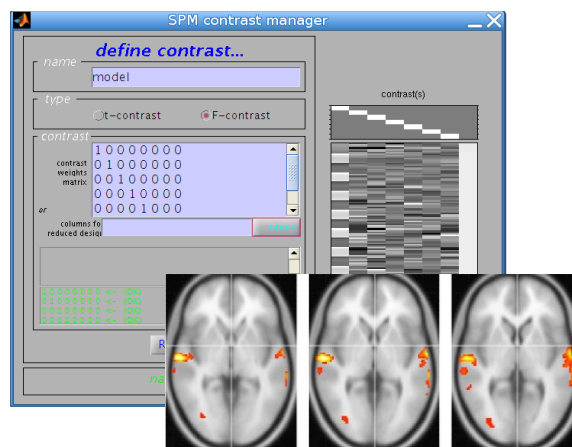
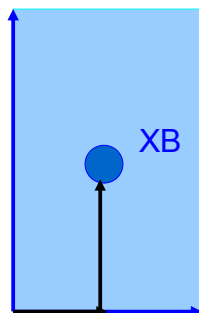
Model = R^2

- Geometrical perspective
- Y = 3 observations X = 2 regressors
- $Y = XB + E$



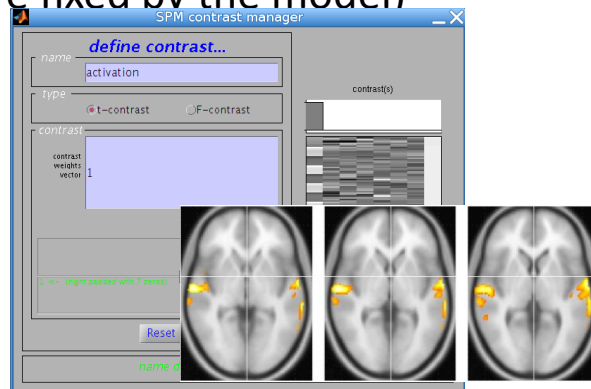
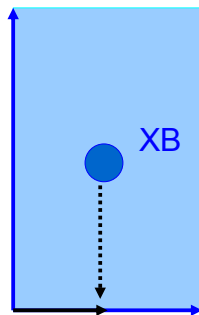
Model = R^2

- Where does the model fit the data?
- F test for all regressors: $y = 1/2x_1 + 1/2x_2 + \varepsilon$



Contrast = effect to test

- Where does the regressor for activation only explain the data (given the model)
- $y = 1/2x_1 + \varepsilon$ (the orientation of x_1 and value of β_1 are fixed by the model)



Summary

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- Contrasts: F or t test for the effect of 1 or several regressors given the design matrix

Issues with regressors

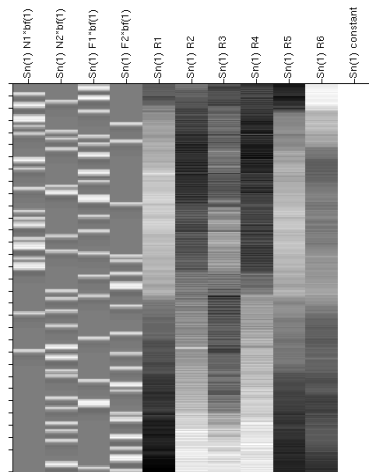
More contrasts

- New experiment: (Famous vs. Nonfamous) x (1st vs 2nd presentation) of faces against baseline of chequerboard
- 2 presentations of 26 Famous and 26 Nonfamous Greyscale photographs, for 0.5s, randomly intermixed, for fame judgment task (one of two right finger key presses).

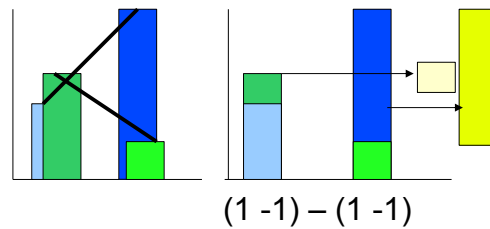
http://www.fil.ion.ucl.ac.uk/spm/data/face_rep/face_rep_SPM5.htm

More contrasts

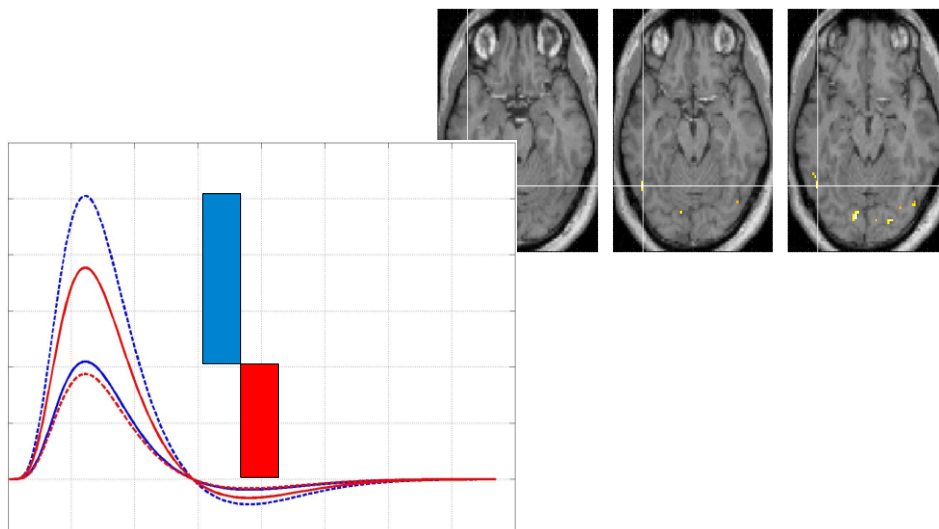
- SPM design matrix



- Questions:
- Main effects
- Fame: $[1 \ 1 \ -1 \ -1 \ 0 \ 0 \ \dots]$
- Rep: $[1 \ -1 \ 1 \ -1 \ 0 \ 0 \ \dots]$
- Interaction $[1 \ -1 \ -1 \ 1 \ 0 \ 0 \ \dots]$

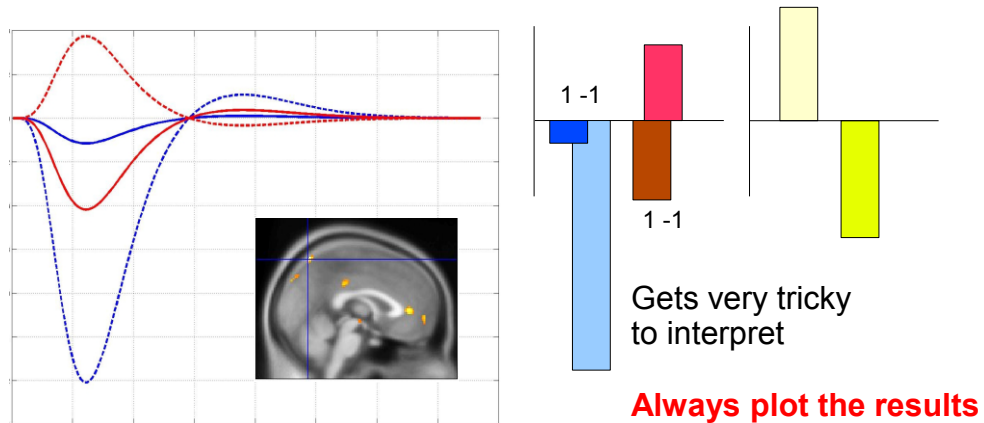


More contrasts



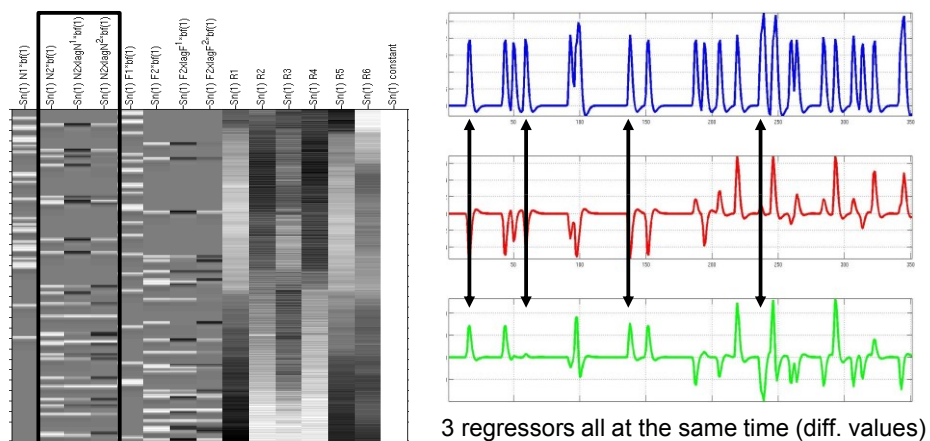
More contrasts

- Search in areas where all regressors are positive or all negative otherwise ...



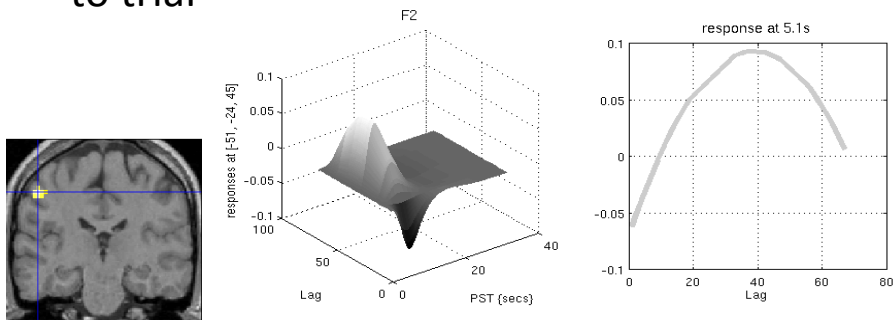
More Regressors

- Same design as before but added a 'parametric' regressor – here the lag between presentations



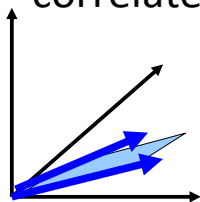
More Regressors

- The parametric regressors express the amplitude of signal as a function of the lag, i.e. the signal amplitude changes from trial to trial



More Regressors: collinearity

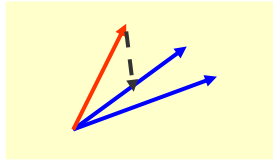
- SPM orthogonalizes the parametric regressors making the regressors non collinear.
- Three or more points are said to be collinear if they lie on a single straight line.
- Regressors are collinear if they are perfectly correlated (note $\text{corr of 2 vectors} = \cos\theta$)



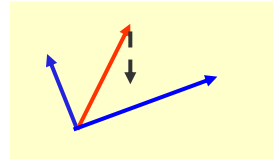
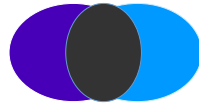
- Can make solution impossible
- Often make the model ok but individual regression values unstable
- Classical height and weight regression pblm

<http://en.wikipedia.org/wiki/Multicollinearity>
<http://mathworld.wolfram.com/Collinear.html>

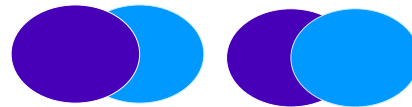
More Regressors: orthogonalization



Lot of variance shared –
because we look for the unique
part of variance, the shared
part goes into the error



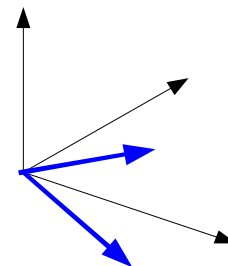
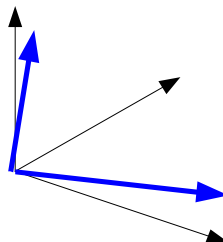
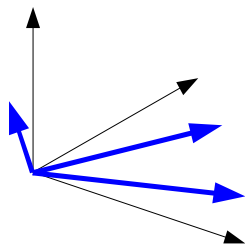
Orthogonalization ($\theta = 90^\circ$) removes shared
variance BUT order
matters !



http://en.wikipedia.org/wiki/Gram-Schmidt_process

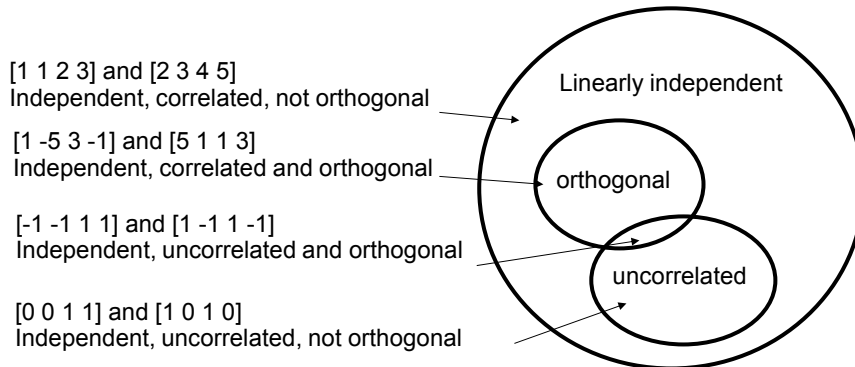
More regressors

- Linearly independent ($X_2 \neq aX_1$), orthogonal ($X_1'Y_2 = 0$)
and uncorrelated ($(X_1 - \text{mean}(X_1))'(X_2 - \text{mean}(X_2)) = 0$)
variables



More regressors

- Linearly independent ($X_2 \neq aX_1$), orthogonal ($X_1'Y_2 = 0$) and uncorrelated ($(X_1 - \text{mean}(X_1))'(X_2 - \text{mean}(X_2)) = 0$) variables



<http://www.jstor.org/pss/2683250>

Summary

- Linear model: $y = \beta_1 x_1 + \beta_2 x_2$ (output = additivity and scaling of input)
- GLM: $Y = XB + E$ (matrix formulation, works for any statistics, express the data Y as a function of the design matrix X)
- Contrasts: F or t test for the effect of 1 or several regressors given the design matrix
- More regressor is better as it captures more of the signal but it may bring instability if regressors are collinear (and cost df) – SPM orthogonalizes parametric regressors