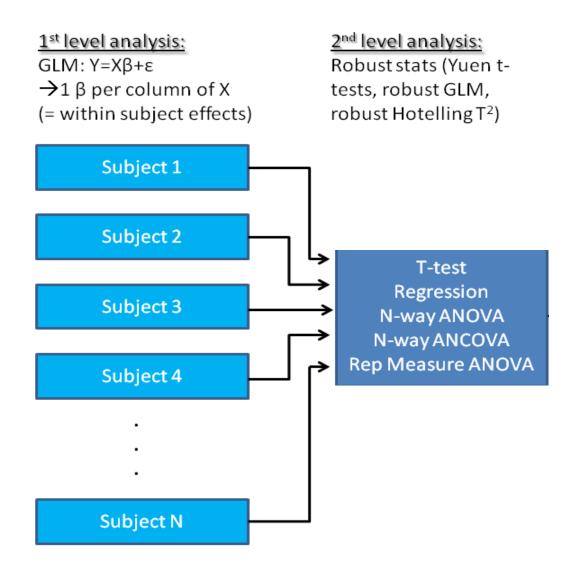
## Robust Linear Modelling of EEG data: the LIMO EEG plug-in

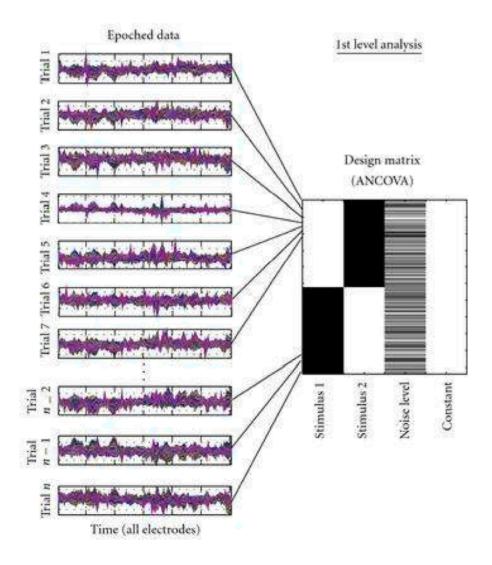
**Arnaud Delorme** 

with Cyril Pernet, PhD, Edinburgh Imaging & Centre for Clinical Brain Sciences

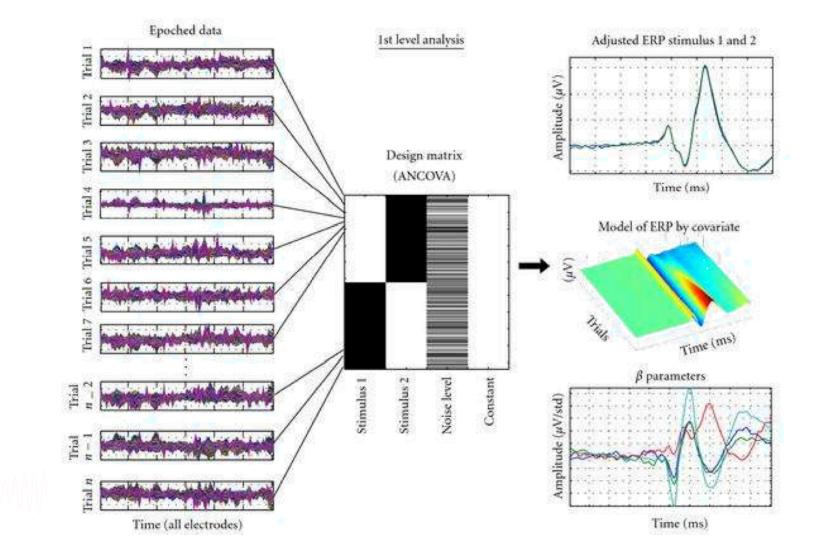
## **Hierarchical Linear Model**



## Linear Modeling of EEG data



## Linear Modeling of EEG data

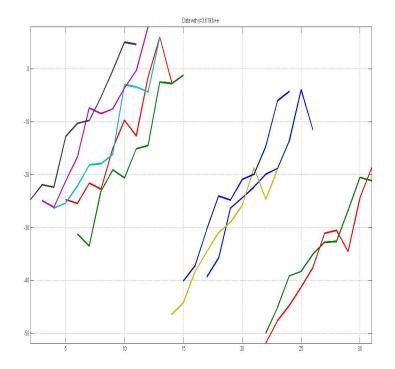


## Random Effect Model

Model the data with fixed effects (the experimental conditions) and a random effect (subjects are allowed to have different overall values – considering subjects as a random variable)

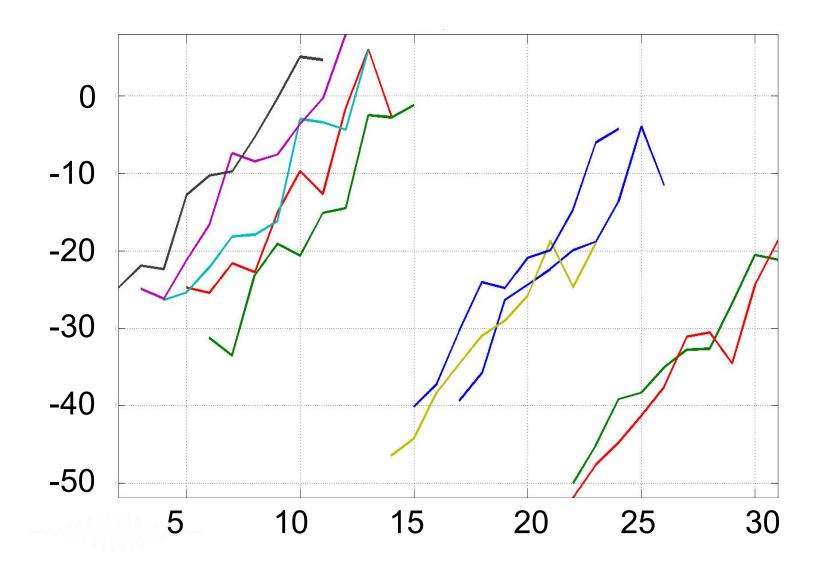
Example: present stimuli from intensity -5 units to +5 units around the subject perceptual threshold and measure RT

 $\rightarrow$  Plot the data per intensity

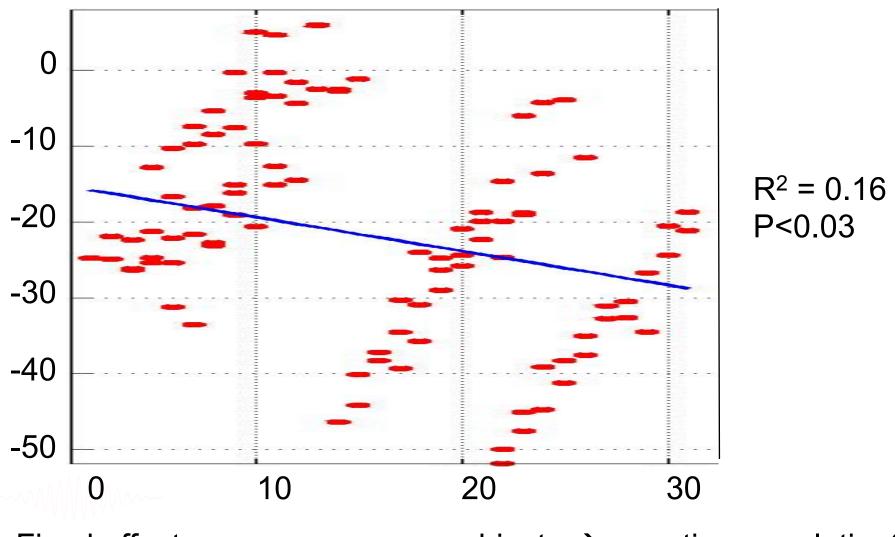




### **Fixed Effect**

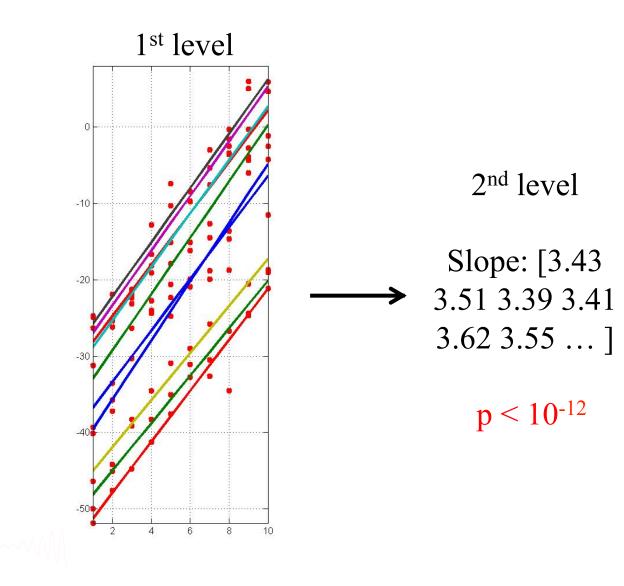


## **Fixed Effect**



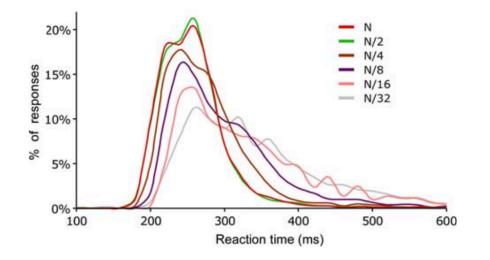
Fixed effect = average across subjects  $\rightarrow$  negative correlation?

## Random Effect Model



#### Varying factor: Contrast of image

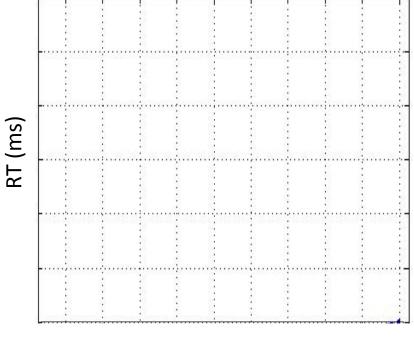
Outcome: Reaction time





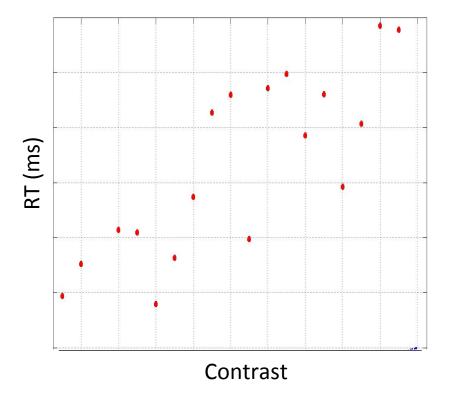
Mace, M., Delorme, A., Richard, G., Fabre-Thorpe, M. (2010) Spotting animals in natural scenes: efficiency of humans and monkeys at very low contrasts. *Animal Cognition*, 13(3):405-18.

• We have an experimental measure x (e.g. contrast)

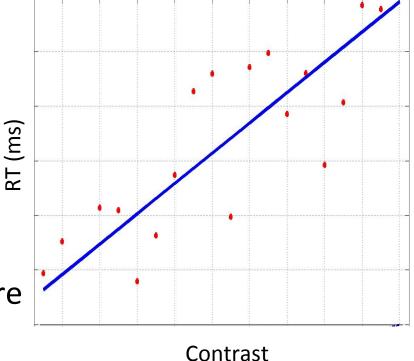


Contrast

- We have an experimental measure x (e.g. contrast)
- We then do the expe and collect data RT (e.g. reaction time)



- We have an experimental measure x (e.g. contrast)
- We then do the expe and collect data RT (e.g. reaction time)
- Model:  $\mathbf{RT} = \beta_0 + \mathbf{x}\beta_1 + \epsilon$
- Do some maths / run a software to find  $\beta_1$  and  $\beta_0$



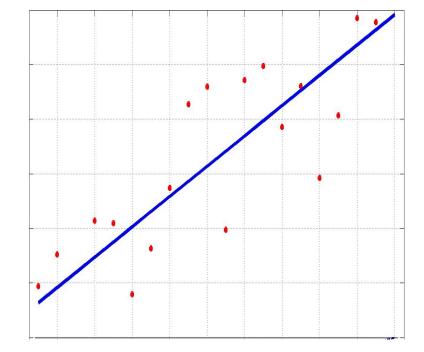
•  $RT^{*} = 23.6 + 2.7x$ 

### For each trial

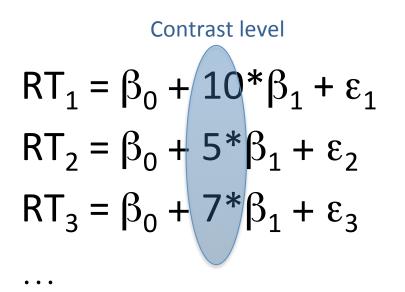
$$RT_{1} = \beta_{0} + 10^{*}\beta_{1} + \varepsilon_{1}$$

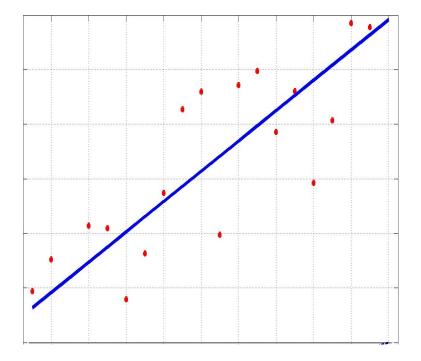
$$RT_{2} = \beta_{0} + 5^{*}\beta_{1} + \varepsilon_{2}$$

$$RT_{3} = \beta_{0} + 7^{*}\beta_{1} + \varepsilon_{3}$$



### For each trial





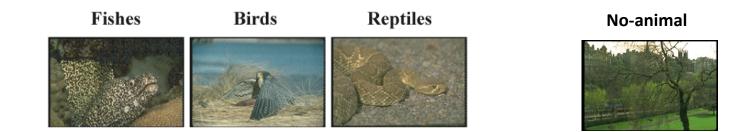
To test for significance compare the original regression model  $RT_i = \beta_0 + c_i^*\beta_1 + \varepsilon_i$  with the simplified model  $RT_i = \beta_0 + \varepsilon_i$ 

Compare these errors

## An ANOVA is a linear model

Varying factor: Type of image

**Outcome:** Reaction time (go/no-go)



**Delorme, A.**, Richard, G., Fabre-Thorpe, M. (2010). Key visual features for rapid categorization of animals in natural scenes. *Frontier in psychology*, 1:21

$$\mathsf{RT}_{i,j} = \beta_0 + \beta_i + \varepsilon_{i,j}$$

$$\mathsf{RT}_{i,j} = \beta_0 + \beta_i + \varepsilon_{i,j}$$

2

For trial 4 (for example first trial of birds) we have

 $\mathsf{RT}_{2,1} = \beta_0 + 0^*\beta_1 + 1^*\beta_2 + 0^*\beta_3 + \varepsilon_{2,1}$ 

$$\mathsf{RT}_{i,j} = \beta_0 + \beta_i + \varepsilon_{i,j}$$

For trial 4 (for example first trial of birds) we have

 $\mathsf{RT}_{2,1} = \beta_0 + 0^*\beta_1 + 1^*\beta_2 + 0^*\beta_3 + \varepsilon_{2,1}$ 

For trial 13 (for example second trial of birds) we have

 $\mathsf{RT}_{2,2} = \beta_0 + 0^* \beta_1 + 1^* \beta_2 + 0^* \beta_3 + \varepsilon_{2,2}$ 

)

$$\mathsf{RT}_{\mathsf{i},\mathsf{j}} = \beta_0 + \beta_\mathsf{i} + \varepsilon_{\mathsf{i},\mathsf{j}}$$

For trial 4 (for example first trial of birds) we have

 $\mathsf{RT}_{2,1} = \beta_0 + 0^*\beta_1 + 1^*\beta_2 + 0^*\beta_3 + \varepsilon_{2,1}$ 

For trial 13 (for example second trial of birds) we have

 $\mathsf{RT}_{2,2} = \beta_0 + 0^*\beta_1 + 1^*\beta_2 + 0^*\beta_3 + \varepsilon_{2,2}$ 

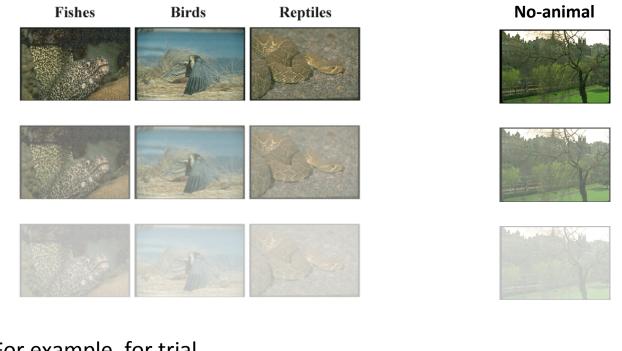
Statistics: if there is an effect of treatment then error of the simplified model  $RT_{i,j} = \beta_0 + \varepsilon_{i,j}$  should be lower than the original model  $RT_{i,j} = \beta_0 + \beta_i + \varepsilon_{i,j}$ 

#### Compare these errors

This is a GLM that is also strictly equivalent to running an ANOVA

## The GLM can do both a Regression and an ANOVA

#### Varying factor: Type of image AND contrast Outcome: Reaction time (go/no-go)

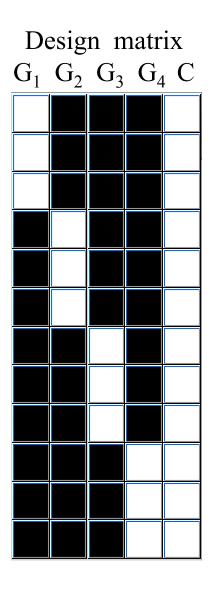


For example, for trial (first bird with contrast  $c_{2,1}$ ) we have  $RT_{2,1} = \beta_0 + \underbrace{0*\beta_1 + 1*\beta_2 + 0*\beta_3 + 0*\beta_2}_{Categorical var.} + \underbrace{c_{2,1}*\beta_4 + \varepsilon_{2,1}}_{REGRESSION}$ 

## The design matrix

. . .

Y	Gp
8	1
9	1
7	1
5	2
7	2
3	2
3	3
4	3
1	3
6	4
4	4
9	4



 $y(1..3) = 1x\beta1 + 0x\beta2 + 0x\beta3 + 0x\beta4 + c + error$  $y(4..6) = 0x\beta1 + 1x\beta2 + 0x\beta3 + 0x\beta4 + c + error$  $y(7..9) = 0x\beta1 + 0x\beta2 + 1x\beta3 + 0x\beta4 + c + error$  $y(10..12) = 0x\beta1 + 0x\beta2 + 0x\beta3 + 1x\beta4 + c + error$ 

$$\left\{\begin{array}{c}
8\\9\\7\\5\\7\\3\\4\\1\\6\\4\\9\end{array}\right\} = \left\{\begin{array}{c}
10001\\10001\\01001\\01001\\00101\\00101\\00101\\00011\\00011\\00011\\00011\end{array}\right\} * \left\{\begin{array}{c}
\beta_1\\\beta_2\\\beta_3\\\beta_4\\c\end{array}\right\} + \left\{\begin{array}{c}
e^1\\\beta_2\\\beta_3\\\beta_4\\c\end{array}\right\}$$

### Design considerations

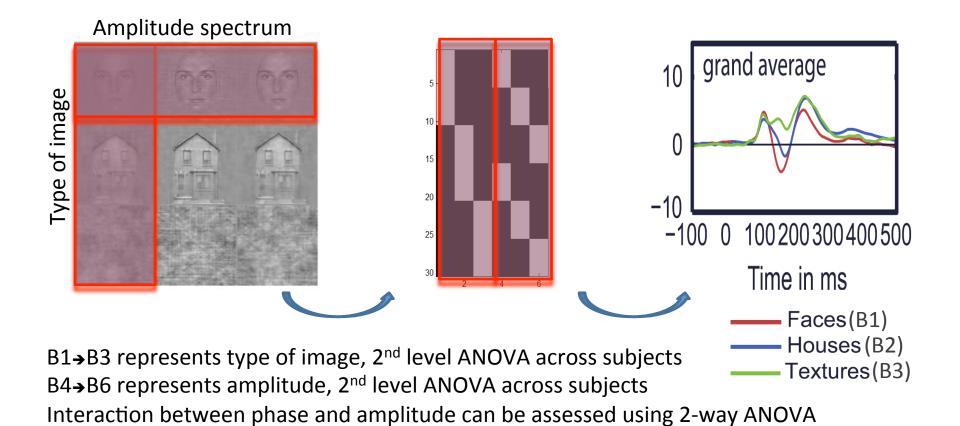
Illustration with a set of studies looking at the effect of stimulus amplitude and phase information



Rousselet, Pernet, Bennet, Sekuler (2008). Face phase processing. BMC Neuroscience 9:98

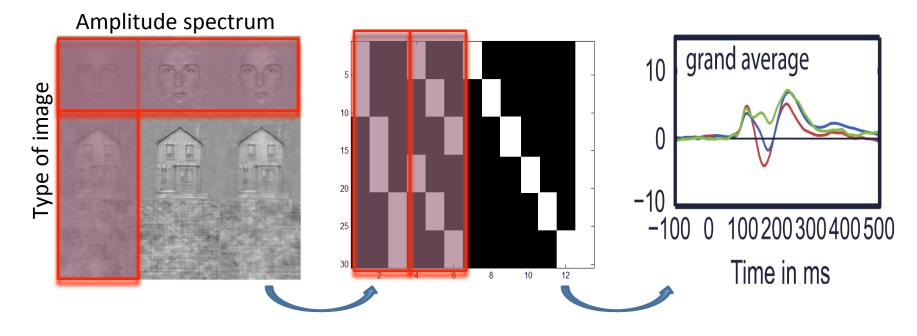
## Factorial Designs: N\*N\*N\*...

Categorical designs: Group level analyses of course but also Individual analyses with bootstrap



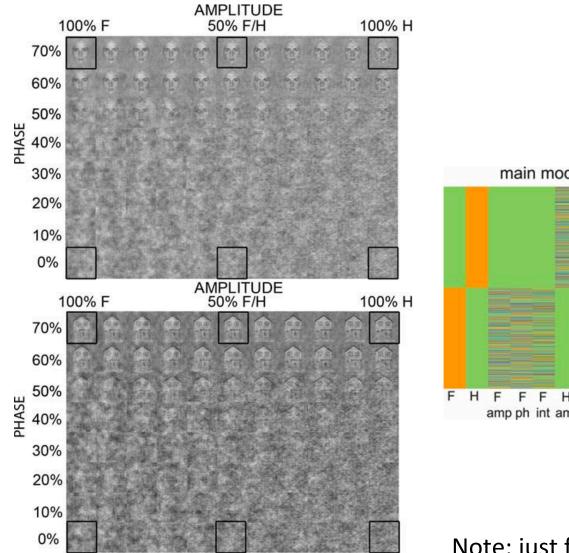
## Factorial Designs: N\*N\*N\*...

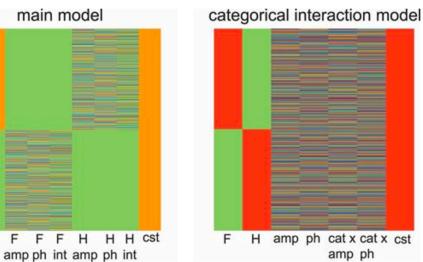
## Categorical designs: Group level analyses of course but also Individual analyses with bootstrap



Interaction between phase and amplitude can be assessed using 1-way ANOVA on B7 to B12. There is no interaction left between B1-B3 and B4-B6 <u>Bienek, Pernet, Rousselet (2012). Phase vs Amplitude Spectrum. Journal of Vision 12(13), 1–24</u>

### **Continuous designs**

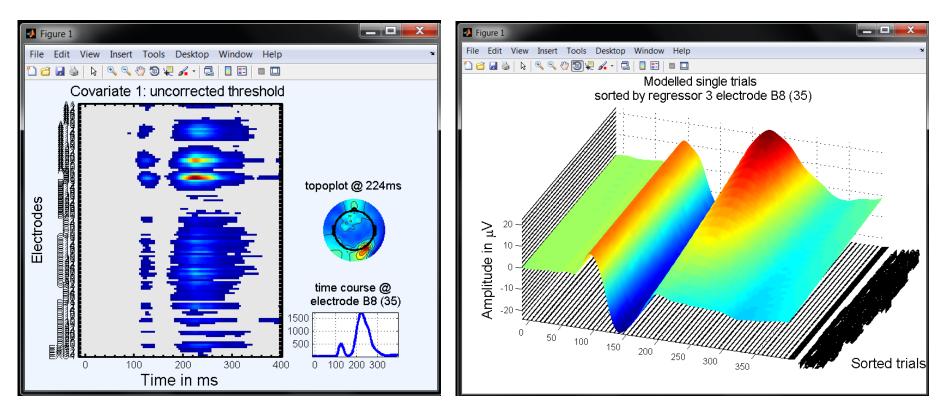




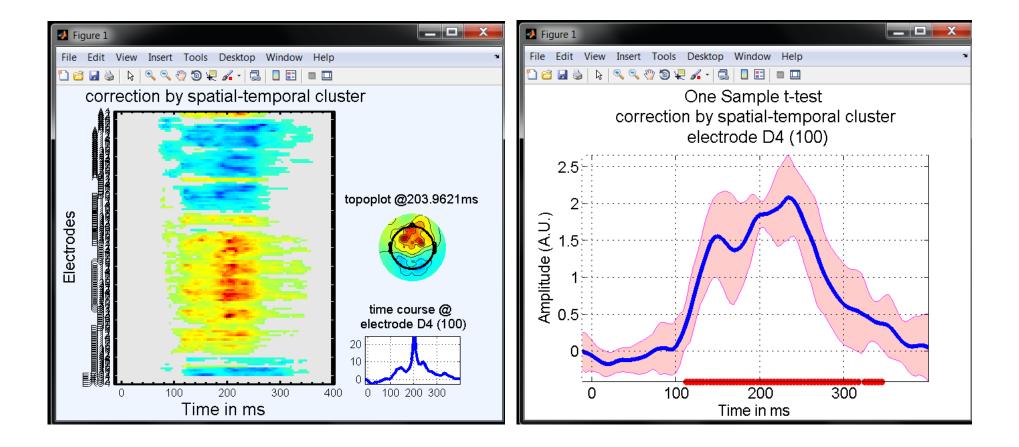
Note: just face and houses (no noise here)

## What have we done: results

- Image all (R2, condition, covariate)
- Course plots, topoplots



## **Review group level results**



## **Design questions!**

- Let's think how to analyse your data!
- Nb of conditions / covariates
- contrasts
- 1<sup>st</sup> level covariates
- 2<sup>nd</sup> level covariates