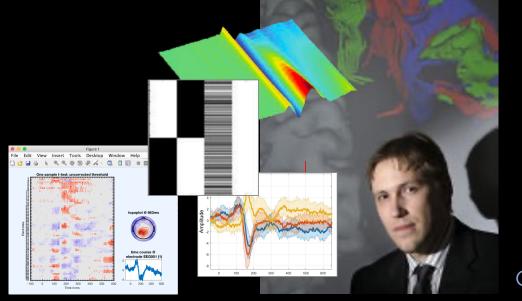
General Linear Modeling in EEGLAB/LIMO

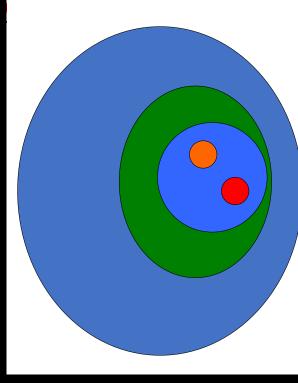
Basic Theory

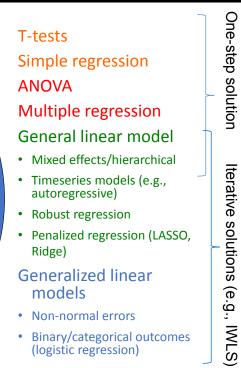
Arnaud Delorme



Cyril Pernet

The GLM family

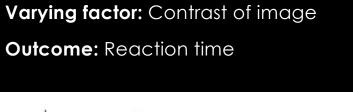


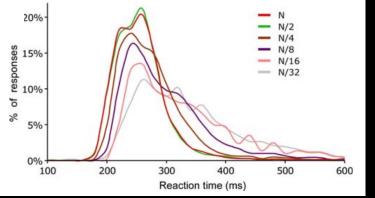


Iterative solutions e ġ.

Slide from Tor Wager

A regression is a linear model





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.

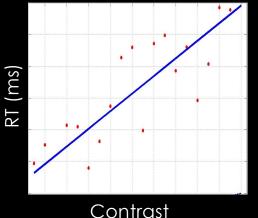
A regression is a linear model

► Given an experimental measure x (e.g. contrast)

► We then do the experiment and collect data RT (e.g. reaction time)

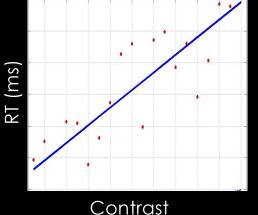
• Model: $RT = \beta_0 + x\beta_1 + \varepsilon$

► Do some maths / run a software to find β_1 and β_0 ► RT = 23.6 + 2.7x



A regression is a linear model

For each trial $RT_{1} = \beta_{0} + 10^{*}\beta_{1} + \varepsilon_{1}$ $RT_{2} = \beta_{0} + 5^{*}\beta_{1} + \varepsilon_{2}$ $RT_{3} = \beta_{0} + 7^{*}\beta_{1} + \varepsilon_{3}$



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 the fit Test if 0 included in confidence interval

An ANOVA is a linear model

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





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

that is to say the data (e.g. RT) = a constant term (grand mean β_0) + the effect of a treatment (β_1 for fishes 1 and β_2 , β_3 for birds and reptiles) and the error term ($\epsilon_{i,i}$)

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

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

This is a GLM that is equivalent to an ANOVA

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

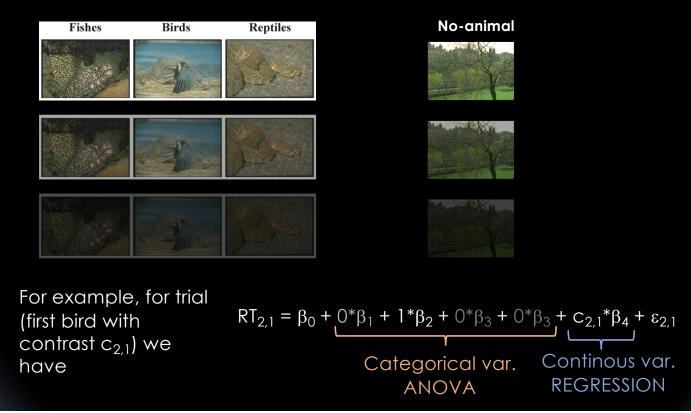
 $\mathsf{RT}_{2,2} = \beta_0 + 0^* \beta_1 + 1^* \beta_2 + 0^* \beta_3 + \varepsilon_{2,2}$

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

Compare the fit

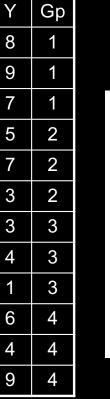
A GLM can do both a Regression and an ANOVA (ANCOVA)

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



The design matrix

 $y(1..3) = 1x\beta 1 + 0x\beta 2 + 0x\beta 3 + 0x\beta 4 + c + error$ $y(4..6) = 0x\beta 1 + 1x\beta 2 + 0x\beta 3 + 0x\beta 4 + c + error$ $y(7..9) = 0x\beta 1 + 0x\beta 2 + 1x\beta 3 + 0x\beta 4 + c + error$ $y(10..12) = 0x\beta 1 + 0x\beta 2 + 0x\beta 3 + 1x\beta 4 + c + error$ Design matrix $G_1 G_2 G_3 G_4 C$ é1) 8 9 7



5

7 3

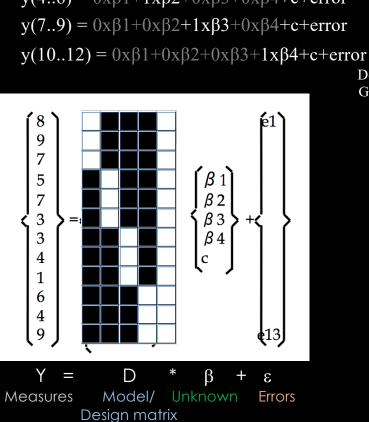
3

4

1

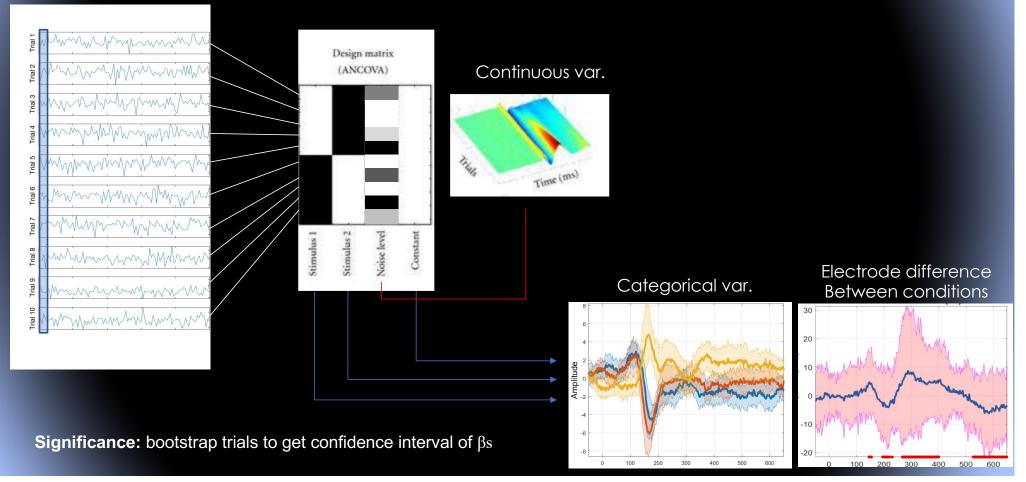
6

4 9

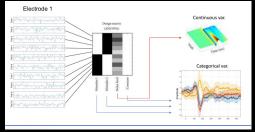




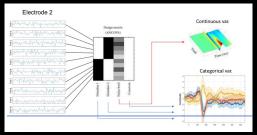
Electrode 1



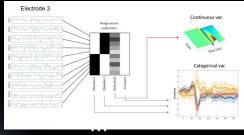
Electrode 1



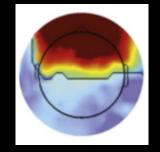
Electrode 2



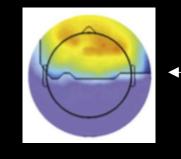
Electrode 3



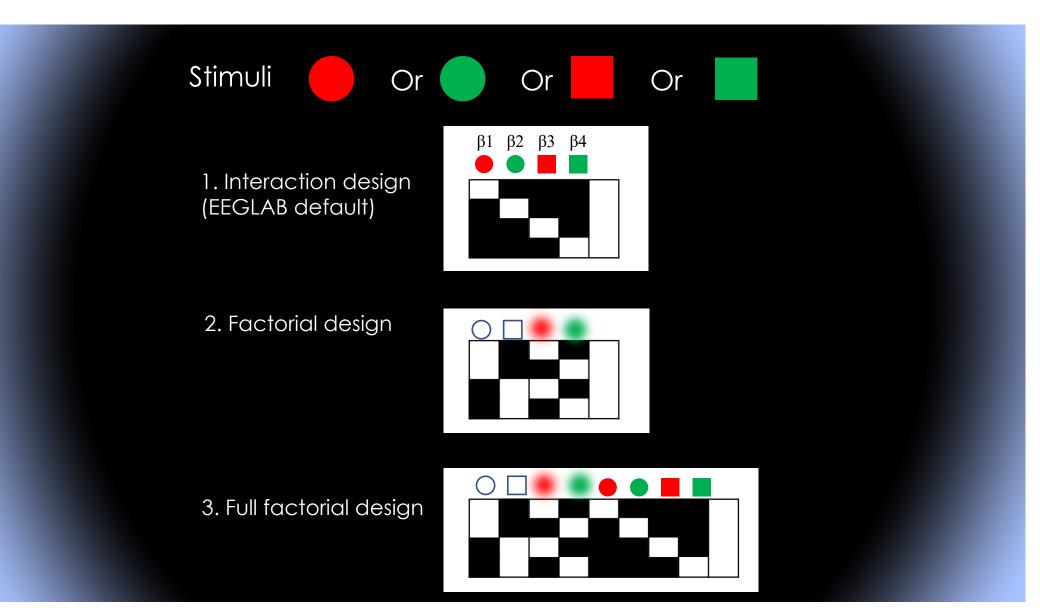
Scalp topography of **beta difference** at a given latency

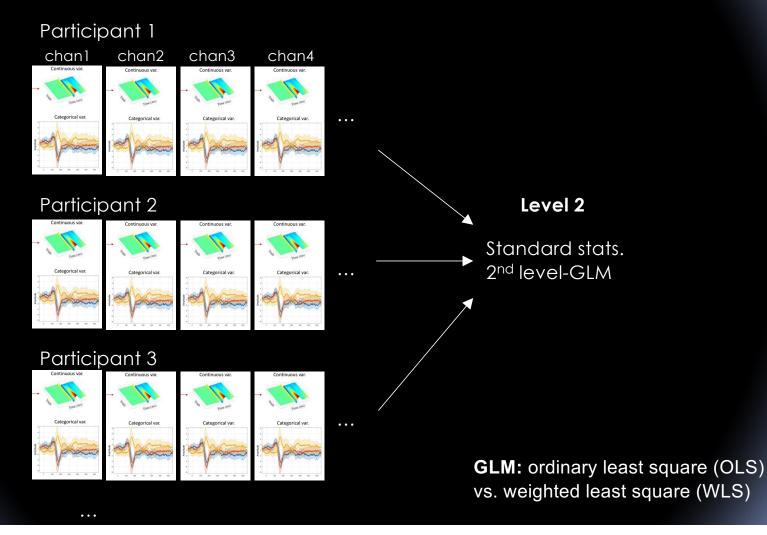


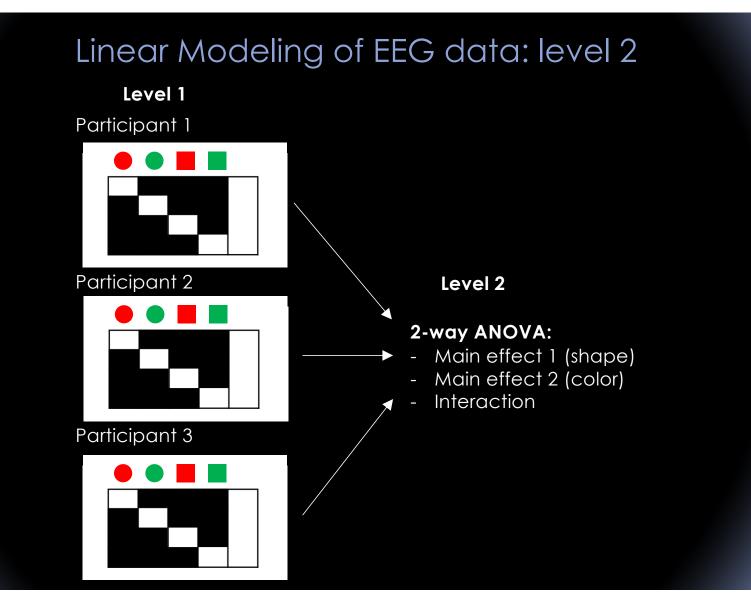
Scalp topography of **potential difference** (masked using beta signif.)



Limit of the regions



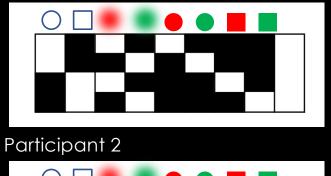




X

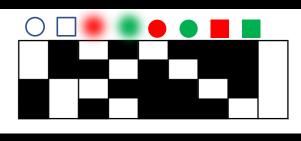
 \geq

Participant 1





Participant 3



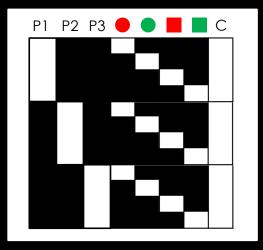
Shape: Group t-test on βο− β□

Color Group t-test on $\beta * - \beta *$

Interaction - One sample t-test on

β●, **β●**, **β■**, **β**■

Mixed effect model (still a GLM)



Hierarchical GLM

