

Neuroimaging for Machine Learners

Validation and inference

PRoNTo course
June 2017

Christophe Phillips, Jr. PhD.

Univariate analysis:

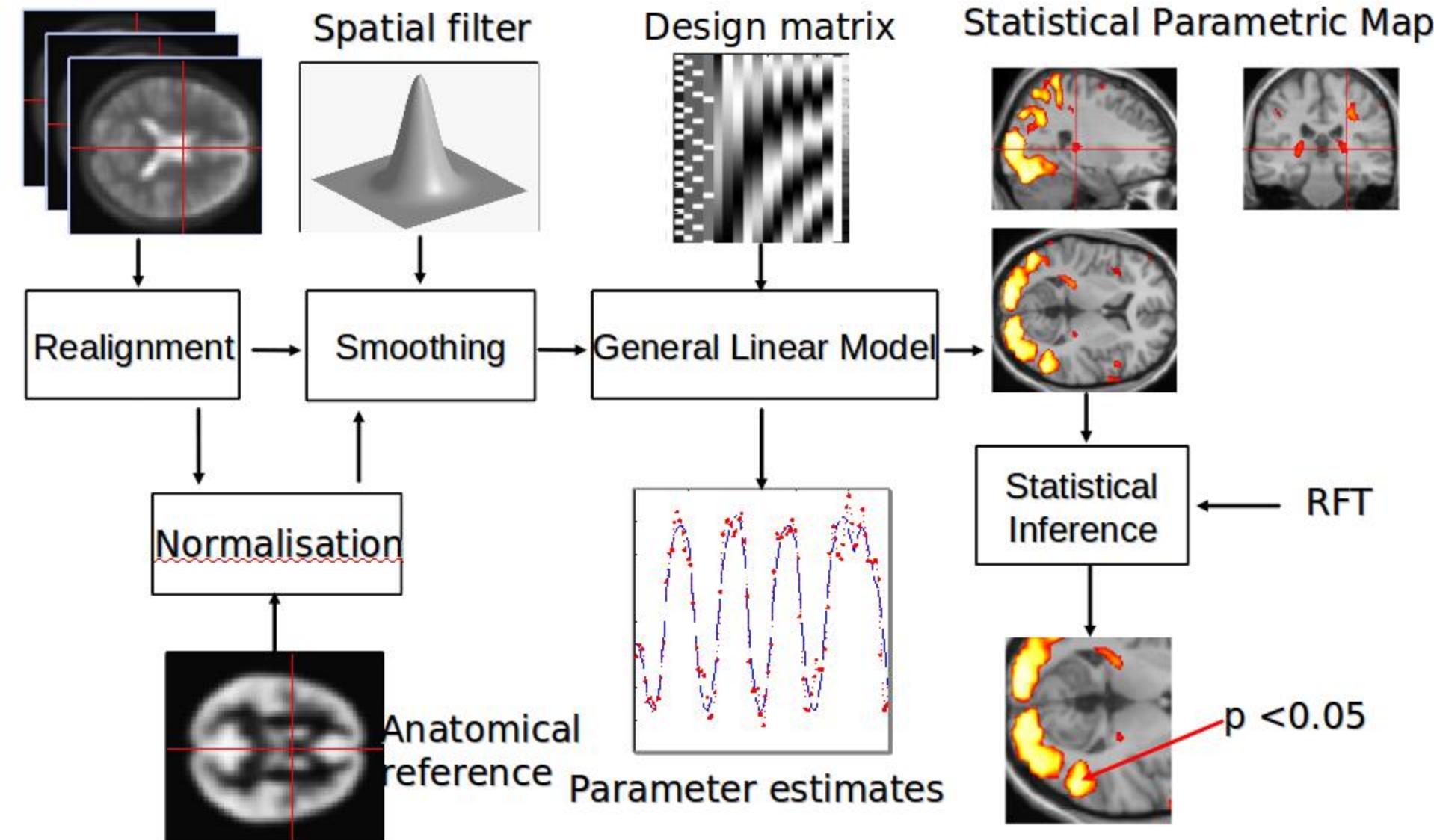
- Introduction: within & between subjects analysis
- GLM, contrasts & Statistical inference
- Multiple comparison problem & other inference levels
- Conclusion

Univariate analysis:

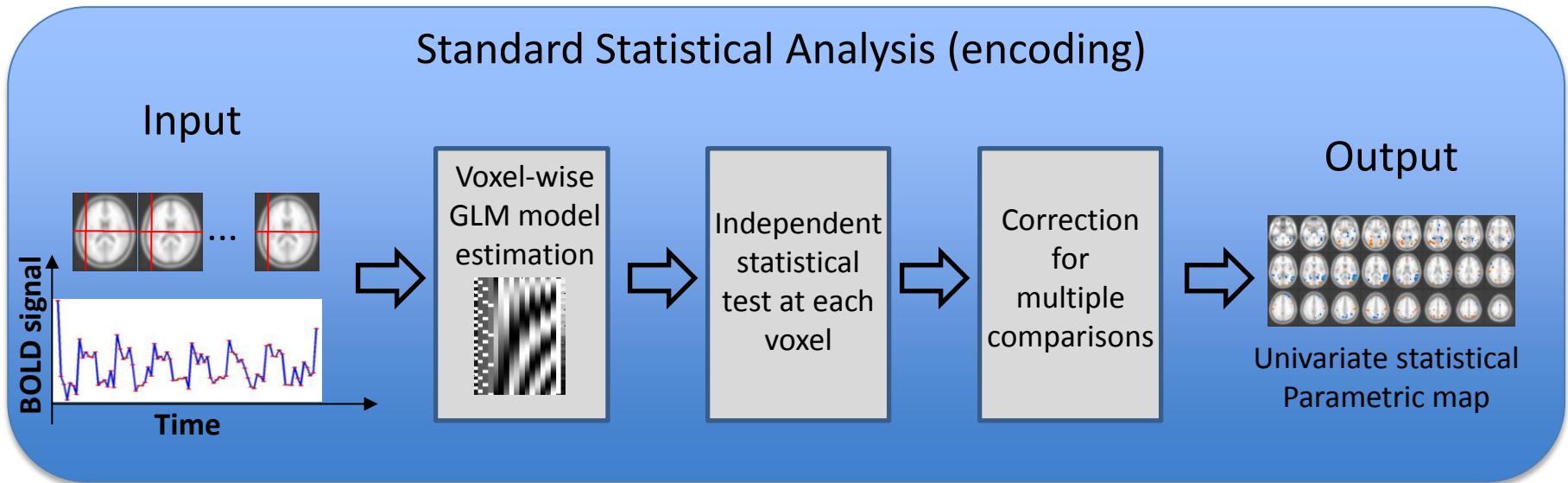
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Image time-series



Standard univariate approach



Find the mapping g from explanatory variable X to observed data Y

$$g: X \rightarrow Y$$

Within subject analysis :

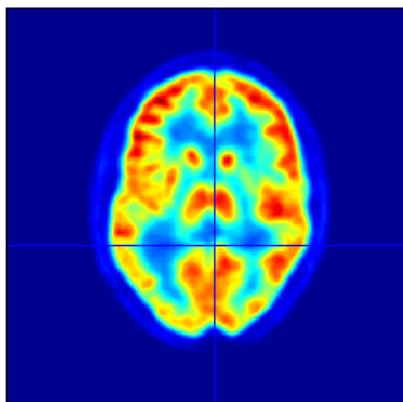
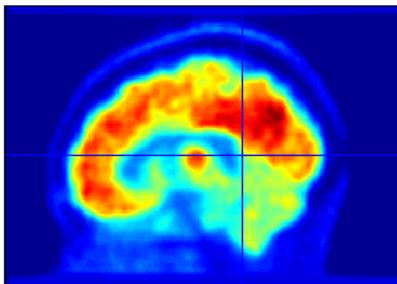
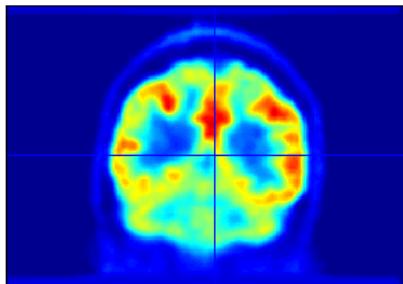
- « first level » analysis
- data = series of measurements from 1 subject, typically « functional MRI » (fMRI)
- modelling BOLD response time series
 - activation task or resting connectivity
- sometimes with parametric modulation and/or confounding regressor
- output = contrast images, statistical maps, connectivity maps, and « cleaned » signal

Between subjects analysis :

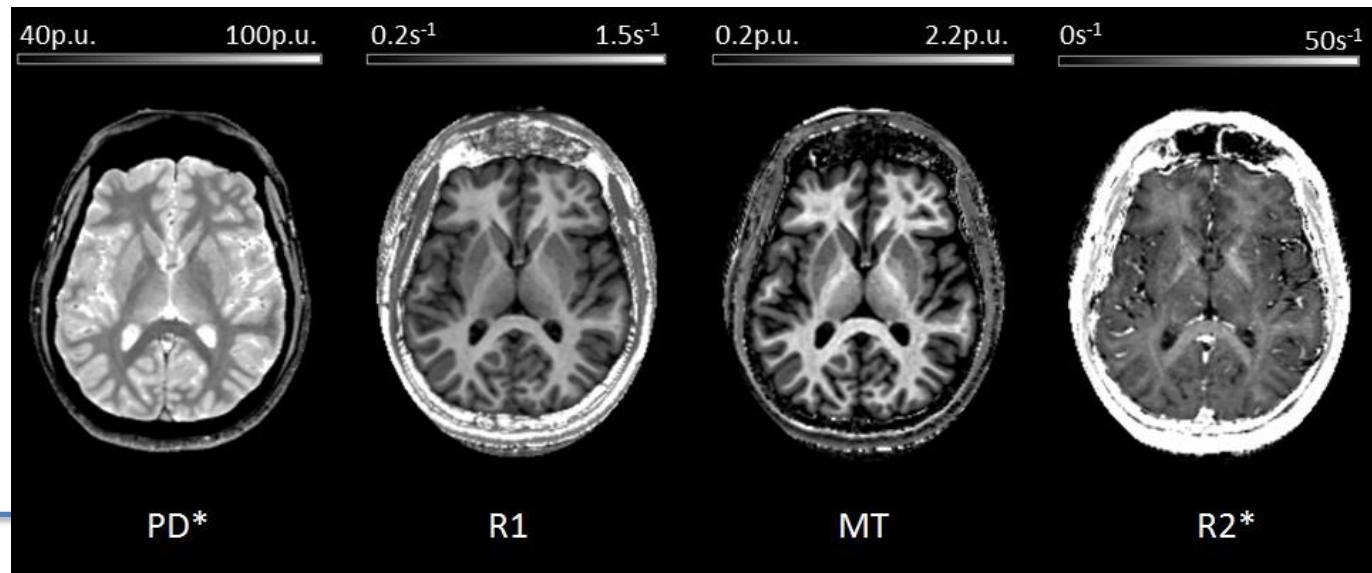
- « second level » analysis
- data = 1 (or few) image(s)/subject from group(s) of subjects, e.g. contrast image, tissue density, deformation map, FDG-PET scan,...
- modelling group effect/differences or regressing subjects' score (confound/interest)
- output = « raw » signal, contrast images and statistical maps

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FDG-PET scan



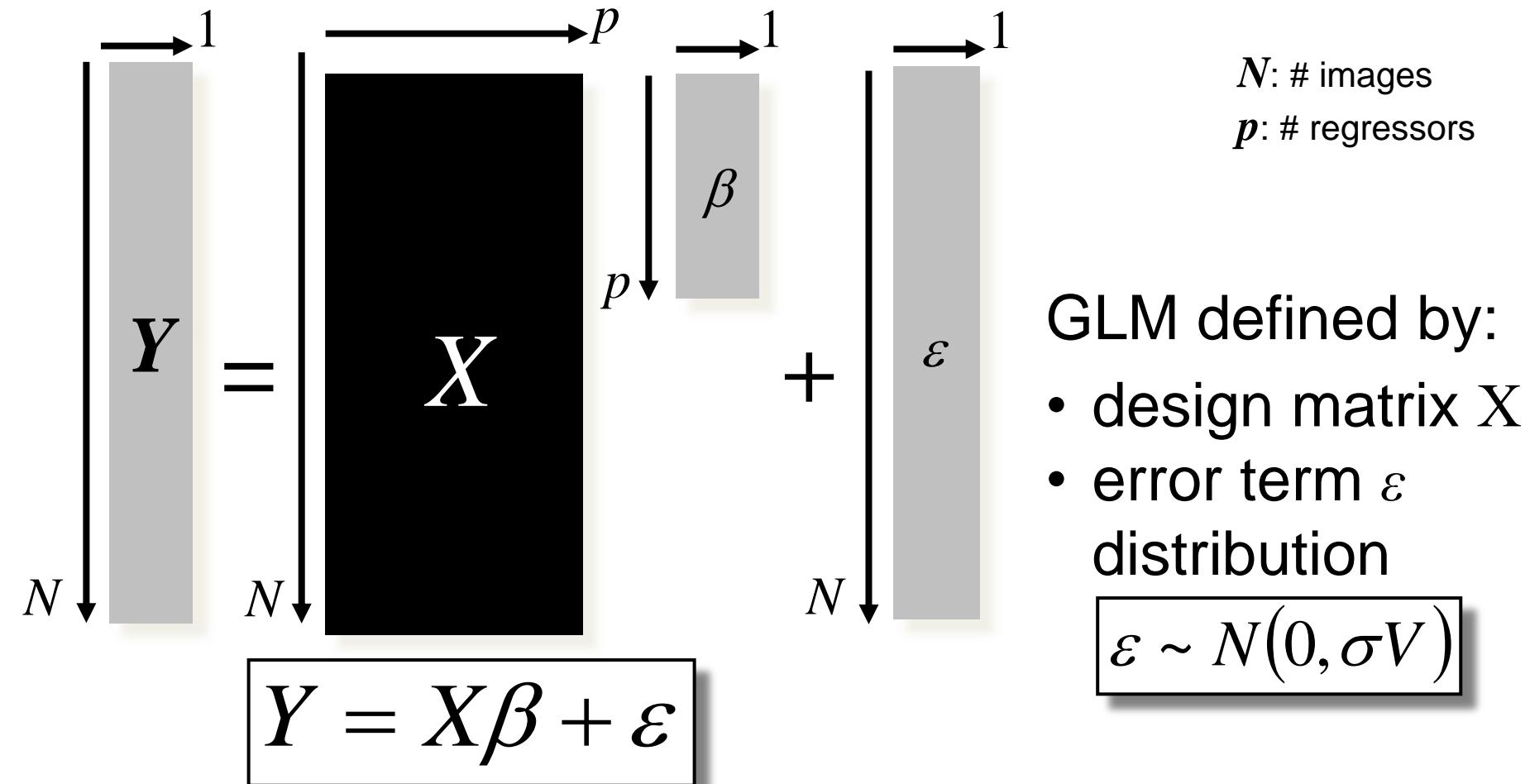
Quantitative MRI



Univariate analysis:

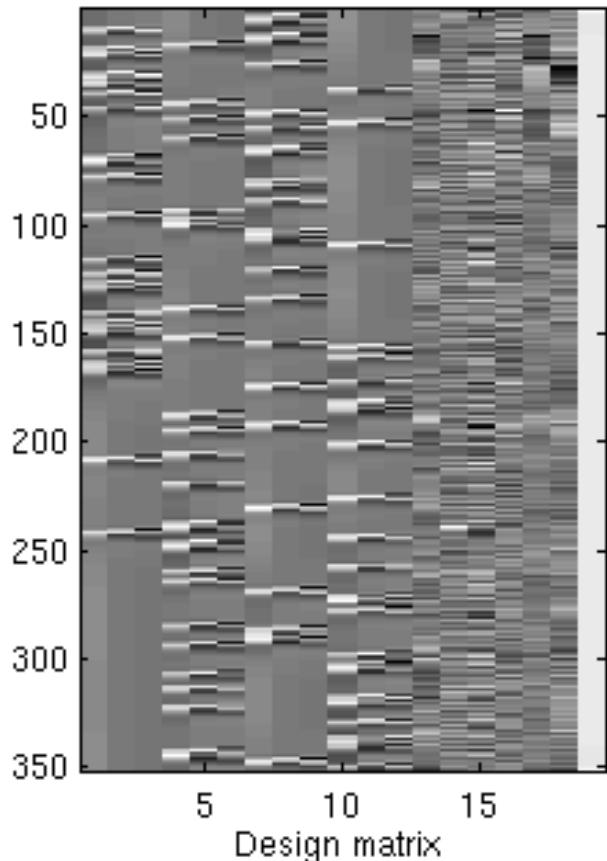
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General Linear Model



Design matrix example

Single subject
fMRI time series



Aim:

model BOLD signal variation
over time/scans, with

- 4 conditions + movement parameters + mean
- each condition event modelled by HRF with 2 derivatives (\approx Taylor expansion)

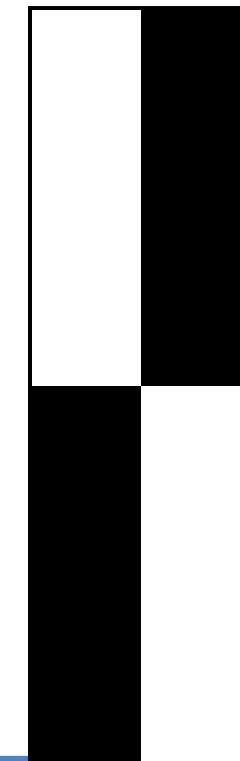
Design matrix example

Group comparison
with two-sample t-test

Aim:

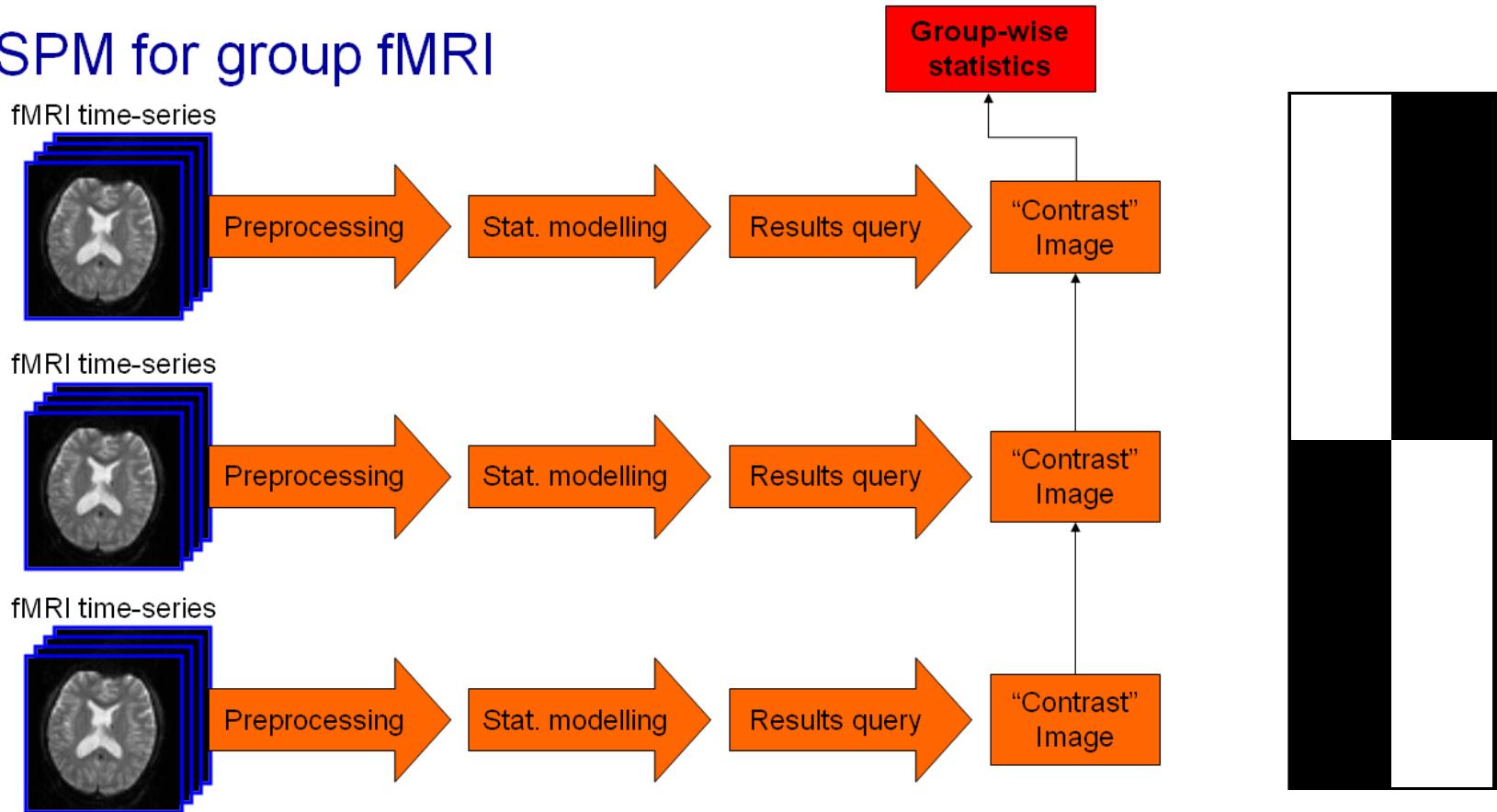
model “group effect“ with:

- Separate group averages



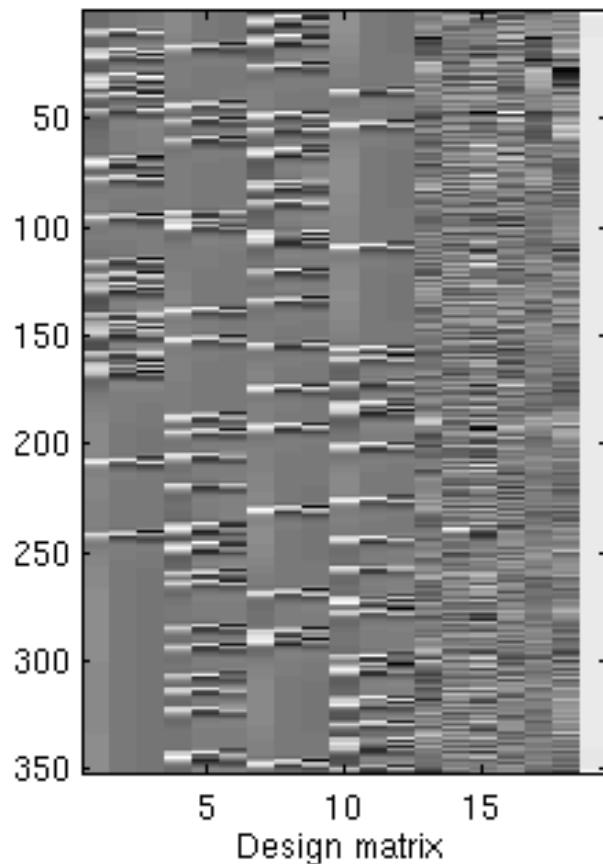
Design matrix example

SPM for group fMRI

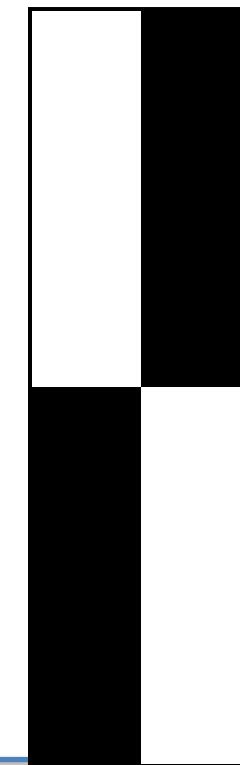


Design matrix example

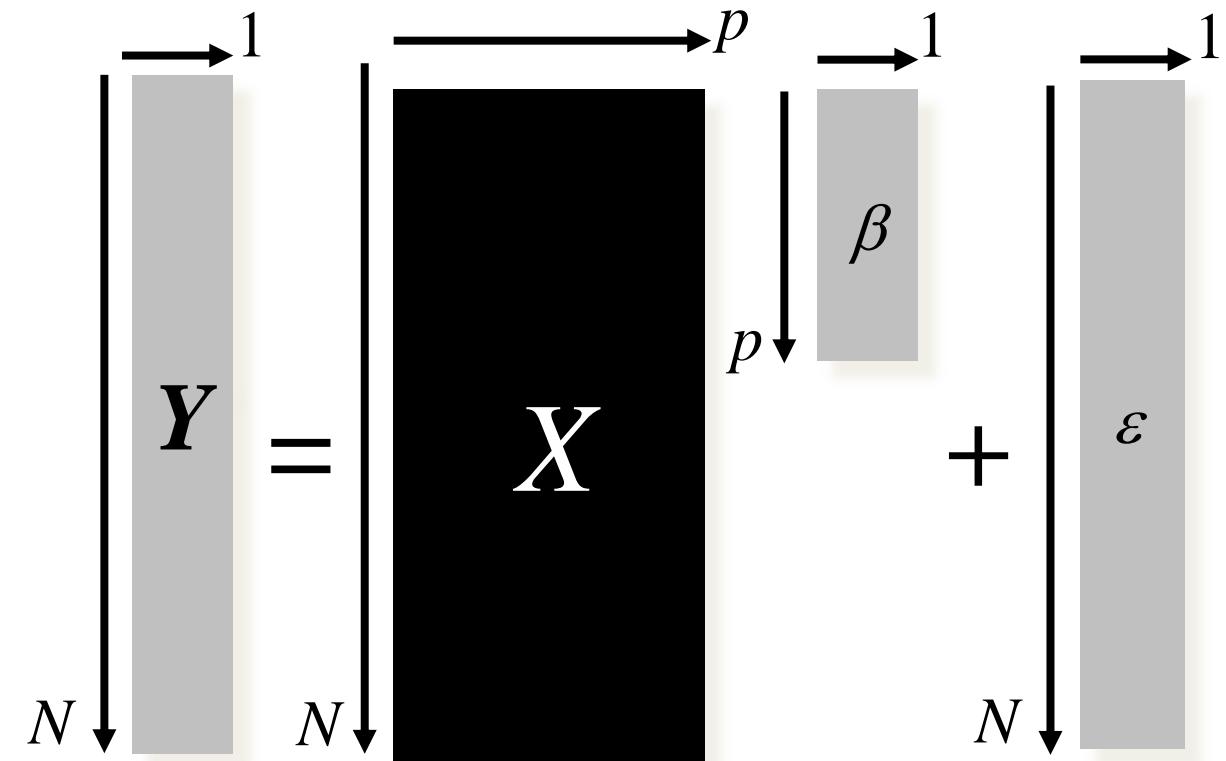
Single subject
fMRI time series



Group comparison
with two-sample t-test



General Linear Model



NOTE:

- GLM applied to each and every voxel *individually!*
- Estimation of the β 's in a (weighted) least-square sense.

$$Y = X\beta + \varepsilon$$

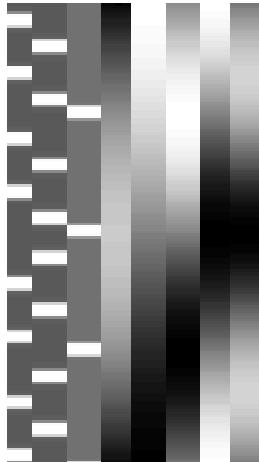
$$\varepsilon \sim N(0, \sigma V)$$

T-test & contrast

A contrast selects a specific effect of interest:

⇒ contrast c = vector of length p .

⇒ $c^T \beta$ = linear combination of regression coefficients β .



$$c^T = [1 \ 0 \ 0 \ 0 \ 0 \ \dots]$$

$$c^T \beta = 1 \times \beta_1 + 0 \times \beta_2 + 0 \times \beta_3 + 0 \times \beta_4 + 0 \times \beta_5 + \dots$$

$$c^T = [0 \ -1 \ 1 \ 0 \ 0 \ \dots]$$

$$c^T \beta = 0 \times \beta_1 + -1 \times \beta_2 + 1 \times \beta_3 + 0 \times \beta_4 + 0 \times \beta_5 + \dots$$

Under i.i.d assumptions $\varepsilon \sim N(0, \sigma I)$

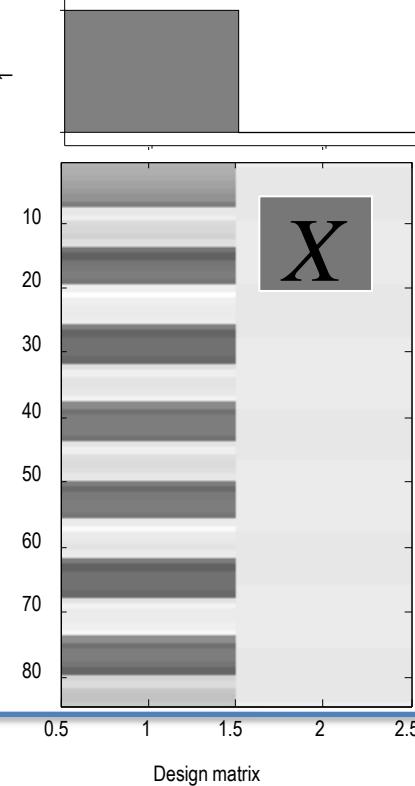
$$c^T \hat{\beta} \sim N(c^T \beta, \sigma^2 c^T (X^T X)^{-1} c)$$

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t-test example:

Passive word listening versus rest

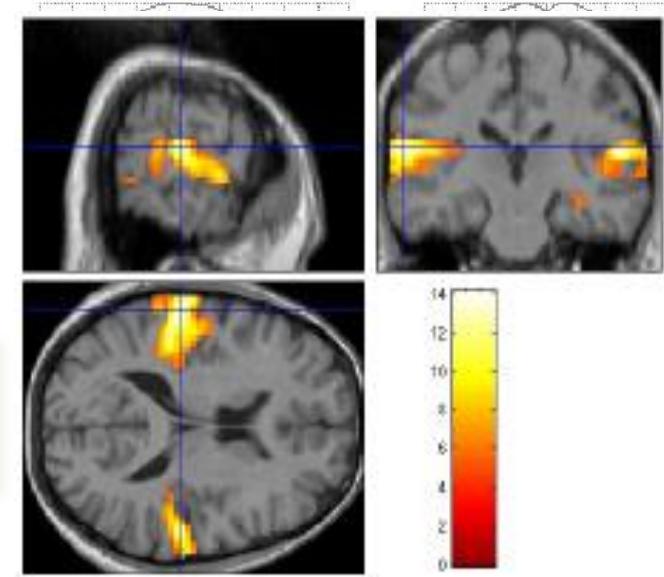
$$c^T = [1 \quad 0]$$



Q: activation during listening ?

Null hypothesis: $\beta_1 = 0$

$$t = \frac{c^T \hat{\beta}}{Std(c^T \hat{\beta})}$$



SPM results:

Height threshold T = 3.2057 {p<0.001}

voxel-level

		T	(Z)	p uncorrected	mm mm mm
Statistics:	p-values adjusted for search volume	13.94	11.82	0.000	-63 -27 15
set-level	cluster-level	Inf	Inf	0.000	-48 -33 12
p	c	p corrected	p EBA	p uncorrected	mm mm mm
0.000	10	0.000	13.72	0.000	57 0.021 12
			12.29	0.000	53 0.000 12
			9.26	0.000	51 0.000 12
			7.39	0.000	57 0.000 12
			6.84	0.000	51 0.000 12
			6.36	0.000	51 0.000 12
			6.19	0.000	51 0.000 12
			5.96	0.000	51 0.000 12
			5.84	0.000	51 0.000 12
			5.44	0.000	48 0.000 12
			5.44	0.000	48 0.000 12
			5.32	0.000	36 -27 17 42

Classical inference

The Null Hypothesis H_0

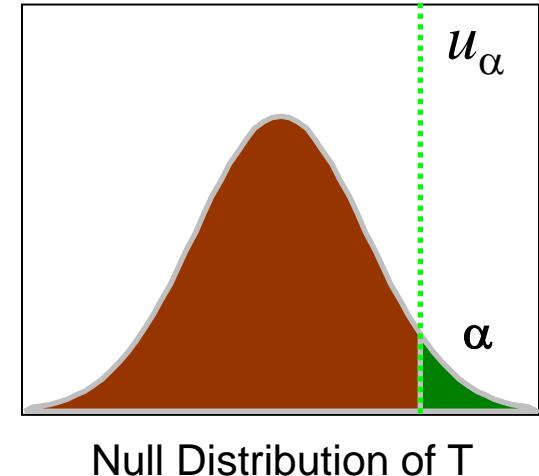
= what we want to disprove (no effect).

⇒ The Alternative Hypothesis H_A
 = outcome of interest.

Significance level α : $\alpha = p(T > u_\alpha \mid H_0)$

Acceptable *false positive rate* α .

⇒ threshold u_α



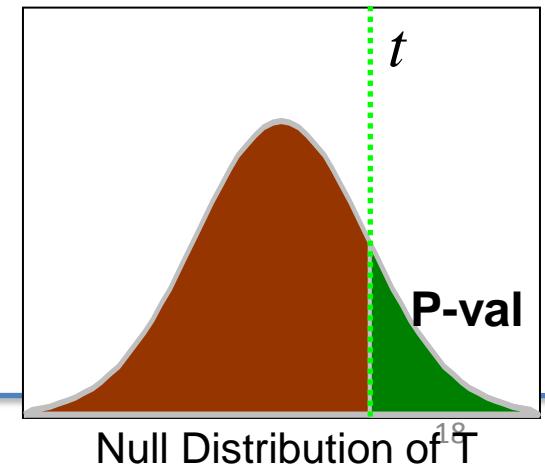
Observation of test statistic t

= a realisation of T

⇒ Reject H_0 in favour of H_A if $t > u_\alpha$

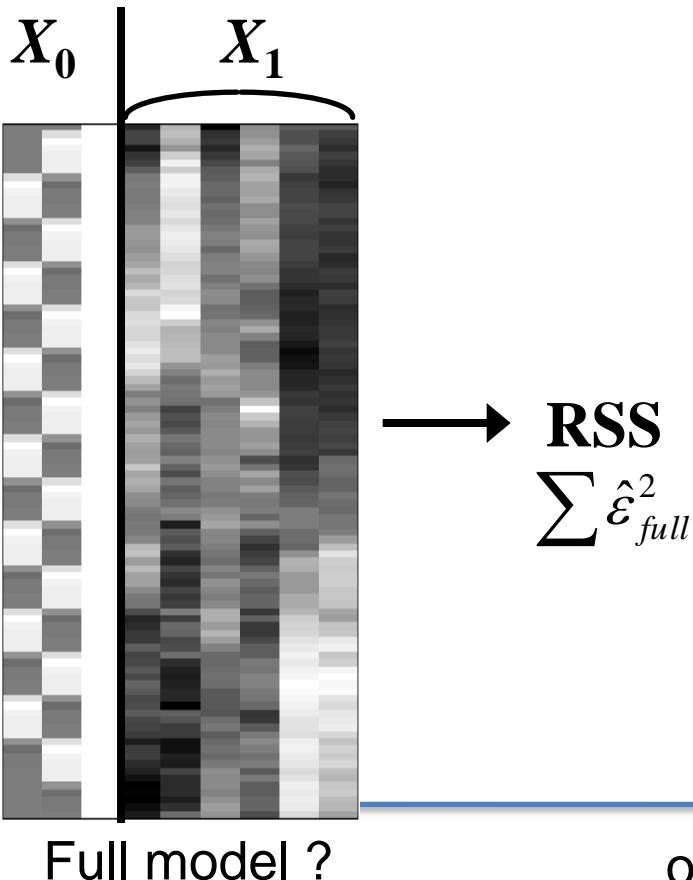
p-value = evidence against H_0 .

$$p(T > t \mid H_0)$$



F-test & contrast

Null Hypothesis H_0 : True model is X_0 (reduced model)



Test statistic: ratio of explained variability and unexplained variability (error)

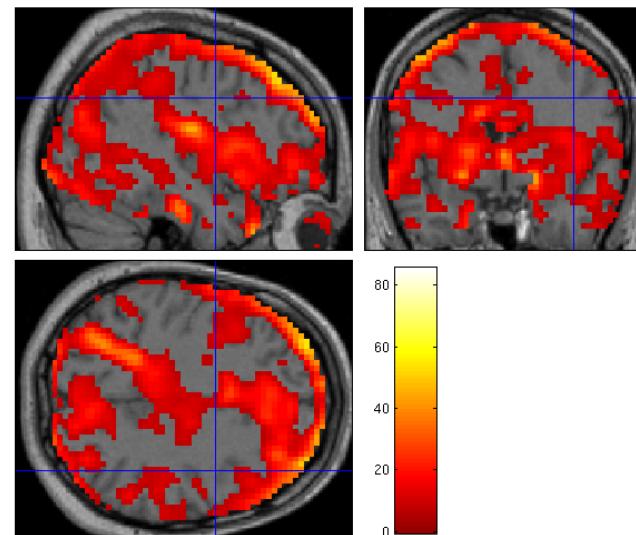
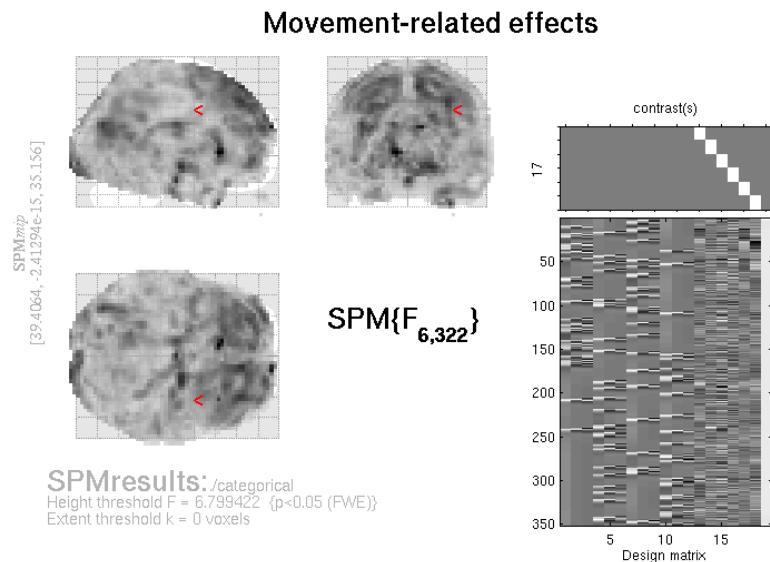
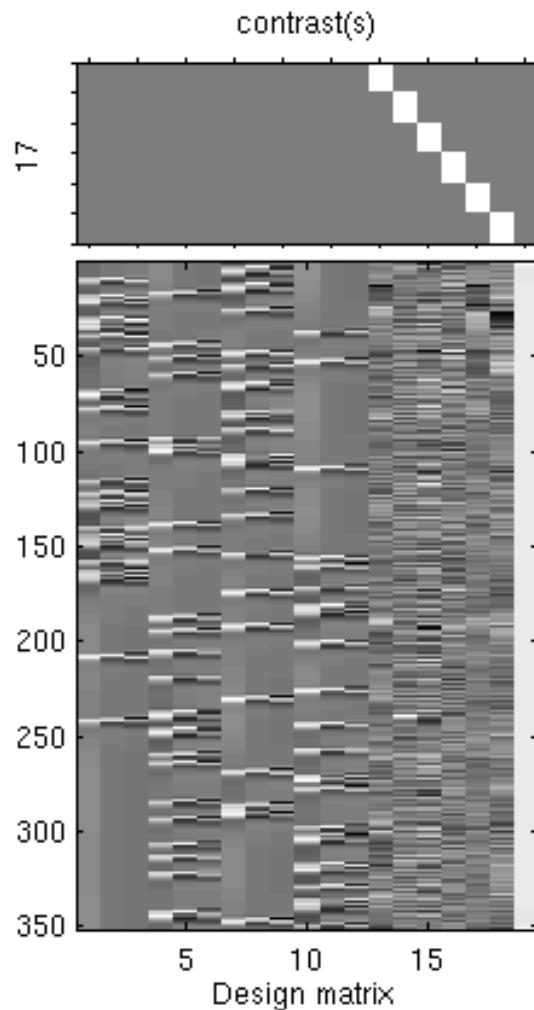
$$F \propto \frac{RSS_0 - RSS}{RSS}$$

$$F \propto \frac{ESS}{RSS} \sim F_{v_1, v_2}$$

$$v_1 = \text{rank}(X) - \text{rank}(X_0)$$

$$v_2 = N - \text{rank}(X)$$

F-test example: movement related effects

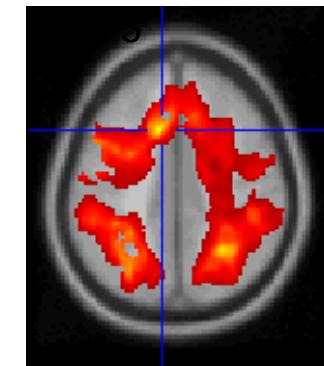
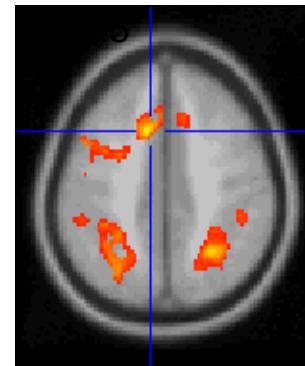
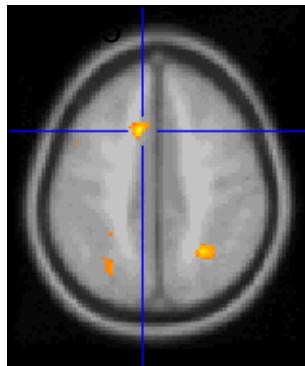


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Multiple comparison problem

High Threshold → Med. Threshold → Low Threshold



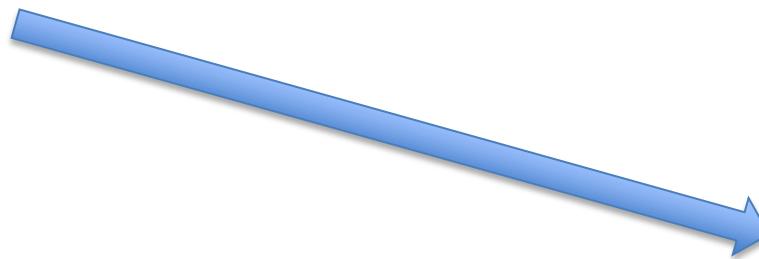
Good Specificity

Poor Power
(risk of false
negatives)

Poor Specificity

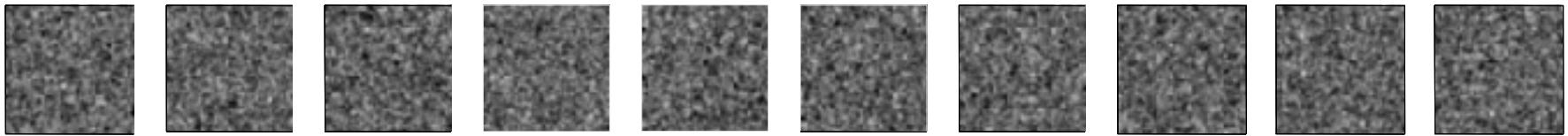
(risk of false
positives)

Good Power

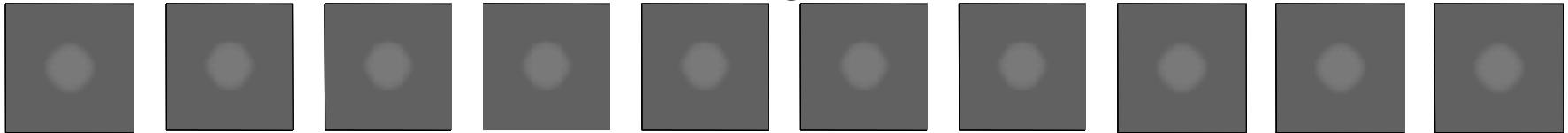


Multiple comparison problem

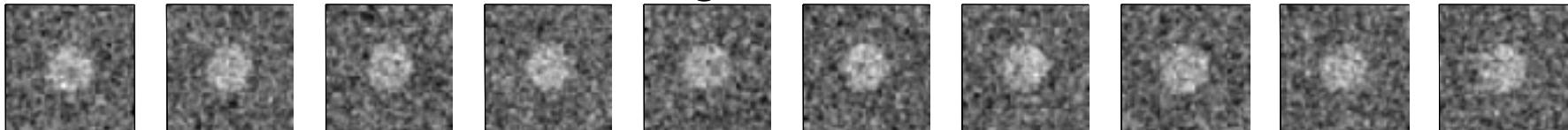
Noise



Signal

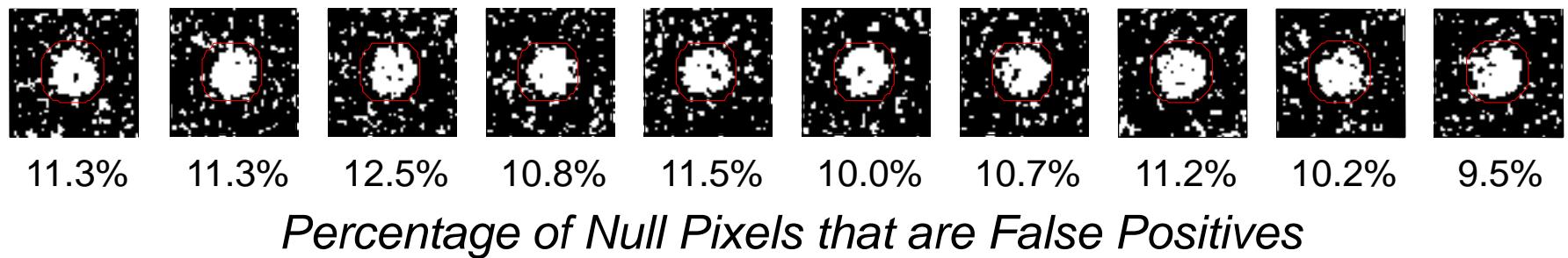


Signal+Noise

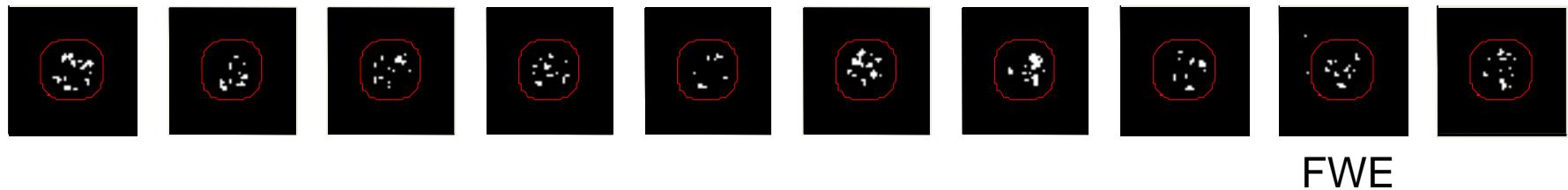


Multiple comparison problem

Use of ‘*uncorrected*’ p-value, $\alpha = 0.1$



Use of ‘*corrected*’ p-value, $\alpha=0.1$

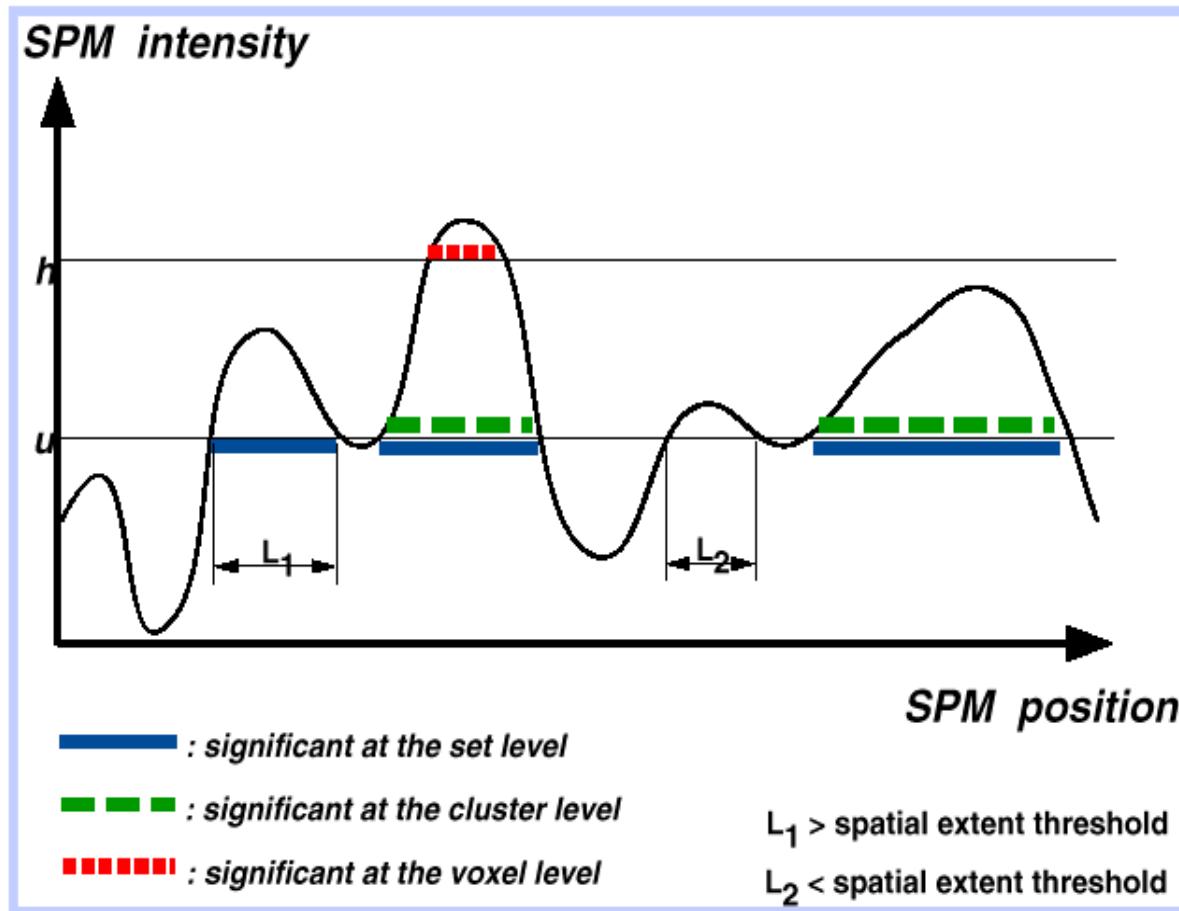


Multiple comparison problem

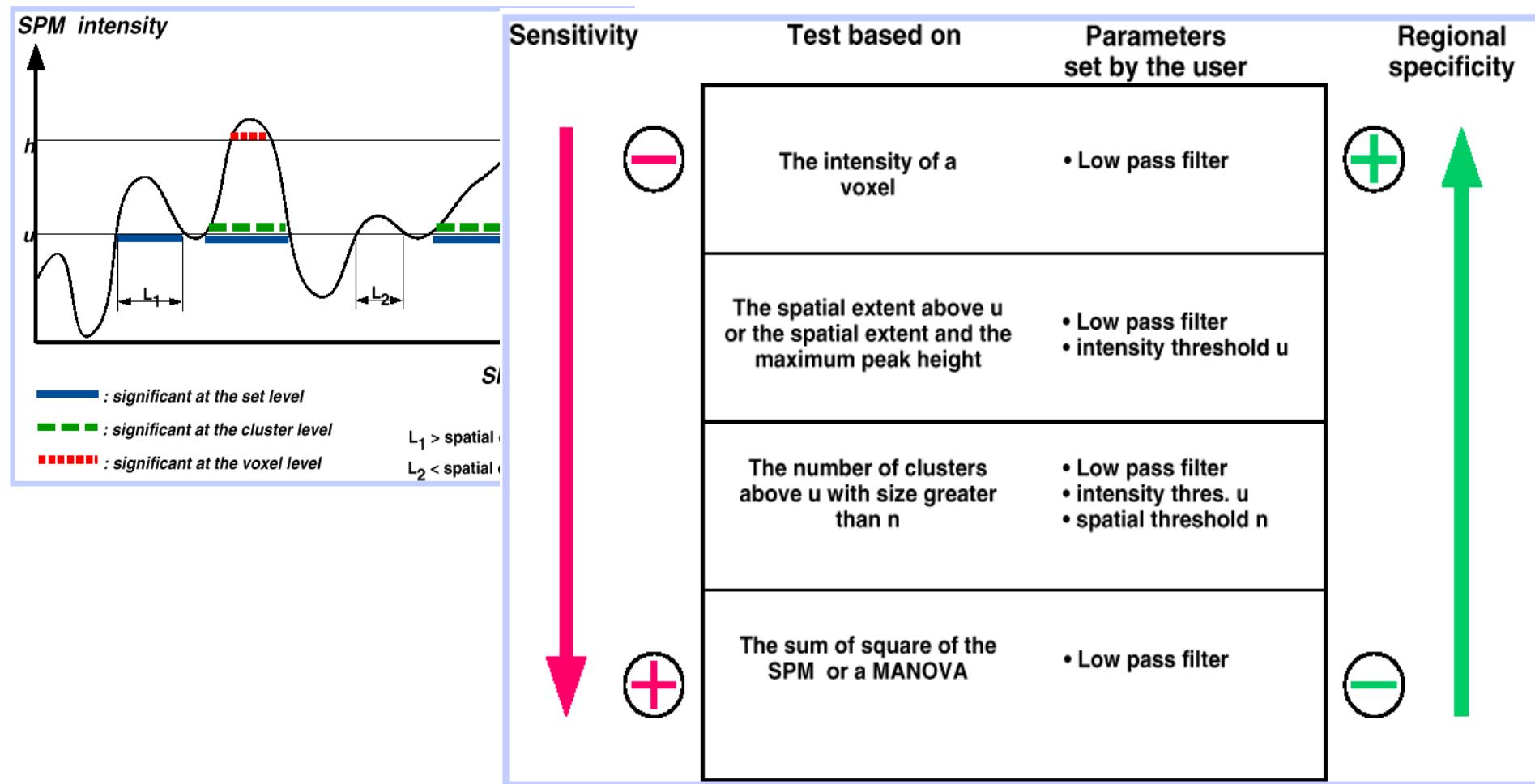
Inference with multiple comparison in neuroimaging depends on:

- search volume = # tests performed
 - smoothness of the data
 - size & shape of volume of interest
 - statistics employed (t- or F-)
 - + degrees of freedom
- usually derived from data and test performed

More inference levels



More inference levels



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Conclusion

From a ‘neuroimager’ perspectives, GLM is useful for:

- Localising an effect of interest, while accounting for
 - effects of no interest
 - confounding effects
 - HRF shape (for fMRI)
- Clean signal & functional connectivity

Conclusion

From a ‘machine learner’ perspectives,
GLM is useful for:

- Check for « *is there any effect?* »
- Data filtering (e.g. confound removal)
- HRF deconvolution for fMRI (single event modeling) or subject’s response extraction (contrast building)
- Feature selection/masking

Thank you for your attention!

Any question?

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