Second level analysis of fMRI

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Topics

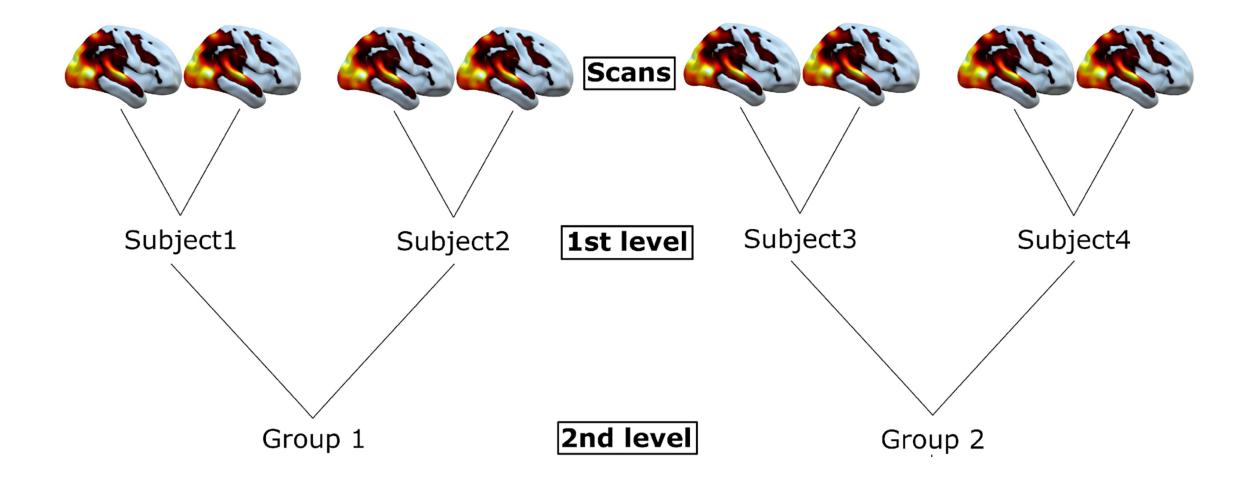
Introduction to the second level analysis

Theoretical framework of the group analysis

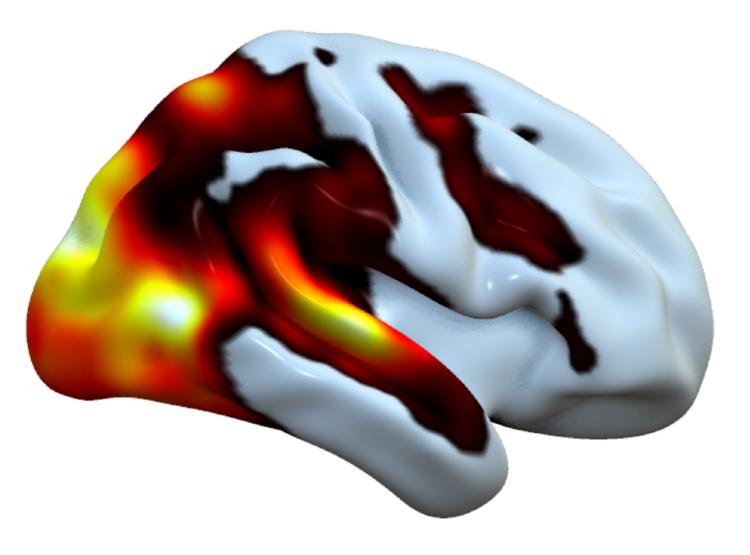
Second level models

Multiple comparisons problem

fMRI data are hierarchical



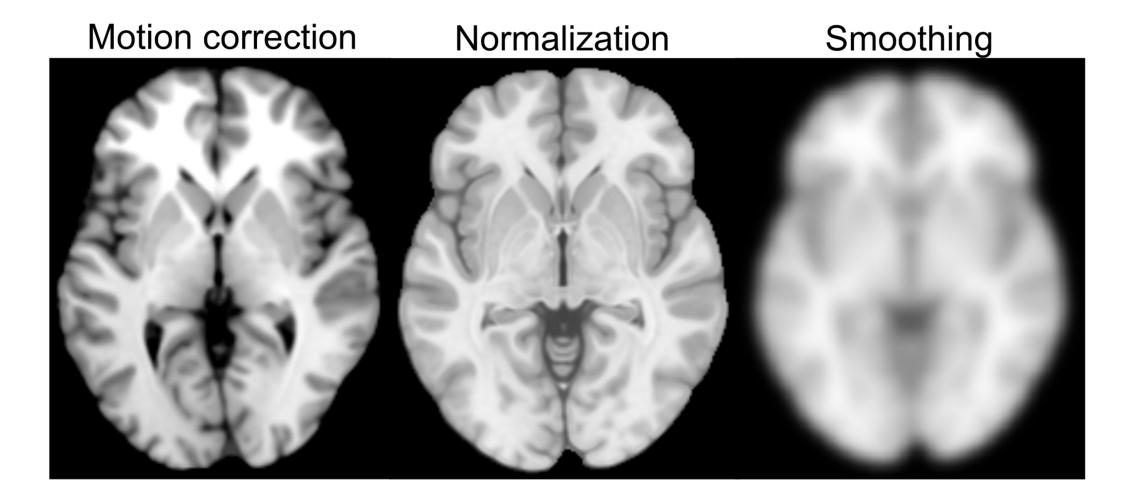
Whole brain analysis Voxel-level analysis Massive univariate analysis



Region-of-interest (ROI) analysis

Temporal Pole Supe		
Tomporal Johol He	schl	•
Temporal lobe He Temporal Supe		
Temporal Mic		•
Occipital Infe		•
Occipital Mid	ddle	•
Occipital Supe		•
Occipital lobe	gual	
Calca		•
Cun		•
Fusif	orm	
Parietal Supe	erior	*
Parietal Infe	erior	
Supramarg	inal	
Rolandic Opercu	lum	
Parietal lobe Precun	eus	
Ang	ular	
Paracentral Lot	bule	
Postcer	ntral	
Precer		
Supplementary Motor A		
Cingulate Poste		•
		•
		•
Frontal lobe Frontal Superior Me	Cingulate Middle Cingulate Anterior Frontal Superior	
Floritativit		
Frontal Inferior Opercu		
Frontal Inferior Or		
Orbitofrontal Poste		
Orbitofrontal Lat		
Olfac		
Parahippocan		•
Hippocam		
	sula	•
Subcortex Amygo		
India		
Pallic		
Cau		
Putar	men	

Preprocessing



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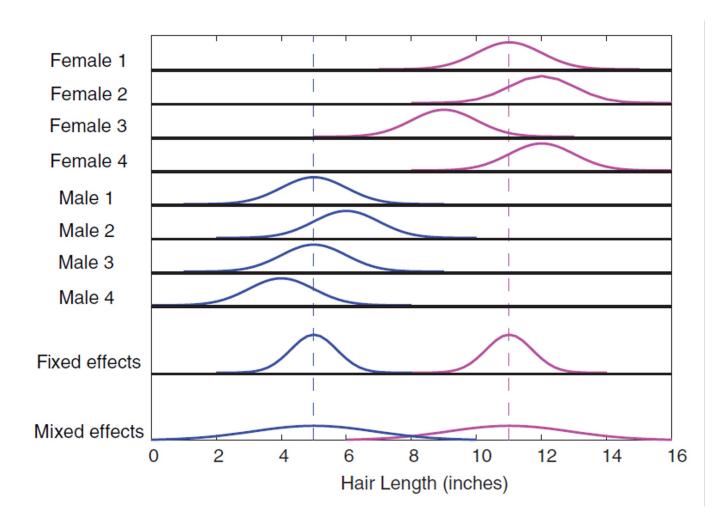
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Sources of variation in fMRI (or hair length)

Var^W = within-subject variance Var^B = between-subject variance



(Poldrack, Nichols, & Mumford, 2011)

Fixed effects model

Mixed effects model

Var^w

Variance used in the analysis

Var^w + Var^B

Describe study sample only



Generalize to population

Combine repeated measures within subjects



Group analysis of fMRI

Mixed effects model mathematically

• Within subject variance estimation (1st level)

$$Y = \beta X + \varepsilon, \qquad \quad \varepsilon \sim N(0, \sigma)$$
 (1)

• Between subject variance estimation (2nd level)

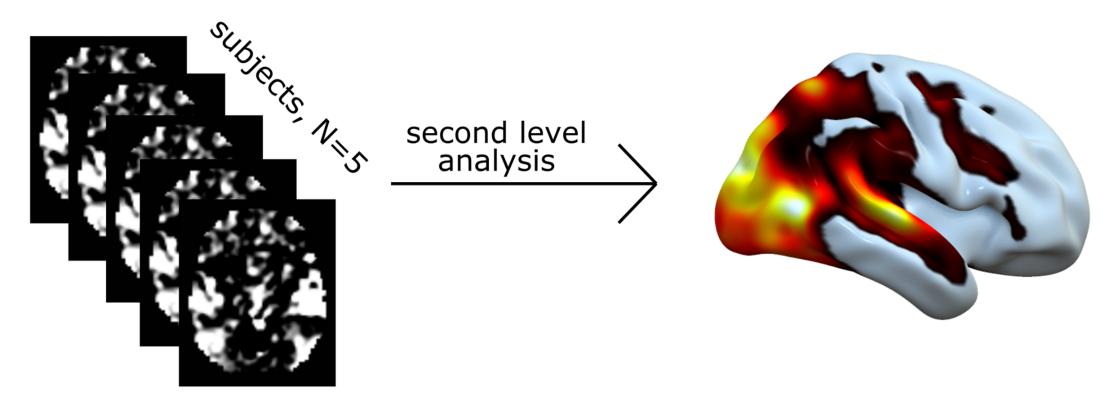
$$\beta = \beta_b X_b + \varepsilon_b \qquad \varepsilon_b \sim N(0, \sigma_b)$$
 (2)

Full mixed effects model

$$Y = (\beta_b X_b + \varepsilon_b) X + \varepsilon$$
(3)

Computationally demanding to estimate!

Summary statistics approach (mixed effect model)



First-level statistical parametric maps

Population level results

Topics

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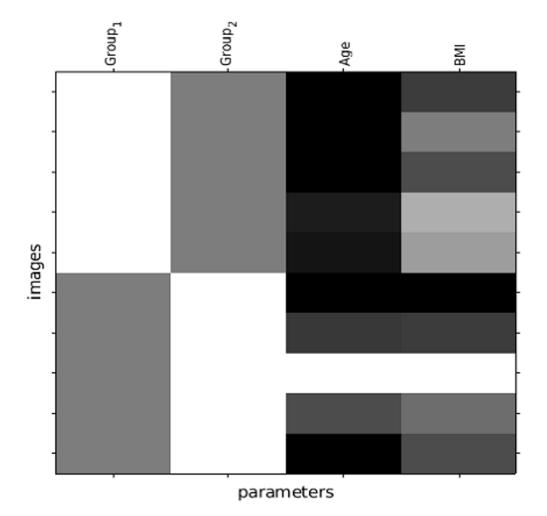
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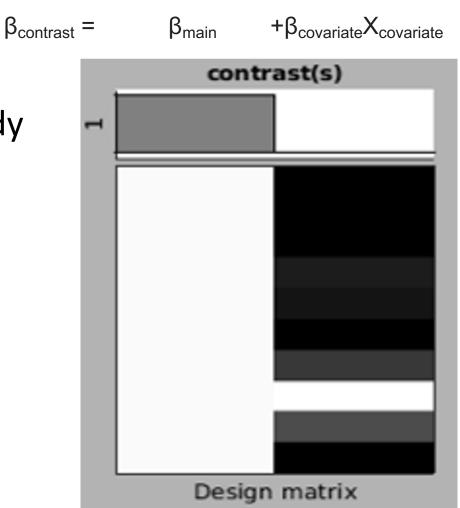
Second level design matrix in SPM

- Dependent variable:
 - β values from 1st level analysis
- Rows
 - subjects
- Columns:
 - Variables describing between subject differences
 - Effect of each column will be estimated



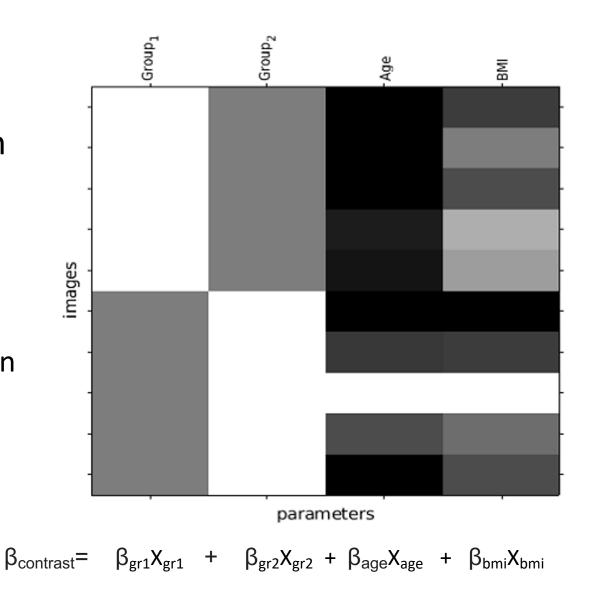
One sample T-test

- H: Brain activity is associated with the study condition somewhere in the brain
- Contrast (specified in the 1st level)
 - Main effect
 - Condition1 Condition2
- Covariates of interest
 - Positive association [0 1] = $\beta_{covariate}$
 - Negative association $[0 1] = -\beta_{covariate}$
- Main effect [1 0] = β_{main}



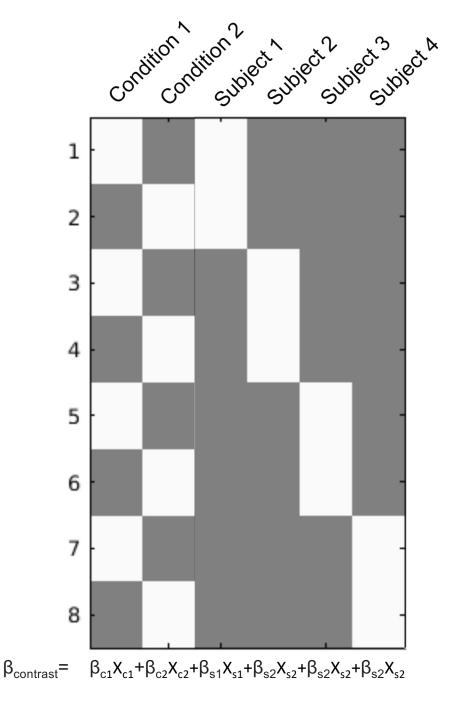
Two sample T-test

- H: Brain activity is different between two groups of subjects
- Group1 > Group2 with nuisance covariates
 - $[1 1 0 0] = \beta_{gr1} \beta_{gr2}$
 - "Whether females have increased brain response for the condition than males when age and BMI are controlled for"



Paired T-test

- H: There is a difference between conditions in the brain response
- 4 subjects, 2 scans per subject
- Condition
 - Cross-sectional
 - Baseline (1) vs. Stimulus (2)
 - Longitudinal
 - Baseline (1) vs. After treatment (2)
- Condition 2 > Condition 1
 - [-1 1 0 0 0 0 0] = - β_{c1} + β_{c2}
 - "Brain response associated with happy faces is higher after treatment"



Topics

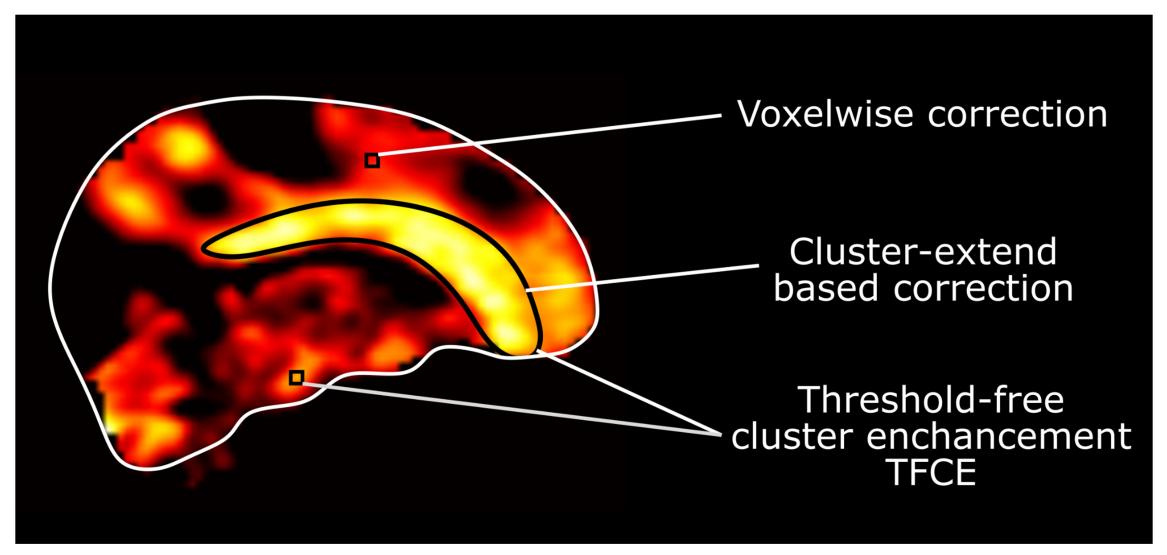
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Multiple comparisons correction methods



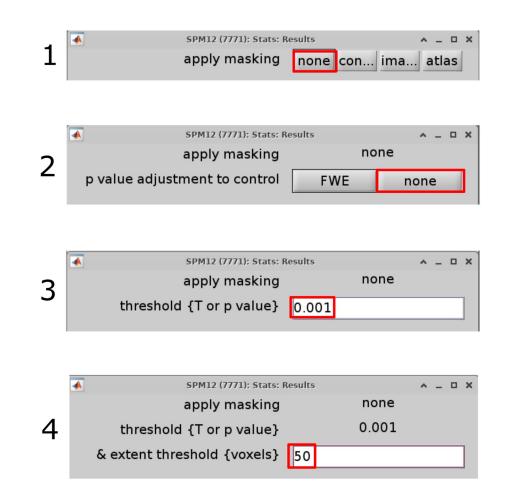
(Review: Lindquist & Mejia, 2015)

Voxelwise multiple comparisons correction

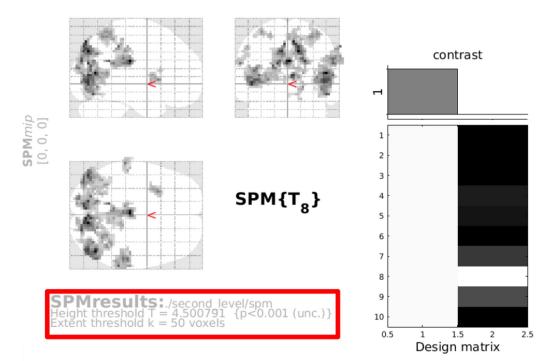
- Family-wise error rate (FWER) (Lindquist & Mejia, 2015)
 - Probability of making one or more false positives
 - Bonferroni correction
 - "There is a 5% probability of making at least one false positive finding"
 - 0.05 / number of tests = corrected p-value threshold
- False discovery rate (Benjamini & Hochberg, 1995)
 - "No more than 5% of our findings are false positives"

Cluster-extend based correction (Lindquist & Mejia, 2015)

- Accounts for the spatial dependency between voxels
- "What is the probability to observe a activating cluster of this size under the null hypothesis of no activation"
- Three-step procedure
 - Choose primary voxel-level threshold e.g. p < 0.001
 - 2. Choose minimum size of the cluster e.g. 50
 - 3. Control for FWER on a cluster level
- The approach may be problematic



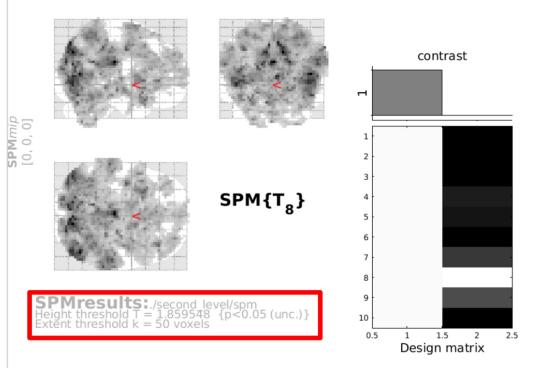
Main effect



Statistics:	p-values	adjusted	for	search	volume	
						-

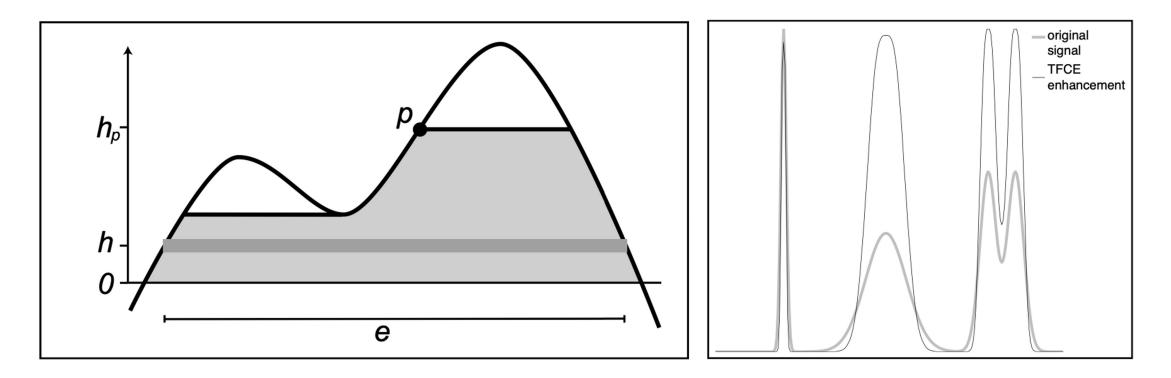
set-level		cluster-le	evel		peak-level					mm mm mm
p c	p _{FWE-corr} q _{FDR-corr} k _E p _{unco}			p _{uncorr}	$p_{\text{FWE-corr}} q_{\text{FDR-corr}} T$ (Z _E)				p _{uncorr}	
0.000 10	0.000	0.000	453	0.000	0.014	0.507	14.69	5.04	0.000	48 - 69 12
					0.099	0.507	11.33	4.65	0.000	45 - 78 3
					0.188	0.581	10.40	4.52	0.000	48 -72 -15
	0.000	0.000	166	0.000	0.016	0.507	14.39	5.01	0.000	0 -24 27
					0.242	0.581	10.06	4.46	0.000	-12 -51 27
					0.593	0.581	8.91	4.27	0.000	0-33 24
	0.000	0.000	1011	0.000	0.019	0.507	14.10	4.98	0.000	12 - 78 39
					0.040	0.507	12.77	4.83	0.000	6 -81 51
					0.061	0.507	12.06	4.75	0.000	3-78 9
	0.000	0.000	202	0.000	0.296	0.581	9.79	4.42	0.000	66 - 39 45
					0.296	0.581	9.79	4.42	0.000	57 - 27 24
					0.425	0.581	9.32	4.34	0.000	57 - 33 30
	0.000	0.000	57	0.000	0.536	0.581	9.03	4.29	0.000	-63 -30 24
					0.999	0.606	7.34	3.94	0.000	-54 -27 45
					1.000	0.824	5.45	3.43	0.000	-57 -33 33
	0.000	0.000	68	0.000	0.764	0.581	8.61	4.21	0.000	-27 12 3
					1.000	0.641	6.70	3.79	0.000	-30 3 3
	0.000				1.000	0.719	6.05	3.61	0.000	-21 15 15
	0.000	0.000	55	0.000	0.831	0.591	8.51	4.19	0.000	18 -66 -6
					1.000	0.694	6.24	3.66	0.000	18 - 60 0
	0.000	0.000	120	0 000	1.000	0.705	6.20	3.65	0.000	12 - 72 - 9
	0.000	0.000	130	0.000	0.991	0.606	8.17	4.12	0.000	30-60 57

Main effect



set-level cluster-level					peak-level						mm mm mr		
p c	P _{FWE-co}	rr q _{FDR-corr}	k _E	p _{uncon}	P _{FWE-co}	p _{FWE-corr} q _{FDR-corr} T (Z _E							
1.000 3	0.000	0.000		0.000	0.014	0.309	14.69	5.04	0.000	48 ·	-69	12	
					0.016	0.309	14.39	5.01	0.000	0 -	-24	27	
					0.019	0.309	14.10	4.98	0.000	12 -	- 78	39	
	0.003	0.000	668	0.000	0.536	0.355	9.03	4.29	0.000	-63 -	-30	2	
					0.999	0.370	7.34	3.94	0.000	-54 -	- 27	4	
					1.000	0.503	5.45	3.43	0.000	-57 -	- 33	3	
	0.893	0.253	163	0.013	1.000	0.490	5.60	3.48	0.000	-36	39	24	
					1.000	0.503	5.34	3.39	0.000	- 39	45	3	
					1.000	0.658	4.22	2.98	0.001	-24	42	3	

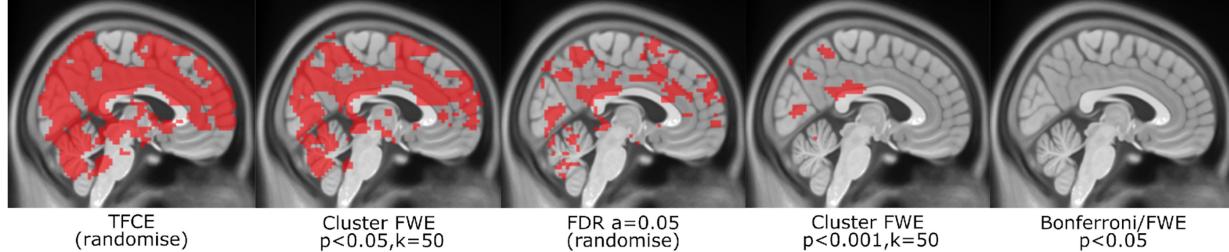
Threshold-free cluster enchancement (TFCE)



TFCE = h_p (voxelwise t-value) * e (amount of supporting voxels)

The voxelwise significance is adjusted by the amount supporting voxels Significance of each voxel is assessed with permutations and then corrected for multiple comparisons

(Smith & Nichols, 2009)



(randomise)

p<0.001,k=50

TFCE (randomise)

False negatives False positives

Conservative correction inflates effect size estimates

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- Motivation for non-parametric tests in group analyses (Eklund, Nichols, & Knutsson, 2016)
 - 1. Parametric tests may produce false positive findings in group level analyses after multiple comparisons corrections
 - 2. Voxelwise multiple comparisons methods may produce too conservative findings and cluster-based methods false positives
 - 3. Non-parametric tests have been shown to correct better for multiple comparisons.
- Tools for non-parametric tests
 - SnPM (Doc: https://warwick.ac.uk/fac/sci/statistics/staff/academic-research/nichols/software/snpm)
 - FSL Randomise (Winkler, Ridgway, Webster, Smith & Nichols, 2014)
 - One and two sample (unpaired/paired) T-tests, repeated measures anova
 - Easy to output statistical result maps with various different multiple comparisons methods
 - Includes TFCE method
 - Doc: <u>https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/Randomise</u>

References

Benjamini, Y., & Hochberg, Y. (1995). Controlling the False Discovery Rate - a Practical and Powerful Approach to Multiple Testing. Journal of the Royal Statistical Society Series B-Statistical Methodology, 57(1), 289-300. doi:DOI 10.1111/j.2517-6161.1995.tb02031.x

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Lindquist, M. A., & Mejia, A. (2015). Zen and the art of multiple comparisons. *Psychosom Med*, 77(2), 114-125. doi:10.1097/PSY.0000000000000148

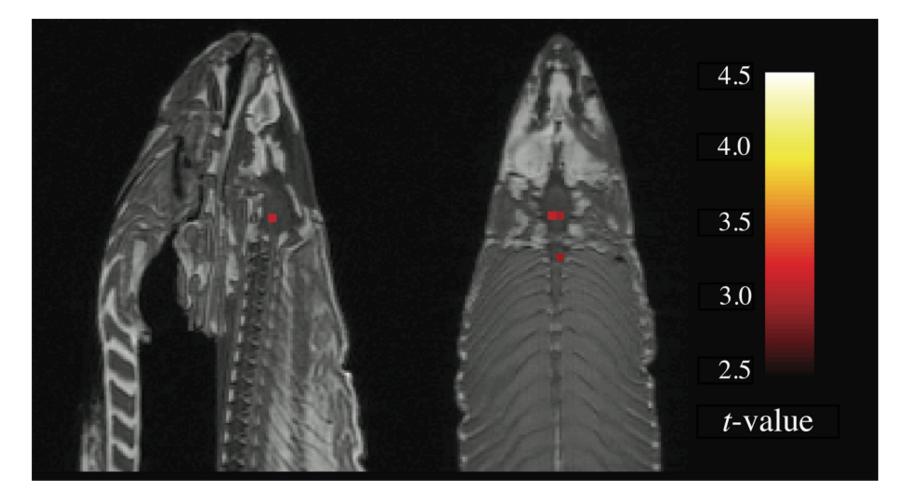
Poldrack, R. A., Nichols, T., & Mumford, J. (2011). Handbook of Functional MRI Data Analysis. *Cambridge: Cambridge University Press*. doi:10.1017/cbo9780511895029

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Winkler, A. M., Ridgway, G. R., Webster, M. A., Smith, S. M., & Nichols, T. E. (2014). Permutation inference for the general linear model. *Neuroimage*, *92*, 381-397. doi:10.1016/j.neuroimage.2014.01.060

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Thank you!



(Bennet, 2011)