## Permutation-based true discovery guarantee by sum tests

28 Sept. 2022 – XXX Conference of the Italian Association of Psychology

Anna Vesely

Joint work with Livio Finos and Jelle L Goeman

University of Padova anna.vesely@unipd.it



## Overview

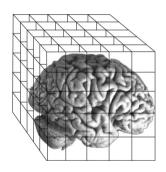
## **Functional Magnetic Resonance Imaging**

fMRI measures brain activation as changes in blood flow (BOLD) under a sequence of stimuli

Activation is measured in voxels  $\approx 200,000$  highly correlated volume units

#### Voxel i:

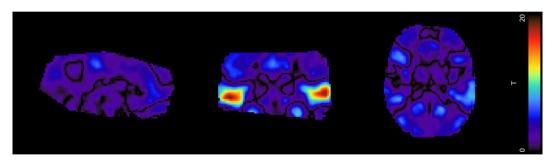
- null hypothesis  $H_i$ : no activation in voxel i
- test statistic  $T_i$  from first-level analysis



## **Auditory data**

140 subjects passively listening to vocal and non-vocal sounds (168,211 voxels)<sup>1</sup>

 $H_i$ : no activation in voxel i  $T_i = \text{one-sample t-statistic}$ 



<sup>&</sup>lt;sup>1</sup>Pernet et al. The human voice areas: spatial organisation and inter-individual variability in temporal and extra-temporal cortices. OpenNeuro dataset, 2019.

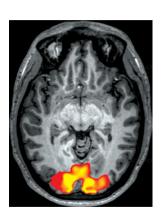
## **Functional Magnetic Resonance Imaging**

Generally interest lies in clusters, brain regions of (contiguous) voxels

# Supra-threshold clusters: regions of connected voxels *i* with $T_i >$ threshold

#### Clusters S:

- null hypothesis  $H_S = \bigcap_{i \in S} H_i$ : no activation in cluster S
- global test statistic  $T_S$



## Multiple hypothesis testing

## Multiple hypothesis testing

#### Single null hypothesis $H_i$ :

- type I error = wrongly reject  $H_i$  when it is true
- significance level  $\alpha = P(\text{type I error})$

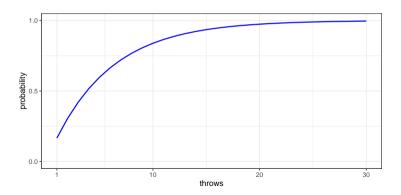
#### Multiple hypotheses together:

the probability of making at least one type I error can be much greater than  $\boldsymbol{\alpha}$ 

#### Intuition: m throws of a dice

The probability of getting at least one 6 is

- $1/6 \approx 0.167$  for m = 1
- $1 (5/6)^m$  in general



## Standard cluster inference

## Cluster extent thresholding

#### Standard method for cluster inference that

- adapts to the unknow correlation structure
- strongly controls the FWER at cluster level

 $H_S$  is rejected  $\Longrightarrow S$  contains at least one active voxel

#### No information on

- the proportion of active voxels (TDP)
- their spatial location

## Cluster extent thresholding

The following statements are not supported:

## 'A large significant cluster...'

- '... contains a substantial number of active voxels.'
   It contains at least one.
- '... is a stronger finding than a small significant cluster.'
   Spatial specificity paradox.
   Follow-up inference leads to inflated type I error rates.
- '... indicates activity in an anatomical area, if there is substantial overlap.'

  If the cluster is not completely contained, activity may lie outside.

## True discovery guarantee

## True discovery guarantee

TDP(S) = proportion of truly active voxels in cluster S

Lower  $(1 - \alpha)$ -confidence bounds:

$$P(\mathsf{TDP}(S) \geq \mathsf{bound}(S)) \geq 1 - \alpha$$

Closed testing 12 gives simultaneous bounds:

$$P(\mathsf{TDP}(S) \ge \mathsf{bound}(S) \text{ for each cluster } S) \ge 1 - \alpha$$

<sup>&</sup>lt;sup>1</sup>Genovese and Wasserman. Exceedance control of the false discovery proportion. *JASA*, 2006.

<sup>&</sup>lt;sup>2</sup>Goeman and Solari. Multiple testing for exploratory research. *Stat. Sci.*, 2011.

#### sumSome

Computational complexity of closed testing: exponential in the number of voxels

ightarrow infeasible for high-dimensional data

We provide a shortcut<sup>1</sup>, valid in many cases, that

- makes inference on the TDP of clusters
- allows for post-hoc selection and follow-up inference
- adapts to the unknown correlation structure of voxels

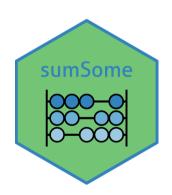
<sup>&</sup>lt;sup>1</sup>Vesely et al. Permutation-based true discovery guarantee by sum tests. arXiv:2102.11759, 2021.

## Results

## sumSome package

The method is implemented in the R package sumSome<sup>1</sup>, with underlying code in C++.

```
permT <- brainScores(copes, alpha=0.05, ...)
brainAnalysis(permT, clusters)</pre>
```

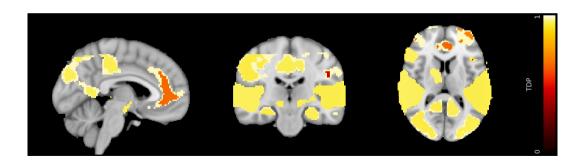


<sup>&</sup>lt;sup>1</sup>https://CRAN.R-project.org/package=sumSome

## **Auditory data**

We obtain simultaneous confidence bounds for the TDP of clusters:

- clusters with  $T_i > 3.2$
- sub-clusters with  $T_i > 4$



## **Auditory data**

cluster	threshold	size	TDP (%)	coordinates		
S	thr	S	lower conf. bound	X	У	Z
FP/CG/SFG/TOF/LO/LG	3.2	40,094	98.21	-30	-34	-16
OFG/ITG/SG/AG/NA						
Left LO/TOF	4	8,983	94.79	-30	-34	-16
Right LO/LG/ITG	4	7,653	93.85	28	-30	-18
Left SFG/FP	4	1,523	69.67	-28	34	42
CG	4	1,341	65.62	6	40	-2
Right FP	4	1,327	66.01	30	56	28
Left SG/AG	4	859	47.85	-50	-56	36
Right STG/PT/MTG	3.2	12,540	95.41	60	-10	0
HG/PrG/T						
STG/PT/MTG/HG	4	9,533	95.17	60	-10	0
PrG	4	485	25.15	52	0	48
Left STG/PT/MTG/	3.2	10,833	94.66	-60	-12	2
HG/IFG/T						
HG/PT/MTG/STG	4	7,894	94.20	-60	-12	2
IFG	4	667	38.98	-40	14	26

#### **Conclusion**

Making inference on the TDP allows to quantify and localize brain activation in clusters.

sumSome is a general closed tesing method to give lower  $(1 - \alpha)$ -confidence bounds for the TDP, simultaneously over all clusters.

This way, results are valid even if

- the cluster of interest is chosen post hoc
- we make follow-up inference inside sub-clusters

## Questions

