# Simplified power and sample size calculations using prevalence \& magnitude of active peaks. Joke Durnez ${ }^{a}$, Beatrijs Moerkerke ${ }^{a}$, Ruth Seurinck ${ }^{a}$ and Thomas Nichols ${ }^{b}$ 

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## Introduction

- There is increasing concern about statistical power in neuroscience research:
- Low power decreases the chance of detecting a true effect
- Low power reduces the chance that a statistically significant result indicates a true effect (loannidis, 2005).
$■ \Rightarrow$ A power analysis is a critical component of any study
- Power analyses for fMRI are difficult: need to specify magnitude, spatial extent and location of a hypothesized effect.
- We present a simple way to characterize the spatial signal in a fMRI study, and a direct way to estimate power based on an existing pilot study.
- We must estimate (or have prior knowledge of):
- the volume of the brain that is activated
- the average effect size in activated brain regions
- This allows the calculation of power for given sample size, brain volume and smoothness.


## Methods

- We start from peak $p$-values in a group level analysis. We estimate $\pi_{1}$, proportion of peak $p$-values thare are non-null, as described in Durnez, Moerkerke \& Nichols (2014).
- We assume that the null distribution of peak values is an exponential distribution (Worsley, 2007).
- We assume that the alternative distribution of peak values is a truncated normal distribution (truncated at excursion threshold $u$ ).
- Therefore, the distribution of peak values can be written as a mixture:

$$
f\left(x \mid \pi_{0}, \mu_{1}, \sigma_{1}, u\right)=\left(1-\pi_{1}\right) u \exp (-u(x-u))+\pi_{1} \frac{\frac{1}{\sigma_{1}} \varphi\left(\frac{x-\mu_{1}}{\sigma_{1}}\right)}{1-\Phi\left(\frac{u-\mu_{1}}{\sigma_{1}}\right)}
$$

- $\mu_{1}$ and $\sigma_{1}$ can be estimated using maximum likelihood, where $\mu_{1}$ is the expected peak height in activated regions.
- Power can be estimated for a given threshold $t$ as $P\left(T>t \mid H_{a}\right)$ with $T$ the $T$-statistic of the peak.


## Example

- Data from Seurinck et al. (2011). The original study was analyzed voxelwise with FWER-control at the 5\% significance level. We reanalyzed the data peakwise with FDR-control at the $5 \%$ significance level.
- Sample size: 13
- Step 1: compute prevalence of activation
- Step 2: estimate truncated distributions

- Expected peak height under activation $\mu_{1}=\delta=4.04$
- Effect size $\mu / \sigma=\delta / \sqrt{n}=1.13$
- Step 3: estimate power for given sample size, estimated peak height under activation and prevalence of activation $\left(\pi_{1}\right)$



## Simulations

- 100 full-brain datasets with smooth Gaussian noise (3 voxels) superimposed with activation (3 \% BOLD change) in 4 foci ( $3 \%$ of total brain volume)
- Prevalence of activation: For small sample sizes: very conservative estimates (close to 0 ).


- Effect size estimation: Rather conservative estimates.

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- Alternative distribution: Constant effect size $(\mu / \sigma)$ over different sample sizes to be used in power analyses.

- Power calculations: Good estimates for realistic values of power with FWER control at the 5\% significance level.



## References and acknowledgements

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