# Analysis of family-wise error rates in statistical parametric mapping using random field theory

Guillaume Flandin and Karl J. Friston

Wellcome Trust Centre for Neuroimaging University College London, UK

27th June 2016

#### Abstract

This technical report revisits the analysis of family-wise error rates in statistical parametric mapping – using random field theory – reported in (Eklund et al., 2015). Contrary to the understandable spin that these sorts of analyses attract, a review of their results suggests that they endorse the use of parametric assumptions – and random field theory – in the analysis of functional neuroimaging data. We briefly rehearse the advantages parametric analyses offer over nonparametric alternatives and then unpack the implications of (Eklund et al., 2015) for parametric procedures.

### 1 Introduction

Random field theory has been at the heart of statistical parametric mapping in neuroimaging – and its various implementations in academic software – for over two decades. With technical advances in data acquisition, its validity has been revisited every few years to ensure it is fit for purpose (Worsley et al., 1996; Hayasaka and Nichols, 2003; Hayasaka et al., 2004; Pantazis et al., 2005; Bennett et al., 2009; Nichols, 2012; Woo et al., 2014), particularly in relation to controlling family-wise error. The statistical validity of procedures based on random field theory is important because random field theory offers an efficient and reproducible alternative to nonparametric testing. The advantages of parametric approaches over nonparametric approaches include the following:

- Parametric approaches are more efficient than their nonparametric counterpart by the Neyman-Pearson lemma. This follows because the most efficient test is based upon the odds ratio inherent in parametric tests. This means that any nonparametric test can only be as efficient as a parametric test or less efficient.
- Parametric approaches are reproducible. In other words, one obtains the same result when repeating the analysis, unlike the *p*-values based upon samples of the null distribution used in nonparametric tests.

- Parametric approaches eschew the problem of complying with the exchangeability criteria of nonparametric procedures. These criteria make it difficult to apply nonparametric tests to data that have serial correlations or when using hierarchical models.
- Parametric approaches are computationally more efficient because they
  use distributional assumptions to eschew computationally intensive sampling from a null distribution.

These advantages rest upon distributional assumptions that, if violated, render parametric tests inexact. In these instances, one could consider using non-parametric tests. It is therefore important to ensure that parametric tests and random field theory are robust to any violations. The analyses reported by (Eklund et al., 2015) speak to this issue. So what conclusions can be drawn from these analyses?

# 2 A review of the Eklund et al simulation results

(Eklund et al., 2015) assess the family-wise error rate using parametric and nonparametric tests and a variety of regressors to analyse (publicly available) resting state fMRI data from two sites. They manipulate a number of factors including: (i) inference based upon peak height versus spatial extent; (ii) spatial extent inference based upon high versus low cluster forming thresholds; (iii) under different levels of spatial smoothing for (iv) block versus event-related regressors, using (v) one- and two-sample t-tests.

In brief, they show that parametric inference based upon peak height is well-behaved and provides acceptable family-wise error control. In contrast, parametric inference based upon spatial extent is not valid when, and only when, a low cluster forming threshold is employed. This failure is well known and is consistent with random field theory: the null distribution for spatial extent is based upon the Nosko conjecture that provides a distributional form for the spatial extent of a cluster (Friston et al., 1994). The parameter of this distributional form is fixed using approximations to the expected number of maxima and the total volume above a threshold (see (Flandin and Friston, 2015) for a brief review). Crucially, both the distributional form for the spatial extent and the expected number of maxima (the Euler characteristic) are approximations that are only true in the limit of high thresholds (see Figure 1 in (Friston et al., 1994)). This means that tests based upon spatial extent become inexact at low thresholds – as verified numerically by (Eklund et al., 2015).

The effects of smoothing reported in (Eklund et al., 2015) are consistent with random field theory, which assumes a good lattice approximation to a continuous random field. This assumption means that the data have to be smoother than the size of voxels. In other words, increasing the smoothness will lead to more exact inference. Again, this is verified numerically by (Eklund et al., 2015).

The effect of one versus two-sample t-tests is slightly more difficult to interpret. This is because the authors used the same regressor for all subjects. Arguably, this was a mistake because any systematic fluctuation in resting state timeseries – that correlates with the regressor – will lead to significant one-sample t-tests against the null hypothesis of zero (e.g., magnetic equilibration

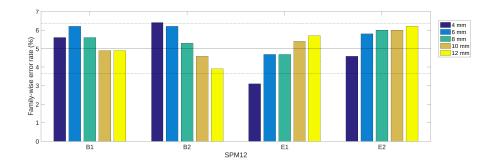


Figure 1: Cluster-level inference results for a two-sample t-test (two groups of ten random subjects, repeated a thousand times) with the Beijing dataset using a cluster-forming threshold of p < 0.001 and the SPM12 software (r6685). Five levels of spatial smoothing were evaluated (4, 6, 8, 10 and 12 mm isotropic Gaussian kernels) with four different regressors (see (Eklund et al., 2015) for details).

effects). This effect is particularly marked for a regressor (called E1) that represents a fast and inefficiently estimated event-related response every few seconds. Crucially, the nonparametric false positive rates are beyond the 95% confidence intervals. This means that this effect is actually expressed in the data over subjects and therefore fails as a model of the null behaviour.

This failure is finessed when comparing parameter estimates between two groups using a two-sample t-test. In this instance, inferences based upon spatial extent fall to acceptable family-wise error rates. We confirmed this by reproducing the analysis (using the same data and regressors) reported in (Eklund et al., 2015) (see Figure 1). These analyses use the close to original (3 mm) voxels sizes – as opposed to the upsampled (2 mm voxel) data as analysed in (Eklund et al., 2015).

### 3 Conclusion

The results provided by (Eklund et al., 2015) endorse the use of random field theory and reveal behaviour that is consistent with the underlying distributional assumptions. Having said this, there was a problem with the construction of the null distributions presented in (Eklund et al., 2015). This is because they used the same regressors for every subject. However, this problem is finessed by the use of two-sample t-tests. The results of these analyses (two-sample t-test at high thresholds) show that the random field theory provides valid inference based on spatial extent, provided its distributional assumptions are not violated (through the use of low cluster forming thresholds or smoothing). This conclusion is important because the issues addressed in (Eklund et al., 2015) underwrite good practice in imaging neuroscience.

# 4 Acknowledgements

We would like to thank our colleagues Chris Mathys and Will Penny for their help in the preparation of this report. We are also grateful to the 1000 Functional Connectomes Project<sup>1</sup>, the Neuroimaging Informatics Tools and Resources Clearinghouse (NITRC) and the corresponding researchers for making publicly available the resting state datasets used in our analyses.

## References

- Bennett, C. M., Wolford, G. L., and Miller, M. B. (2009). The principled control of false positives in neuroimaging. *Social Cognitive and Affective Neuroscience*, 4:417–422.
- Eklund, A., Nichols, T., and Knutsson, H. (2015). Can parametric statistical methods be trusted for fMRI based group studies? arXiv:1511.01863.
- Flandin, G. and Friston, K. J. (2015). Topological inference. In Toga, A., editor, *Brain Mapping: An Encyclopedic Reference*, volume 1, pages 495–500. Academic Press.
- Friston, K. J., Worsley, K. J., Frackowiak, R. S. J., Mazziotta, J. C., and Evans, A. C. (1994). Assessing the significance of focal activations using their spatial extent. *Human Brain Mapping*, 1:210–220.
- Hayasaka, S. and Nichols, T. E. (2003). Validating cluster size inference: random field and permutation methods. *NeuroImage*, 20:2343–2356.
- Hayasaka, S., Phan, K. L., Liberzon, I., Worsley, K. J., and Nichols, T. E. (2004). Non-stationary cluster size inference with random field and permutation methods. *NeuroImage*, 22:676–687.
- Nichols, T. E. (2012). Multiple testing corrections, nonparametric methods, and random field theory. *NeuroImage*, 62:811–815.
- Pantazis, D., Nichols, T. E., Baillet, S., and Leahy, R. M. (2005). A comparison of random field theory and permutation methods for the statistical analysis of MEG data. *NeuroImage*, 25:383–394.
- Woo, C.-W., Krishnan, A., and Wager, T. (2014). Cluster-extent based thresholding in fMRI analyses: pitfalls and recommendations. *NeuroImage*, 1(91):412–419.
- Worsley, K. J., Marrett, S., Neelin, P., Vandal, A. C., Friston, K. J., and Evans, A. C. (1996). A unified statistical approach for determining significant voxels in images of cerebral activation. *Human Brain Mapping*, 4:58–73.

http://fcon\_1000.projects.nitrc.org/