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Inflated false positive rates in fMRI depend on the voxel size of normalized images

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Introduction

- In a recent manuscript, Eklund et al. [1] reported inflated false positive rates in functional magnetic resonance imaging (fMRI) using common software packages including SPM, FSL, and AFNI.
- Briefly, a nominal family-wise error rate of 5% in the parametric statistical evaluation was shown to be conservative for voxel-wise inference but not for cluster-wise inference.
- As a cause of the observed invalid cluster inferences, the authors suggested that the spatial autocorrelation functions do not follow the assumed Gaussian shape.
- We would like to draw attention to an important aspect that was not addressed in this publication: **Statistical inferences obtained using the Gauss random field approach depend heavily on a pre-processing parameter that was not included in the analysis performed by Eklund et al. [1], namely the spatial resolution to which the data are resampled and interpolated during pre-processing.**
- Eklund et al. [1] used the common default setting of $2 \times 2 \times 2 \text{ mm}^3$.
- In response to the paper by Eklund et al., Flandin and Friston [2] used a different setting of this parameter, namely $3 \times 3 \times 3 \text{ mm}^3$. Together with a more stringent initial cluster-forming threshold, they did not observe inflated false positive rates.
- However, a spatial resolution of $2 \times 2 \times 2 \text{ mm}^3$ is the default value in two major software packages (SPM, FSL) and, hence, it is likely to be used for processing fMRI data by these packages.
- In previous work, Friston and colleagues [3] stated that resampling to $2 \times 2 \times 2 \text{ mm}^3$ renders the analysis “more sensitive”.
- It is, thus, completely unclear what a valid setting for this parameter should be. Therefore, it is of substantial relevance to systematically assess its influence on statistical inference.

Methods

- We analyzed 47 resting-state fMRI data sets, each acquired at a nominal spatial resolution of $3 \times 3 \times 4 \text{ mm}^3$ with 300 volumes.
- Using a strategy analogous to that of Eklund et al. [1] we imposed various fake designs including block- and event-related types.
- Analysis was performed using SPM12.
- Normalization included a resampling with three different voxel sizes: $3 \times 3 \times 3 \text{ mm}^3$, $2 \times 2 \times 2 \text{ mm}^3$, and $1 \times 1 \times 1 \text{ mm}^3$.
- Two normalization pipelines: (a) normalizing the raw data and performing the individual statistics with normalized data, and (b) normalizing the contrast images.
- Using a family-wise error (FWE) correction for multiple comparisons based on the Gauss Random Field approach, we first evaluated each data set separately. Thereafter, we performed a group-level inference in which all 47 data sets were pooled.

Results & Discussion

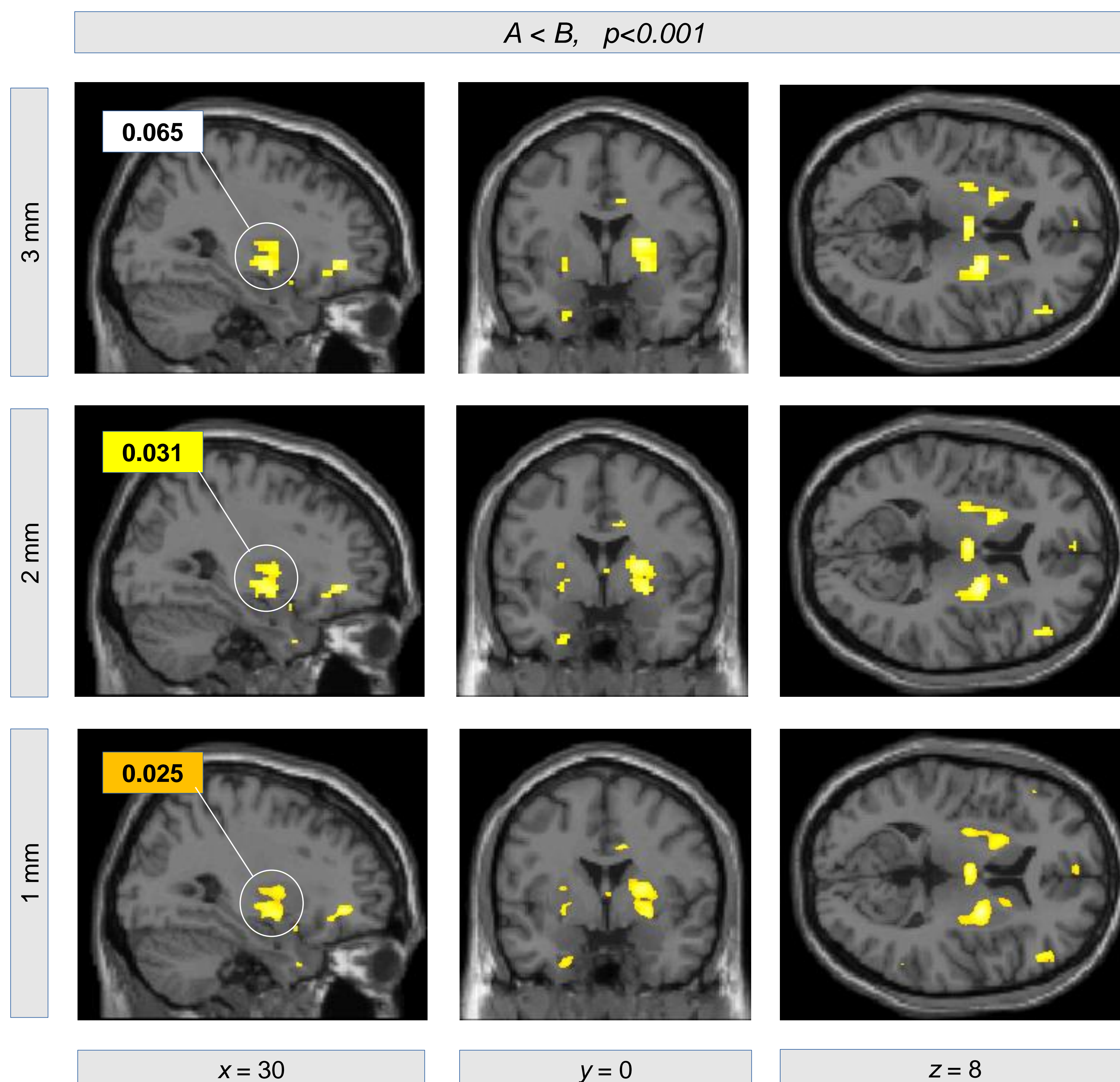


Figure 1. Orthogonal brain sections showing brain activity differences between two experimental conditions A and B using a fake design with resting-state fMRI data. During normalization into the standard space, data were scaled to $3 \times 3 \times 3 \text{ mm}^3$, $2 \times 2 \times 2 \text{ mm}^3$, and $1 \times 1 \times 1 \text{ mm}^3$. Family-wise error (FWE) corrected p -values become smaller when using a higher upscaling. The white box shows the FWE-corrected p -value of a selected cluster that reaches significance that is a false positive result.

- On the individual level, we found that with higher resampling resolutions, the FWE-corrected p -values decrease systematically so that more and more false positives occur.
- As expected by our null hypothesis, we did not obtain any positive clusters with $3 \times 3 \times 3 \text{ mm}^3$ resolution. However, with $2 \times 2 \times 2 \text{ mm}^3$, the p -values are already smaller leading to a significant cluster. With a resolution of $1 \times 1 \times 1 \text{ mm}^3$, the p -value decreased again.
- Figure 1 shows the effect using a fake event-related design with two experimental conditions.
- We obtained the same effect when using an arbitrary on/off-design with a block length of 20 s.
- It appears that there is a **systematic** dependence of the false positive rate on the resampling parameter with smaller voxel sizes leading to smaller FWE-corrected p -values and hence more false positives.
- While some dependence on pre-processing parameters may be inevitable, a systematic dependence of this type is clearly worrisome, because researchers may be tempted to interpolate their data until the desired statistical significance level is reached.
- Statistical inference should certainly not depend in such a systematic way on a pre-processing parameter that can be set *ad libitum*.
- Clearly, this issue requires further in-depth analysis.

References

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