

A Self-Supervised Voxel Shuffling Framework for Kernel-Based fMRI activation detection

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Introduction:

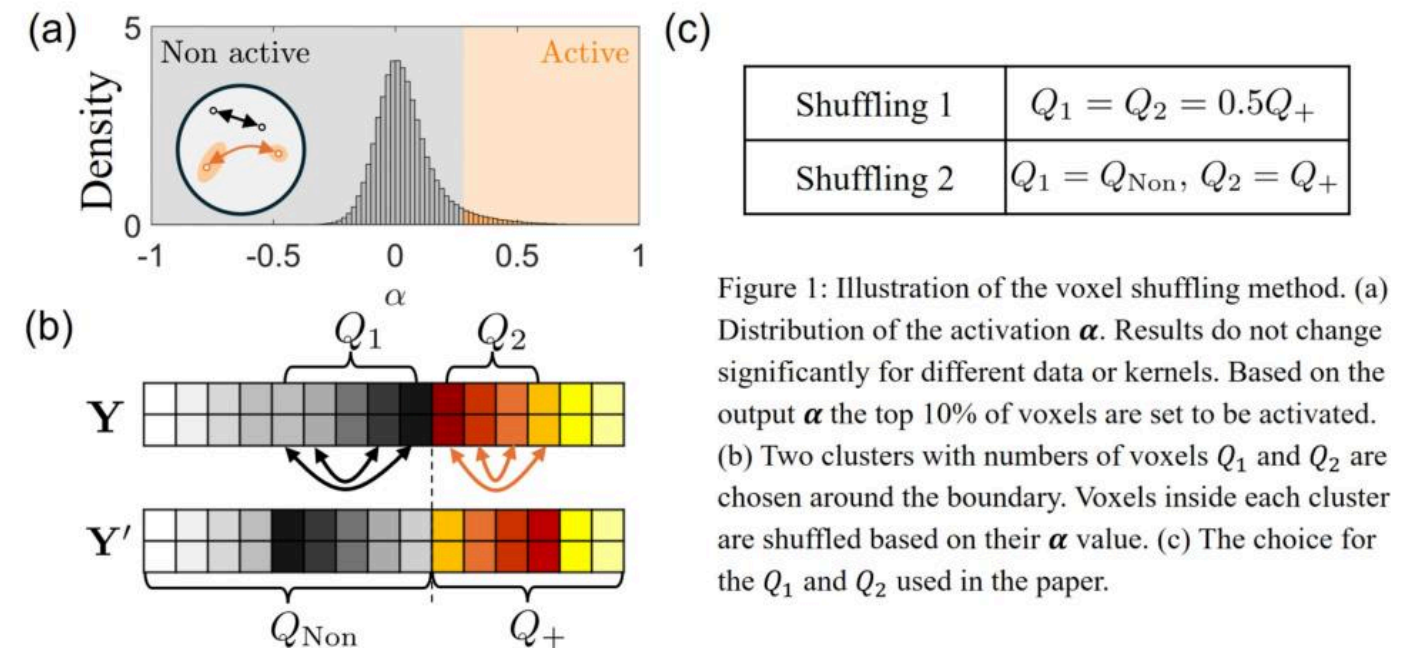
Kernel-based methods are powerful for dimension reduction in fMRI analysis. Previous methods used supervised optimization (Xifra-Porxas, 2021) or an additional dataset serving as negative pairs (Yang, 2018) to determine the unknown kernel mapping, which is less efficient and hard to expand to real datasets when there is no ground truth. Inspired by self-supervised learning in image processing (Chen, 2020), we propose a data augmentation method that could generate supervisory examples by voxel shuffling. By maximizing the similarity before and after augmentation, we show our method could reduce overfitting and increase accuracy even for complicated kernels. The results

are validated using real fMRI datasets from two different sources for the activation detection problem.

Methods:

Suppose fMRI data with $Y \in \mathbb{R}^{T \times Q}$, where T indicates the time and Q is the total number of voxels, given a certain design signal $X_{\text{eff}} \in \mathbb{R}^{T \times 1}$. Kernel canonical correlation analysis (KCCA) solves this problem by mapping the original data to kernel space and maximizing the correlation (Hardoon, 2004). This process could involve unknown functions or regularization parameters. For example, the linear kernel contains one additional regularization parameter γ to reduce overfitting (Yang, 2018). Our goal is to find the augmented data Y' , which could serve to determine the unknowns.

Figure 1 shows our proposed algorithm. Starting from arbitrary kernel mapping and performing KCCA, we can divide the output α into activated and non-activated clusters; we use $Q_+ = 0.1Q$ and $Q_{\text{Non}} = 0.9Q$ to indicate the number of voxels in each cluster. Our shuffling algorithm involves switching the voxel locations inside each cluster along the decision boundary. Specifically, we select Q_1 voxels in the non-activated cluster and Q_2 voxels in the activated cluster. Within the Q_1 voxels, the voxel with the highest α is changed to the lowest α , and the same for the second highest/lowest voxels. The same shuffling is repeated for Q_2 . The detailed choice for Q_1 and Q_2 is shown in Figure 1. Practically, we use two shufflings and take the average. The data after shuffling is indicated by Y' , which is then input to the KCCA to get another activation pattern α' . The similarity is measured by treating α as ground truth, then using different thresholds to compute the Receiver Operating Characteristic (ROC) curve, with area under the curve (AUC) with False Positive Rate (FPR) smaller than 0.1 defined as similarity. We implement the surrogate optimization algorithm in MATLAB to maximize it.



All the data are minimally preprocessed using the SPM12 package (J. Ashburner, 2008), normalized to the MNI atlas (F. F. Glasser, 2013), with Gaussian smoothing FWHM=4 mm and 7 spatial orientation filters for kernel methods with the same smoothing level (Yang, 2018). We set the criteria to be Gray Matter (GM) overlapping. Using gray matter as ground truth, different thresholds for α are used to evaluate the ROC curve. The results for two different datasets are shown in Figure 2 (a) and (b). Relationship between shuffling robustness and gray matter overlapping is observed.

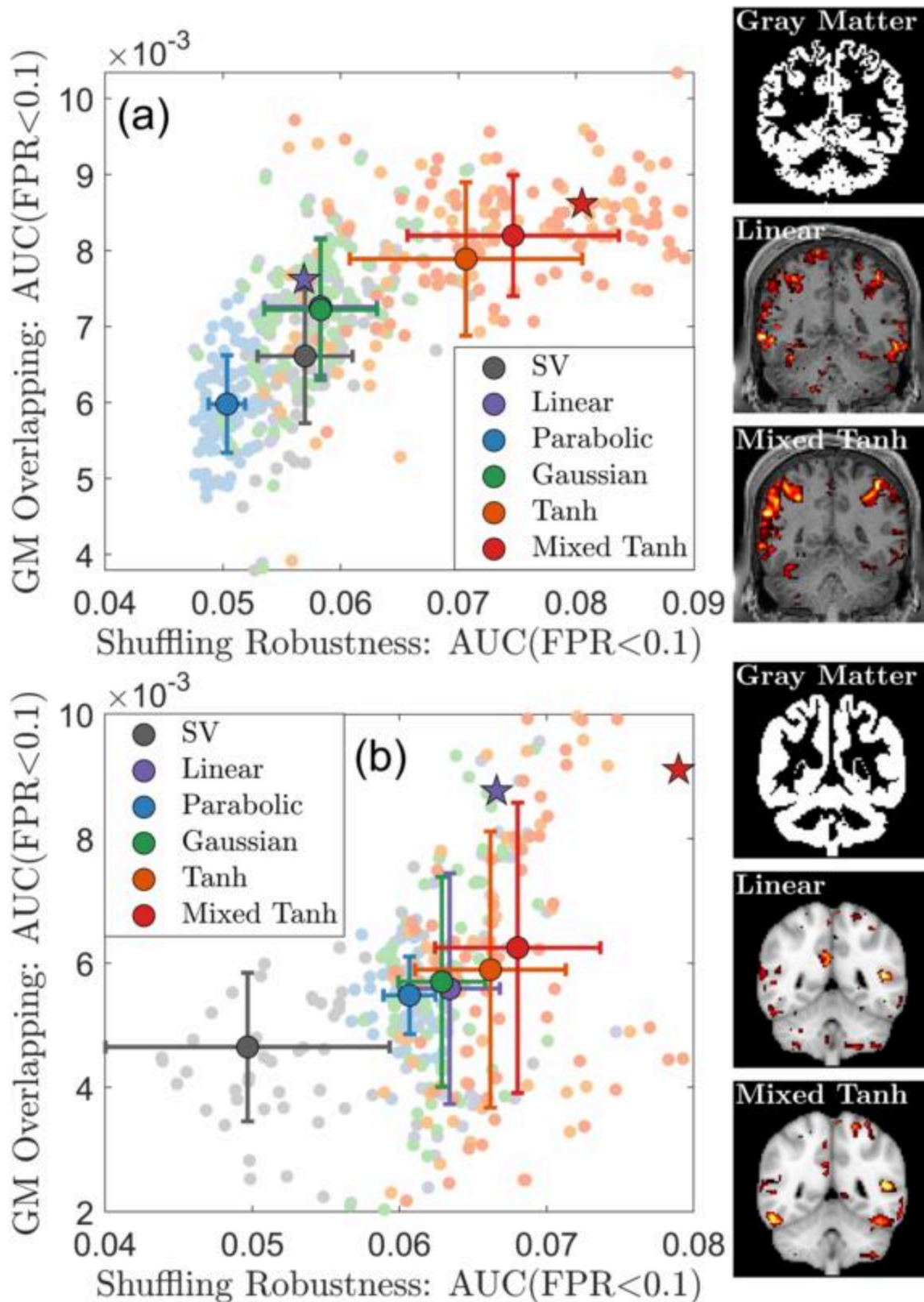


Figure 2: The relationship between shuffling robustness and gray matter overlapping for task fMRI. Each dot represents one subject analyzed using the specified methods. Panels (a) and (b) correspond to the HCP dataset and in-house scans, respectively. In each dataset, one subject was selected (indicated by a star), and the corresponding activation patterns are displayed on the right. Compared with SV, the mixed hyperbolic tangent kernel increased the AUC by 25.74% and 27.14% for HCP and in-house scans, respectively.

37.1470 for fMRI and in-house scans, respectively.

Conclusions:

The main finding of the study is that we have found a robust way to do data augmentation. We find that by maximizing the similarities before and after augmentation, we can determine the best kernel mapping without using the spatial ground truth or additional datasets.

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Modeling and Analysis Methods:

Activation (eg. BOLD task-fMRI) ¹

Methods Development

Novel Imaging Acquisition Methods:

BOLD fMRI ²

Keywords:

FUNCTIONAL MRI

^{1|2}Indicates the priority used for review

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Not applicable

Please indicate which methods were used in your research:

Functional MRI

For human MRI, what field strength scanner do you use?

3.0T

Provide references using APA citation style.

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