Application of Self-organizing Maps in Functional Magnetic Resonance Imaging

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Abstract. In the present work, we used Kohonen's self-organizing map algorithm (SOM) to analyze functional magnetic resonance imaging (fMRI) data. As a first step to increase computational efficiency in data handling by the SOM algorithm, we performed an entropy analysis on the input dataset. The resulting map allowed us to define the pattern of active voxels correlated with auditory stimulation in the data matrix. The validity of the algorithm was tested using both real and simulated data.

1 Introduction

Functional magnetic resonance imaging (fMRI) is a non-invasive tool widely used for studying the human brain in action. The fMRI has been applied to cognitive studies and also in a clinical setting to monitor tumour growth, pre-surgical mapping, and to diagnose epilepsy, Alzheimer's disease, etc [1]. The fMRI measurements are based on blood-oxygen-level-dependent correlations (BOLD) [2],[3], with hemoglobin being used as endogenous contrast agent, due to the magnetic properties of oxy-hemoglobin (diamagnetic) and deoxy-hemoglobin (paramagnetic) [4].

The BOLD signal was obtained using two experimental paradigms. The first used a blocked design, with the subject being exposed to alternating periods of stimulation and rest. The event-related paradigm, on the other hand, required that the subject performs a simple task, intercalated by long resting intervals.

A fMRI dataset consists of images in 3D space $(x \times y \times z)$, with each image point, named a voxel, changing a long time (t). Most fMRI analysis try to identify how signal related to voxels in a region of interest (ROI) vary in time and to find out whether these variations are somehow correlated with the stimulus. This analysis, however, is a computational challenge due to the low signal to noise ratio in the BOLD response and the usually large amount of data that needs to be processed. Many analytical methods have been developed to deal with this complexity, some of them were created earlier to analyze positron emission topography-generated signals (PET).

Most methods available in the literature use statistical techniques to identify active regions, including *Student*'s t test [5], crossed correlation [6], and the general linear

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In Proceedings of the 6th International Workshop on Artificial Neural Networks and Intelligent Information Processing, pages 72-80 Copyright © SciTePress model (GLM) [7]. These methods are based on the standard hemodynamic function, which models the BOLD response in the brain. Other popular methods are independent component analysis (ICA) [8], [9] and principal component analysis (PCA) [10].

Clustering techniques have also been used successfully, including K-means, fuzzy cluster and hierarchical clustering. Clustering techniques are based on the similarity observed in voxel's time series. The present study uses the Kohonen's self-organizing maps algorithm (SOM) to analyse fMRI data. The SOM [11] is a type of clustering technique which transforms a signal input pattern of an arbitrary dimension into a discrete map and implements transformations in a topologically organized way.

2 Material and Methods

2.1 Simulated Data

We simulated the fMRI experiment (64×64) depicted in Figure 1 with 120 slices by convoluting a block-like stimulus function with the canonical hemodynamic response function generated as a sum of two distribution functions [12]:

$$h(t) = \left(\frac{t}{d}\right)^a \exp\left(\frac{-(t-d_1)}{b}\right) - c\left(\frac{t}{d'}\right)^{a'} \exp\left(\frac{-(t-d')}{b'}\right),\tag{1}$$

with a = 6, a' = 12, b = b' = 0.9 and c = 0.35, having been determined experimentally by Glover in 1999 with auditory stimulation [13].



Fig. 1. Diagram showing the spatial distribution of active voxels and their intensity along time in simulated data.

The active area corresponds to 49 voxels, while 1349 voxels corresponded to the remaining grey matter. The other 2698 voxels corresponded to the background and are not time modulated. We added uniform Gaussian noise to reach a SNR of 2dB, which was calculated with the following expression:

$$SNR = 10 \log\left(\frac{\sigma_S^2}{\sigma_R^2}\right),\tag{2}$$

where $\sigma_S^2 2$ and σ_R^2 are signal and noise variances, respectively.

2.2 Real Data

The fMRI experiment used a 1.5 T Siemens scanner (Magnetom Vision, Erlangen, Germany), with the following parameters for EPI (echo-planar imaging) sequences: TE = 60 ms, TR = 4.6 s, FA = 90, FOV = 220 mm, and slice thickness of 6.25 mm. 64 cerebral volumes with 16 slices each were acquired with a matrix dimension of 128×128 .

During the experimental procedure the subject received auditory stimulation in a blocked design, with 5 stimulation blocks (27.5s each) intercalated with 6 resting blocks (27.5 s each). During the task, the subject listened passively to a complex story with a standard narrative structure. After, the test the subject had to inform to the experimenter its comprehension of the story content.

Acquired images were preprocessed with the software SPM8 (Statistical Parametric Mapping) in order to increase the signal-to-noise ratio (SNR) and to eliminate incident noise associated with the hardware, involuntary movements of the head, cardiac and respiratory rhythms, etc.

2.3 Self-organizing Maps

fMRI data was analyzed with Kohonen's SOMs using an implementation available in the literature (see [14], [15], [16], [17], [18], [19]). Kohonen's SOM is an artificial neural network where neurons are disposed as a uni- or bi-dimensional grid layout. In a bi-dimensional layout, geometriy is free and can be rectangular, hexagonal, triangular etc. In a SOM, each neuron in a grid is represented by a probability distribution function of the input data.

The SOM algorithm responsible for map formation begins initializing the grid neurons weights with random values, which can be obtained from the input data. In the present work we used a bidimensional grid of dimension 10×10 (i = 100) [19]. Each neuron in the grid is connected to every element of the input dataset, i.e., the dimension of weights \mathbf{m}_i is the same as the input dataset:

$$\mathbf{m}_i = [m_{i1}, m_{i2}, \dots, m_{in}]^T \in \Re, \tag{3}$$

where n indicates the total amount of points available in the time series generated by the fMRI experiment.

After each iteration t of the ANN, we selected randomly a vector from the input dataset, given by:

$$\mathbf{x}_i = [x_1, x_2, \dots, x_n]^T \in \Re,\tag{4}$$

which indicates the time series of a given voxel from the fMRI dataset.

Then, x is compared to weights \mathbf{m}_i in the grid using the minimun euclidean distance as criteriom for choosing a winner in the ANN [15],[19]. Since the correlation distance metric, however, seems to be a better method to discern similarities than conventional Euclidean distance [16], the winner neuron is selected by:

$$\mathbf{m}_{c} = \arg \max \left\{ \operatorname{corr}(\mathbf{x}(t), \mathbf{m}_{i}(t)) \right\},$$
(5)

with i = 1..., M where M is the total number of neurons in the grid, $\mathbf{m}_c(t)$ represents the time series of the winner c and $\operatorname{corr}(\mathbf{x}(t), \mathbf{m}_i(t))$ is the correlation coefficient between $\mathbf{x}(t)$ and $\mathbf{m}_i(t)$.

The updating of the weight vector $\mathbf{m}(t+1)$ in time t+1, with t = 0, 1, 2, ... is defined by:

$$\mathbf{m}_{i}(t+1) = \mathbf{m}_{i}(t) + h_{ci}(t)[\mathbf{x}(t) - \mathbf{m}_{i}(t)],$$
(6)

which is applied to every neuron on the grid that is within the topological neighborhoodkernel h_{ci} from the winner. Thus, Equation (6) has the goal of approximating the weight vector \mathbf{m}_i of neuron *i* towards the input vector, following the degree of interation h_{ci} . This approach transforms the grid, after training, in a topologically organized characteristic map, in the sense that adjacent neurons tend to have similar weights.

A function frequently used to represent the topological neighborhood-kernel h_{ci} is the Gaussian function, which is defined by:

$$h_{ci}(t) = \alpha(t) \exp\left\{\frac{-\|\mathbf{r}_c - \mathbf{r}_i\|}{2\sigma^2(t)}\right\},\tag{7}$$

where $\alpha(t)$ is the learning rate, which has to gradually decrease along time to avoid that new data gathered after a long training session could compromise the knowledge already sedimented in the ANN; \mathbf{r}_c and \mathbf{r}_i determine the discrete position of neurons c and i in the grid; and $\sigma(t)$ defines the full-width at half-maximum (FWHM) of the Gaussian kernel. Parameters $\sigma(t)$ and $\alpha(t)$ gradually decrease by t/τ (τ is a time constant) after each iteration t, following an exponential decay.

2.4 Evaluating the SOM Quality

There are several mechanisms that can evaluate the quality of the generated map obtained after the learning process. In the present work we used the quantization error:

$$E_q = \frac{1}{N} \sum \|\mathbf{x} - \mathbf{m}_c\|^2.$$
(8)

The quantization error is defined as the mean error corresponding to the difference between each characteristic vector \mathbf{x} and the winner neuron \mathbf{m}_c , where N is the total number of patterns.

2.5 Analysis of Entropy

Some authors recommend the ad-hoc reduction of voxel samples to optimize the algorithm implementation [20]. Thus, in order to improve analitical efficiency, only signals originating from the brain were actually processed. Besides, we performed an entropy analysis to each voxel of the characteristics set and eliminated all voxels with an entropy level below an empirically determined threshold. The Shannon's entropy, as well as other techniques based on Information Theory, has proved to be satisfactory in fMRI experiments [21].

The Shannon entropy of a random variable **X** with probability vector (p_1, \ldots, p_n) is defined by:

$$H(\mathbf{X}) = -\sum_{i=1}^{n} p_i \cdot \log_2 p_i,\tag{9}$$

with $H(\mathbf{X})$ being the entropy of variable \mathbf{X} . The Shannon entropy [22] measures the uncertainty present in any dataset and allows the comparison of its properties with other datasets of similar dimensions, by representing the amount of information contained in each as a probabilistic event.

To calculate the entropy, the time series of each voxel is divided into two levels of intensity, then is calculated the probabilities of levels of intensity from the amount of time points at each level. Finally, the entropy of each time series is calculated according to Equation (9). The entropy of signals corresponding to non-active voxels tends to have low value because of an irregular configuration of the signal. On the other hand, the signal of a probable active voxel tends to present a high value of entropy, which is associated with a wide distribution of probability.

3 Results

In this work, the configuration parameters of the SOM were initialized according to previous studies [16],[19], both in real and simulated data. In Equation (7), the learning rate was initialized as $\alpha(0) = 0.1$ and the parameter effective width as $\sigma(t) = 7$. The number of SOM iterations was regulated dinamically according to error stabilization (Eq. 8).

3.1 Simulated Data

First, we calculate Shannon's entropy for each voxel of the simulated data. The values of entropy for the 1398 voxels contained in the interior of the artificial brain had varied of 0.4949 to 1, where the 49 voxels with activation signal had presented one high value of entropy (Figure 2a). After that, about 6% of 1398 voxels were eliminated from the input data of SOM, these voxels had presented a value of entropy H < 0.85 (Figure 2b).



Fig. 2. (a) Entropic map of the artificial data; (b) Entropic analysis of synthetic data. The dark dots in the image were eliminated, the equivalent of 6% of the input data.

From the final conformation of the neuronal grid achieved after 100 algorithm iterations (Figure 3), we can observe that voxels with similar temporal patterns are clustered together in the SOM.

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Fig. 3. 10×10 grid of neurons after implementation of the SOM algorithm, the cluster of neurons in yellow match the patterns of activity.

For better visualization of these clusters, there are several clustering methods that can be used, such as K-means [23], fuzzy logic [24] and correlation as a measure of similarity in a hierarchical clustering [16]. We use this last reference utilizing a simple correlation as a measure of similarity between neurons in the grid.

Figure 4 shows the active regions defined using the average signal from the neurons demarcated in the Figure 3. A correlation coefficient (CC) was determined between this average and each voxel in a fMRI dataset, showing only those with CC > 0.7056.



Fig. 4. The dark regions in the brain correspond to active regions as defined by the SOM after 100 iterations.

3.2 Real Data

The same analytical procedure used for simulated data was used to deal with real data. However, we adopted the quantization error (Eq. 8) to estimate the amount of steps of the algorithm and also act as quality controller of learning. Figure 5 shows the evolution of error to each 10 iterations, using normalized data. But even if the error has begun to stabilize at around 100 repetitions, the training is continued until 250 iterations in order to perform a fine tuning of the map features and thus produce a statistically accurate quantization of the input space. Analyzing the same figure, it is possible verify that the magnitude of error for the case where it was applied to analyze the entropic prior to SOM (Fig. 5a) presents lower, also have begun to stabilize somewhat earlier than the case where not used the Shannon's entropy (Fig. 5b).



Fig. 5. Graph of the quantization error for a total of 250 iterations. (a) quantization error with the application of entropy; (b) quantization error without entropy.

Figure 6 reveals the active voxels, according to our method, in the eighth slice of fMRI data CC > 0.6. In it you can see two main regions as a result of the auditory task located in the temporal lobe.



Fig. 6. Active regions correlated with the auditory stimulation defined after 250 iterations with the SOM.

4 Conclusions

The Kohonen's self-organizing map was applied in data of functional magnetic resonance in synthetic and real models, this last one representing an auditory experiment with the paradigm in block. With the purpose of increasing the efficiency of the analysis method was proposed to Shannon's entropy, which eliminated a range of 5 - 10% of the set of input data. The configuration of the data after the entropy analysis allowed more likely to find groups of neurons active in the SOM grid with a smaller number of iterations. Moreover, in the temporal evolution of the quantization error of the SOM, it can be verified that entropy analysis decreased the amplitude of this error and admitted his slightly faster stabilization. The results of SOM, both for simulated data, as for real data, reaffirmed that it can be used as a tool for interpretation of fMRI data. And it has the advantage that the shaped of the hemodynamic response is not considered, that is, a HRF modeled mathematically is not used.

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