Abstract

Recently machine learning methods had gain lots of publicity among researchers in order to analyze the brain images such as functional Magnetic Resonance Imaging (fMRI) to obtain a better understanding of the brain and its related disease such as Alzheimer's disease. Different methods have been deployed in order to discriminate Alzheimer's disease from normal ones which is a hard task, especially in early stages (eMCI) case. The majority of deployed techniques rely on constructing the functional connectivity (FC) for each person and use the vectorized FC as the input for the classifiers which has two main drawbacks: 1) The need for constructing the FC 2) The loss of possible valuable structural information in the vectorization step. Considering these problems and based on multidimensional nature the data, we have came up with a novel framework which omits the FC construction part and preserve the structural integrity of data for the classification. The proposed framework uses the High Order Singular Value Decomposition (HOSVD) in order to prune the classes and select the proper basis for each of them. This framework also allows us to obtain a general FC pattern for normal and eMCI classes but not a single sample which helps us to shed more lights on the brain abnormalities in the Alzheimers disease at its early stages. Extensive experiments using the ADNI dataset demonstrate that our proposed framework effectively boosts the fMRI classification performance and reveals novel connectivity patterns in Alzheimer's disease at its early stages.

Keywords:

1. Introduction

Alzheimers disease (AD) is a progressive neurodegenerative disorder with a long pre-morbid asymptomatic period which affects millions of elderly individuals worldwide[1]. It is predicted that the number of affected people will double in the next 20 years, and 1 in 85 people will be affected by 2050 [2]. The predominant clinical symptoms of AD include a decline in some

important brain cognitive and intellectual abilities, such as memory, thinking, and reasoning. Precise diagnosis of AD, especially at its early warning stage: early Mild Cognitive Impairment (eMCI), enables treatments to delay or even avoid such disorders [3].

In recent years, brain imaging techniques like Positron Emission Tomography (PET)[15], Electroencephalography (EEG)[16] and functional Magnetic Resonance Imaging (fMRI)[17] have been used in analysis of AD. Due to the high spatial resolution and relatively lower costs, fMRI is vastly used among researchers in order to monitor brain activities especially in AD and all its stages in which detecting abnormalities within small brain regions is essential [4]. An fMRI sample is naturally a 4D tensor consisting of 3D voxels moving in time, and each voxel contains an intensity value that is proportional to the strength of the Blood Oxygenation Level Dependent (BOLD) signal, which is a measure of the changes in blood flow, to estimate the activity of different brain regions[5]. Resting-state fMRI(rs-fMRI) is an fMRI technique in which the patient is asked to rest during the whole scan, focuses on the low-frequency (< 0.1Hz) oscillations of BOLD signal, which presents the underlying neuronal activation patterns of brain regions[8][10]. rs-fMRI is usually used in order to analyze brain diseases like AD or autism[27, 28].

Since each fMRI volume consist of hundreds of thousands of voxels which are often highly correlated with the surrounding voxels in the brain volume, parcellation of the brain for further analysis has moved toward the use of anatomical atlases. These atlases are strictly defined using anatomical features of the brain, like locations of common gyri and do not rely on any functional information. To generate data using an Atlas based approach, the BOLD signal from all voxels is averaged within each brain region called Region of Interest(ROI)[6]. By putting together the average time-series for all the ROIs, the *i*th volume would become $X_i \in \mathbb{R}^{T \times R}, i = \{1, 2, \dots, S\}$ in which R, T and S are the number of ROIs, time points and samples respectively. The process of obtaining such a matrix is shown in Figure 1.

There are two major studies associated with rs-fMRI data: finding common brain disorders caused by diseases like Alzheimer's or autism, and more recently detecting patients with brain disorders using classification techniques [29, 30]. Due to the high dimensionality of data and the nature of diseases like eMCI which does not show any reliable clinical symptoms, researchers moved towards advanced machine learning techniques in order to achieve more reliable analysis [31].

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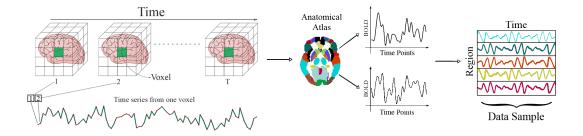


Figure 1: The process of extracting ROI time-series from the original 4D volume.

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A powerful tool that is commonly used in order to achieve aforementioned goals is Functional Connectivity (FC) network. FC is a region \times region matrix X in which \bar{x}_{ij} represents the functional connectivity between the ith and jth ROI. Functional connectivity is an observable phenomenon quantifiable with measures of statistical dependencies, such as correlations, coherence, or transfer entropy [32]. Recent studies have shown that some brain disorders like AD could alter the way that some brain regions interact with each other. For example, compared with the healthy, AD patients have been found decreased functional connectivity between the hippocampus and other brain regions, and MCI patients have been observed increased functional connectivity between the frontal lobe and other brain regions??. So, Finding an FC that highlights the patterns caused by a disease has been a common goal in the rs-fMRI study for a long time. Several approaches exist to find common patterns among different brain scans. Data-driven methods such as kernel-PCA or clustering techniques have been proposed for this taskref?. But ultimately most of them rely on calculating a network for each volume. This may overlook the role of noises or outliers within the data.

In recent years FCs are also used as features in classification. So, instead of using X_i as the i^{th} sample its corresponding FC i.e. \bar{X}_i is used as a feature. Although FCs show promising results, they bring their own challenges. The computational cost of FC usually is high and also its quality has a main effect in the quality of learning process. Also, Since the conventional

classifiers like SVM work on data in vector format, these matrix features should be vectorized in order to feed them to classifiers. This vectorization leads to high-dimensional vectors which produce poor performance due to the phenomena known as the Curse of Dimensionality. Alongside the curse of dimensionality, vectorization also destroys potential information that are embedded in the structure of data. This problem has been studied especially in image data in which vectorization destroys the spatial relations within an image. Rezghi and S.Ahmadi paper

In this paper, based on high order tensor decomposition, we have created a framework in which the mentioned goals i.e. finding an FC for a whole class and detecting a disorder via classification could be achieved via a single High Order Singular Value Decomposition (HOSVD) of each class. Here based on latent variables obtained by HOSVD a general representative pattern of FC for eMCI and normal are obtained. Experiments show that the functional connectivities by the proposed method are confirmed by empirical methods along with novel connectivities. Also, The proposed classifier also outperforms state of the art eMCI classification methods. Viewing each class as a tensor allows us to work with time and region features separately but simultaneously. This multilinear view ables us to design a proper dimension reduction relative to the nature of each feature along with a discriminant function based on linear regression on latent space of samples that uses the test data to enhance the quality of the training set without forcing any a prior knowledge to the classifier, a task which is not possible through well known classifiers like SVM, logistic regression or k-NN. It is also notable that the proposed discriminant function directly works with the X_i s as features. Having the FC calculation step omitted not only heavily affects the computational performance of the method, but it also saves us from the troubles of FCs mentioned before.

To verify our approach, we conduct an extensive experimental study on rs-fMRI data from the benchmark dataset ADNI ¹ As will be seen, the results well demonstrate the effectiveness and advantages of our method. Specifically, the proposed framework, not only grants us superior classification accuracy to that from other methods, it is also much faster and more stable against different data selection schemes. We have also confirmed our achieved whole-class FC matrix using empirical data on the eMCI and Normal func-

¹http://adni.loni.usc.edu/

tional connectivity patterns.

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3. Bibliography

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