Brain Autism Segmentation Using Convolutional Neural Networks In MRI Images Interface

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Abstract- *Functional magnetic resonance imaging or functional AUTISM (AUTISM) is a functional neuro imaging procedure using autism technology that measures brain activity by detecting associated changes in blood flow, uses the Blood-oxygen-level dependent (BOLD) contrast. Current results from neuroscience suggest a modular organization of the brain. To understand the complex interaction patterns among brain regions. In proposed system uses CNN algorithm, an efficient algorithm for partitioning segmentation. A brain region is defined as a set of subjects sharing a similar interaction pattern among their brain regions. An extensive experimental evaluation on benchmark data demonstrates the effectiveness and efficiency of our approach. The results on two real autism studies demonstrate the potential of riemanian approach to contribute to a better understanding of normal brain function and the alternations characteristic for psychiatric disorders it means that Mental disorders are generally defined by a combination of how a person feels, acts, thinks or perceives. This may be associated with particular regions or functions of the brain or rest of the nervous system, often in a social context.*

Keywords- Autism Spectrum Disorder, MRI images, Convolutional Neural Networks, Segmentation

I. INTRODUCTION

Medical imaging is the technique and procedure of creating visual demonstration of the internal of a body for experimental analysis and health intervention. Medical imaging seeks out to disclose internal structures hidden by the skin and bones, as well as to diagnose and treat disease. Medical imaging also establishes a database of normal anatomy and physiology to make it possible to identify abnormality. Although imaging of removed organs and tissues can be performed for medical reasons, such procedures are usually considered part of pathology instead of medical imaging.

II. LITERATURE REVIEW

In this paper proposed a method for clustering of time series based on their structural characteristics. Unlike other alternatives, their proposed method does not cluster point values using a distance metric, rather it clusters based on global features extracted from the time series. The feature measures are obtained from each individual series and can be fed into random clustering algorithms, including an unsupervised neural network algorithm, self-organizing map, or hierarchal clustering algorithm. Global measures describing the time series are obtained by applying statistical operations that best capture the underlying uniqueness: trend, seasonality, periodicity, serial correlation, skewness, kurtosis, chaos, nonlinearity, and self-similarity. Since the method clusters using extracted global measures, it reduces the dimensionality of the time series and is much less sensitive to missing or noisy data. They further provide a search mechanism to find the best selection from the feature set that should be used as the clustering inputs. Their technique has been tested using benchmark time series datasets formerly reported for time series clustering and a set of time series datasets with known distinctiveness.

The empirical results show that their approach is able to yield meaningful clusters. The resulting clusters are comparable to those produced by other methods, but with some promising and interesting variations that can be instinctively explained with knowledge of the global characteristics of the time series. Hirano et al. in proposed an algorithm for clustering the time series medical data. Their paper presents a cluster analysis method for multidimensional time-series data on clinical laboratory examinations. Their method represents the time series of test results as trajectories in multidimensional space, and compares their structural similarity by using the multiscale comparison technique. It enables us to find the part-topart correspondences between two trajectories, taking into account the relationships between different tests.

The resultant distinction can be further used with clustering algorithms for finding the groups of similar cases. The method was applied to the cluster analysis of Albumin-Platelet data in the chronic hepatitis dataset. The experimental results demonstrated that it could form interesting groups of cases that have high correspondence to the fibrotic statistics.

III. EXISTING SYSTEM

Human brain activity is very complex and far from being fully understood. Many psychiatric disorders like Schizophrenia and Somatoform Pain Disorder can so far neither be identified by biomarkers, nor by physiological or histological abnormalities of the brain. Aberrant brain activity often is the only resource to understand psychiatric disorders. Functional magnetic resonance imaging (autism) opens up the opportunity to study human brain function in a noninvasive way. The basic signal of autism relies on the blood-oxygen-level dependent (BOLD) effect, which allows indirectly imaging brain activity by changes in the blood flow related to the energy consumption of brain cells. In a typical autism experiment, the subject performs some cognitive task while in the scanner. . Recently, resting state autism has attracted considerable attention in the neuroscience community. Interaction K-means (IKM) simultaneously Clusters the data and discovers the relevant cluster specific interaction patterns. The algorithm IKM is a general technique for clustering multivariate time series and not limited to autism data. Besides autism, multivariate time series are prevalent in many other applications. Increasing amounts of motion stream data are collected in multimedia applications. Gesture sensing devices, such as a Cyber Glove usually contain multiple sensors to capture human movements. Human motion stream data can also be extracted from video streams. In this application, it makes sense to regard each movement as a data object. A cluster analysis of motion stream data potentially identifies. Clusters with similar movements, usually performed by different person.

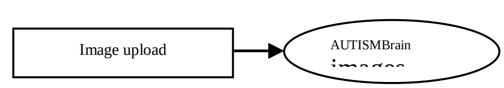
IV. PROPOSED SYSTEM

Human brain activity is very complex and far from being fully understood. Many psychiatric disorders like Schizophrenia and Somatoform Pain Disorder can so far neither be identified by biomarkers, nor by physiological or histological abnormalities of the brain. Aberrant brain activity often is the only resource to understand psychiatric disorders. Functional magnetic resonance imaging (Autism) opens up the opportunity to study human brain function in a noninvasive way. The basic signal of autism relies on the blood-oxygen-level-dependent (BOLD) effect, which allows indirectly imaging brain activity by changes in the blood flow related to the energy consumption of brain cells. Brain tissues are segmented using Conditional Random field approach. It presents a new CNN for segmenting SWI venography datasets. The CNN model aggregates multiple first- and second-order potentials. Specifically, appearance, shape, location, auto-logistic (Ising) interaction and data dependent interaction potentials are combined to produce robust, complete and fully automated SWI genogram segmentation. In a typical autism experiment, the subject to perform some cognitive task while in the scanner. And implement statistical region merging approach to group the similar regions. It is the reconstruction of regions on the observed image, based on an unknown theoretical (true) image on which the true regions we seek are statistical regions whose borders are defined from a simple axiom. Second, we show the existence of a particular blend of statistics and algorithmic to process observed images generated with this model, by region merging, with two statistical properties. With high probability, the algorithm suffers only one source of error for image segmentation: over merging, that is, the fact that some observed region may contain more than one true region. The algorithm does not suffer neither under merging, nor the most frequent hybrid cases where observed regions may partially span several true regions.

4.1 BRAIN IMAGE ACQUISITION

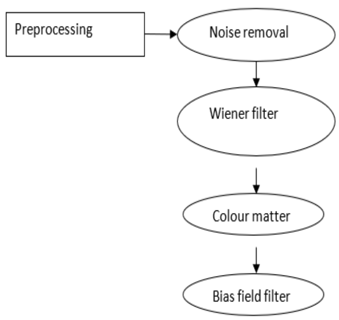
Functional magnetic resonance imaging or functional autism (AUTISM) is a functional neuroimaging procedure using Autism technology that measures brain activity by detecting changes associated with blood flow. This technique relies on the fact that cerebral blood flow and neuronal activation are coupled. When an area of the brain is in use, blood flow to that region also increases. [Citation needed] The primary form of autism uses the blood-oxygen-level dependent (BOLD) contrast, discovered by Seiji Ogawa. This is a type of specialized brain and body scan used to map neural activity in the brain or spinal cord of humans or other animals by imaging the change in blood flow (hemodynamic response) related to energy use by brain cells. Since the early 1990s, autism has come to dominate brain mapping research because it does not require people to undergo shots, surgery, or to ingest substances, or be exposed to ionizing radiation, etc. Other methods of obtaining contrast are arterial spin labeling and diffusion autism.

Upload image:



4.2 PREPROCESSING

In this module we convert the RGB image into gray scale images. The colors of leaves are always green shades and the variety of changes in atmosphere because the color features having low reliability. Therefore, to recognize various plants using their leaves, the obtained leaf image in RGB format will be converted to gray scale beforepre-processing



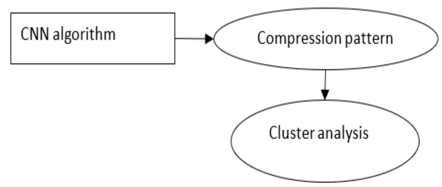
4.3 BIAS FIELD FILTER

Bias field signal is a low-frequency and very smooth signal that corrupts AUTISM images specially those produced by old AUTISM (Magnetic Resonance Imaging) machines. Image pro- cessing algorithms such as segmentation, texture analysis or classification that use the graylevel values of image pixels will not produce satisfactory results. A pre-processing step is needed to correct for the bias field signal before submitting corrupted AUTISM images to such algorithms or the algorithms should be modified. In this report we discuss two approaches to deal with bias field corruption. The first approach can be used as a preprocessing step where the corrupted autism image is restored by dividing it by an estimated bias field signal using a surface fitting approach. The second approach shows how to modify the fuzzy c-means algorithm so that it can be used to segment an autism image corrupted by a bias field signal.

4.4 CNN

In machine learning, a network (CNN, or ConvNet) is a class of deep, feed-forwardartificial neural networks, most commonly applied to analyzing visual imagery. CNNs use a variation of multilayer perceptrons designed to require minimal preprocessing. They are also known as shift invariant or space invariant artificial neural networks (SIANN), based on their shared-weights architecture and translation invariance characteristics.

Convolutional networks were inspired by biological processes in that the connectivity pattern between neurons resembles the organization of the animal visual cortex. Individual cortical neurons respond to stimuli only in a restricted region of the visual field known as the receptive field. The receptive fields of different neurons partially overlap such that they cover the entire visual field .CNNs use relatively little pre-processing compared to other image classification algorithms. This means that the network learns the filters that in traditional algorithms were hand-engineered. This independence from prior knowledge and human effort in feature design is a major advantage.



V. RESULT ANALYSIS

It can evaluate the performance to analyze the effectiveness of our proposed algorithm Accuracy metric is used to evaluate the performance of the system. It can be measured using truly classified pixels

Methods	ACCURACY
CNN	87.30%
FCM	64.22%

VI. CONCLUSION

In this project implement, a novel segmentation notion for multivariate time series. We define a segment as asset of objects sharing a specific interaction pattern among the dimensions. In addition, to propose CNN, an efficient algorithm for interaction-based segmentation. Our experimental evaluation demonstrates that the interaction-based cluster notion is a valuable complement to existing methods for clustering multivariate time series. CNN approach achieves good results on synthetic data and on real world data from various domains, but it cluster only the multivariate time series only so lot of in information loss and it not efficient to cluster the brain images especially results on EEG and autism data. So we propose to consider different models for different regions of the time series using statistical region clustering. We intend to work on methods for suitable initialization of CNN since existing strategies for K-means cannot be straightforwardly transferred to CNN because of the special cluster notion. We are also investigating in feature selection for interaction-based segmentation. From segmentation with statistical region clustering can cluster the different brain region for gathering the more information. Here the statistical region clustering, it can be consider the different regions of brain and then cluster different brain region, and merging the cluster image with database image finally provide the result about the human being is normal or abnormal. Then provide diseases in brain images with improved accuracy rate.

VII. FUTURE WORK

In future work we can extend our work to implement this approach in advance 3D images and 4D images with improved various type of segmentation algorithms to predict the neuron states of brain images. Autism image of autism diseases in brain images with improved accuracy rate and easily find out the disease and predict.

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