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Tumor Detection and Classification of MRI Brain Images using SVM and DNN

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ABSTRACT

The brain is one of the most complex organ in the human body that works with billions of cells. A cerebral tumor occurs when there is an uncontrolled division of cells that form an abnormal group of cells around or within the brain. This cell group can affect the normal functioning of brain activity and can destroy healthy cells. Brain tumors are classified as benign or low-grade (Grade 1 and 2) and malignant tumors or high-grade (Grade 3 and 4). The proposed methodology aims to differentiate between normal brain and tumor brain (Benign or Melign). The proposed method in this paper is automated framework for differentiate between normal brain and tumor brain. Then our method is used to predict the diseases accurately. Then these methods are used to predict the disease is affected or not by using a comparison method. These methodology are validated by a comprehensive set of comparison against competing and well established image registration methods, by using real medical data sets and classic measures typically employed as a benchmark by the medical imaging community our proposed method is mostly used in medical field. It is used to easily detect the diseases. We demonstrate the accuracy and effectiveness of the preset framework throughout a comprehensive set of qualitative comparisons against several influential state-of-the-art methods on various brain image databases.

***KEYWORDS:** Benign, Melign, Acquisition, Pre-processing, Segmentation, Classification, Extraction*

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I. INRODUCTION

MRI is an extensively used technique which facilitates the diagnosis and prognosis of brain tumors in many neurological diseases and conditions. Standard MRI sequences are generally used to differentiate between different types of brain tumors based on visual qualities and contrast texture analysis of the soft tissue. More than 120 classes of brain tumors are known to be classified in four levels according to the level malignancy by the World Health Organization (WHO). The grading from low to high (1-4) are malignant levels from the least aggressive biological tumor to the most aggressive tumors, as shown by histological criteria, for example, vascularity, invasiveness, and tumor growth rate. Gliomas are the most primary cerebral tumor and a pretreatment evaluation grade is necessary; however, the exclusive use of standard MRI sequences may be insufficient for a precise diagnosis. As the support vector machines architectures are becoming more mature, they gradually outperform previous state-of-the-art classical machine learning algorithms. The literature survey has revealed that some of the techniques are invented to obtain segmentation only; some of the techniques are invented to obtain feature extraction and some of the techniques are invented to obtain classification only.

Tumor is an uncontrolled growth of cancer cells in any part of the body. Tumors are of different types and have different characteristics and different treatments. At present, the brain tumors are classified as: primary brain tumors and metastatic brain tumors. The former begin in the brain and tend to stay in the brain, the latter begin as a cancer elsewhere in the body and spreading to the brain. Brain tumors are divided into two types: benign and malignant. In fact, the most widely used grading scheme has been issued by the World Health Organization (WHO). It classifies brain tumors into grade I to IV under the microscope. In general, grade I and grade II are benign brain tumor (low-grade); grade III and grade IV are malignant brain tumor (high-grade). Usually, if low-grade brain tumor is not treated, it is likely to deteriorate to high-grade brain tumor.

Therefore, brain tumor are seriously endangering people's lives and early discovery and treatment have become a necessity. Along with the advance of medical imaging, imaging modalities play an important role in the evaluation of patients with brain tumors and have a significant impact on patient care. Recent years, the emerging new imaging modalities, such as XRay, Ultrasonography, Computed Tomography (CT), Magneto Encephalo Graphy (MEG), Electro Encephalo Graphy (EEG), Positron Emission Tomography (PET), Single-Photon Emission Computed Tomography (SPECT), and Magnetic Resonance Imaging (MRI), not only show the detailed and complete aspects of brain tumors, but also improve clinical doctors to study the mechanism of brain tumors at the aim of better treatment. Clinical doctors play an important role in brain tumor assessment and therapy. This information is very important and critical in deciding between the different forms of therapy such as surgery, radiation, and chemotherapy. Therefore, the evaluation of brain tumors with imaging modalities is now one of the key issues of radiology departments.

II. PROPOSED SYSTEM

In this project we discuss about the brain tumors disease detection by using a method at MRI images. Nowadays many peoples are affected at more Brain cell tumor disease. So using that situation many doctors and hospitals are easily theft more money from patients. The common peoples are mostly affected by this problem. We propose the method is used to predict the disease accurately. Then it detect all the Brain tumor disease easily. That is used to detect all the brain cell related diseases and its provide more accuracy. Then the detection of brain tumor disease at less computation time. Our proposed method is used to predict all the brain cell related diseases like that Tumors. They are easily detected and its used accurately identify the diseases in less processing time.

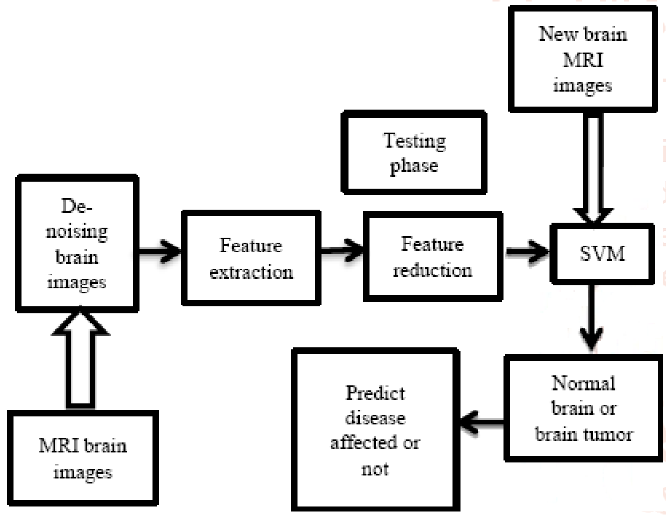

Fig1. Block Diagram

III. SOFTWARE IMPLEMENTATION

A. DATA ACQUISITION

The disease affected image or not affected images are captured from camera or to upload the images. It is used to upload the pair of input images. However, it differs from conventional methods by using thermal images as input.

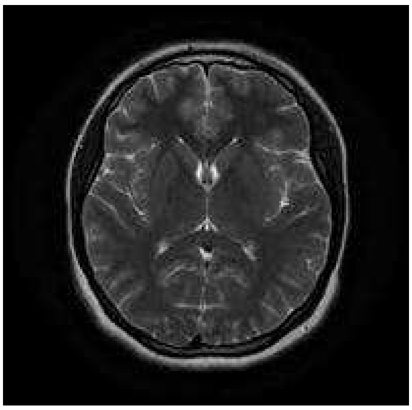

Fig2. Data acquisition

B. PRE PROCESSING

The pair of images is preprocessed it by the alignment process. The main process of the preprocessing is used to separate the meaningful features. The academic community has achieved fruitful breakthroughs in the field of brain image in the past few decades. Processing chain is de-noising medical thresholding techniques in noise presence for various wavelet families. We applied Haar, Symlet, Morlet, and Daubechies in de-noising MRI brain images. Performance estimation and analysis are accomplished using Signal to Noise Ratio (SNR), Peak Signal to Noise Ratio (PSNR) and Mean Square Error (MSE). We are using wavelets to de-noise images. In medical images, edges are places where the image brightness changes fast. Maintaining edges while de-noising an image is severely important for initiative quality. Traditional low pass filtering removes noise, it often smoothens edges and influences image quality. Wavelets are able to remove noise while maintaining important features. From the obtained results it can be confirmed that the wavelets with higher level tend to give good result. Since the SNR value corresponding to the higher level. The same goes for Symlet wavelets. While the best result are obtained using the Haar wavelets, with highest SNR value, least MSE and least entropy. It can be conclude from above results that the wavelet that are higher level show better results.

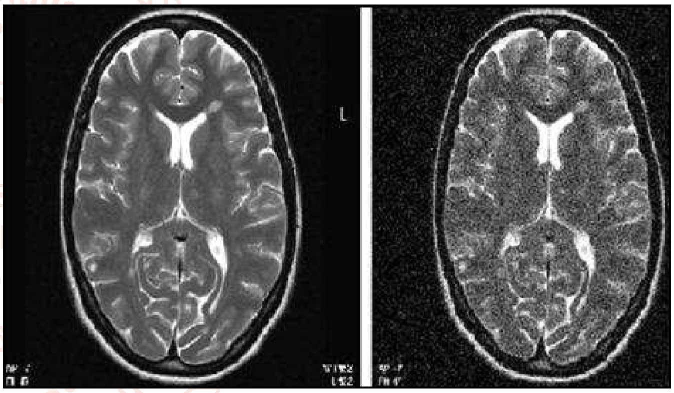

Fig3. Preprocessing

C. IMAGE SEGMENTATION

The segmentation method is described using its free parametric character and unsupervised nature of threshold choice and has the following benefits like the process is very easy but only the zeroth and first order cumulative moments of the Greylevel histogram are used. We can apply the simple extension to multi thresholding problems that is possible by the criteria on which the method is based, we can automatically selecting an optimal threshold or set of threshold, that are not based on the diffraction (i.e. a local property such as valley) but on the integration (i.e. global property) of the histogram.

We can also analyze other aspects for example evaluation of class separabality, estimation of class mean levels, etc. We are able to underline the generality of the method, it covers a large unsupervised decision procedure. That is used to automatically perform clustering- based image thresholding or reduction of grey-level image to a binary image in computer vision and image processing. The algorithm presumes that the image contains two classes of pixels following bimodal histogram, it then calculates the optimum threshold separating the two classes so that their mixed expansion is minimal, or uniformly so that their inter class variance is maximal.

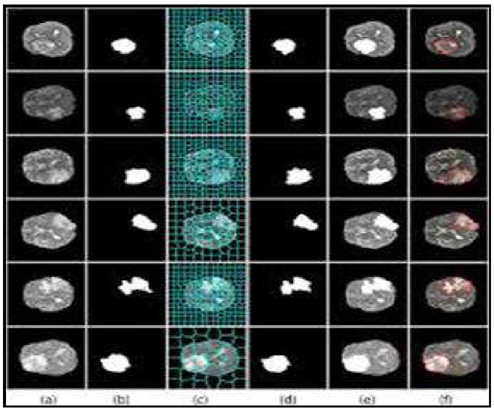

Fig4. Image Segmentation

D. IMAGE CLASSIFICATION

The classification method is used to classify the object easily so we are detecting the object accurately. Then this method is used to classify the disease accurately and refining a background objects accurately. Our data has specifically two classes we can use a support vector machine (SVM). We are using a set of new MRI brain images. Finding the best hyper plane that detaches all data points of one class those of the other class means correct data classification using SVM methods. Establishing the best hyper plane for an SVM means the one with the largest margin between the two classes.

The maximal width of the plate parallel to the hyper plane that has no interior data points determine the margin. Complicated binary classification problems do not have a simple hyper plane as a useful separating criterion. There is a variant of the mathematical approach that retains nearly all the simplicity of an SVM separating hyper plane for those problems.

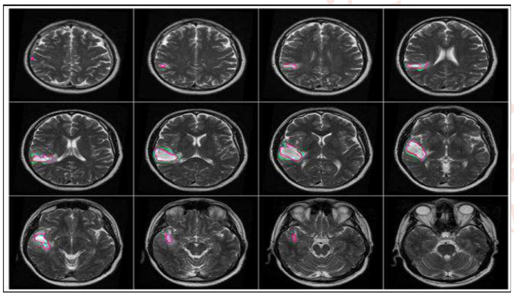

Fig4. Image Classification

E. FEATURE EXTRACTION

This method is used to predict disease accurately. This system identify from the MRI image at the brain is normal or affected. Then the useful disease prediction from the brain images then it provides the range of disease level. We remove the average pooling layers from the pre-trained model and add auxiliary convolution layers to detect large sizes of objects. It will not be surprising for getting a better result after adding a top-down structure in FG stage, but that is beyond the scope of this paper.

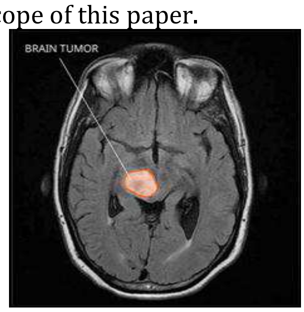

Fig5. Feature Extraction

IV. CONCLUSION

The brain tumor detection and classification system is implemented using CWT, DWT and SVMs. The result shows that SVMs having the proper sets of training data are able to distinguish between abnormal and normal tumor regions and classify them correctly as a benign tumor, malign tumor or healthy brain. In practice, SVMs have significant computational advantages Five times improvement in computation speed (22 vs. 110 ms) for our proposed method.

This classification is very important for the physician in establishing a precise diagnostic and recommending a correct further treatment. If we are interested in de-noising, compression, restoration, then DWT is often more appropriate. A hybrid approach is recommended in solving properly the detection and classification problems in brain tumors. Our method is used to predict the diseases accurately. Then these methods are used to predict the disease is affected or not affected by using a comparison method. These methodology are validated by a comprehensive set of comparisons against competing and well-established image registration methods, by using real medical datasets and classic measures typically employed as a benchmark by the medical imaging community our proposed method is mostly used in medical field. it is used to easily detect the diseases.

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