# Review on Brain Tumor Detection Using Digital Image Processing

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Abstract— Medical image processing is the most challenging and emerging field today. This paper describes the methodology of detection & extraction of brain tumor from patient's MRI scan images of the brain. In this paper, a method for segmentation of brain tumor has been developed on 2D-MRI data which allows the identification of tumor tissue with high accuracy and reproducibility compared to manual techniques.

This method incorporates with some noise removal functions, segmentation and morphological operations which are the basic concepts of image processing. Detection and extraction of tumor from MRI scan images of the brain is done by using MATLAB software. The aim of this work is to design an automated tool for brain tumor quantification using MRI image data sets.

Key words: Brain tumor, grey scale imaging, MRI, MATLAB, morphology, noise removal, segmentation.

# **1 INTRODUCTION**

Tumor is defined as the abnormal growth of the tissues [3]. Epilepsy is a brain disorder in which clusters of nerve cells, or neurons, in the brain sometimes signal abnormally [9]. Neurons normally generate electrochemical impulses that act on other neurons, glands, and muscles to produce human thoughts, feelings, and actions [6]. In epilepsy, the normal pattern of neuronal activity becomes disturbed, causing strange sensations, emotions, and behaviour or sometimes convulsions, muscle spasms, and loss of consciousness [20].

Magnetic Resonance Imaging (MRI) is an advanced medical imaging technique used to produce high quality images of the parts contained in the human body [12]. From these high-resolution images, we can derive detailed anatomical information to examine human brain development and discover abnormalities [30]. MRI consists of T1 weighted, T2 weighted and PD (proton density) weighted images and are processed by a system which integrates fuzzy based technique with multispectral analysis [1].

### 2 METHODOLOGY

The algorithm has two stages, first is pre-processing of given MRI image and after that segmentation and then morphological operations. Steps of algorithm are as following:-

- 1) Give MRI image of brain as input.
- 2) Convert it to gray scale image.
- 3) Apply high pass filter for noise removal.
- 4) Apply median filter to enhance the quality of image.
- 5) Compute threshold segmentation.
- 6) Compute watershed segmentation.

- 7) Compute morphological operation.
- 8) Finally output will be a tumor region.

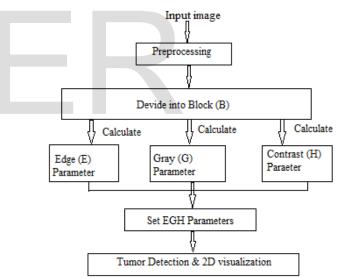


Figure 1. Proposed framework

### 2.1 The proposed framework

The process flow of our proposed methodology may be shown as figure 1.

The first section discusses how images are divided into regions using a block-based method. The second section shows how each classified block is studied individually by calculating its multiple parameter values. In this instance, the multiparameter features refer to the following three specific features: the edges (E), gray values (G), and local contrast (H) of the pixels in the block being analyzed [6].

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### 2.2 Input image

The images we got from MRI are of three types: axial Images, saggital Images, coronal Images. The number of images depends on the resolution of the movement of the MRI magnets [31]. Preprocessing

In preprocessing the first step is to load the MRI image data set on to the MATLAB workspace and after loading they will be processed. After this processing there are only three clips instead of 384 separate images, i.e. one clip for axial images, one clip for saggital images and one clip for coronal images[6].

Axial, saggital and coronal images were loaded and viewed in the MATLAB. After this, axial, saggital and coronal movie clips were prepared and are shown in figure 2.

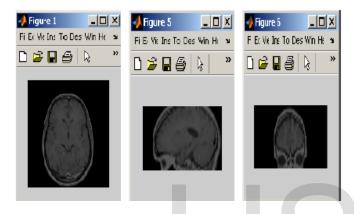


Figure 2 Axial, saggital and coronal image chips

#### 2.3 Multiparameter Calculations

Classifying regions using their multiparameter values makes the study of the regions of physiological and pathological interest easier and more definable. The multiparameter features refer to the following three specific values for the edges (E), gray values (G), and local contrast (H) of the pixels [3].

*Edge* (*E*) *Parameter:* We use sobel edge detection method to detect image edges ( $I_E$ ) is obtained by filtering an input image with two convolution kernels concomitantly, one to detect changes in vertical contrast ( $H_X$ ) and the other to detect horizontal contrast ( $H_Y$ ), shown in equation (1). Image output ( $I_E$ ) is obtained by calculating the gradient magnitude of each pixel, as shown in equation (2). Subsequently, the edge parameter (E) is calculated, whereby E ( $_{r,c}$ ) is increased by one each time when  $I_E$  (x, y) = '1' in a supervised block, as shown in equation (3). [24]

$$H_{X} = \begin{pmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{pmatrix}, H_{Y} = \begin{pmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{pmatrix}$$
(1)

$$I_{E}(x, y) = \sqrt{I_{x}^{2}(x, y) + I_{y}^{2}(x, y)}$$
(2)

$$E(r,c) = \sum_{(x,y)\in B} \left( I_{EP} = 1 \right)$$
(3)

Gray (G) Parameter: The gray parameter (G) for each block of the brain is accumulated, and controlled by a binary image ( $I_T$ ) using the GD value as a threshold, as shown in equation (6). GD value is calculated using the average pixel value ( $Iav_s$ ) of each image slice (S) for total image slices (T) of an image dataset, shown in equation (4) and (5).

$$Iav_{s} = \frac{1}{65536} \sum I_{p} \left( x, y \right) \tag{4}$$

$$G_D = \frac{1}{T} \sum_{s=0}^{T} I_{avs}$$
<sup>(5)</sup>

$$G(r,c) = \sum_{(x,y)\in B} I_p(x,y), \forall \mathbf{I}_{\mathrm{T}}(x,y) = 1$$
(6)

Contrast (H) Parameter: Contrast (H) is often used to characterize the extent of variation in pixel intensity. We adopt the minimum/maximum stretch algorithm for the 8-neighborhood connectivity, where min H and max H represent the minimum and maximum intensity values of the neighbourhood pixel  $C_8(I_H)$ , as shown in equation (7). In the previous studies, tumor cells are often associated with higher value of contrast (H) parameter (Senaratne et al., 2006).  $H_d$  is obtained by totaling the contrast of a supervised block, as shown in equation (8).

$$I_{H}(x,y) = \left(\frac{I(x,y) - \min H}{\max H - \min H}\right) \times \max H | H \in C_{8}(I_{H})$$
(7)

$$H_d(r,c) = \sum_{(x,y)\in B} I_H(x,y)$$
(8)

# 2.4 Segmentation of brain tumor using Region of Interest (ROI) Command

As it has been seen from the above result that high density images have been separated from the MRI images using watershed segmentation. Here main aim is to segment the tumor from the MRI images. This can be done by using the ROI command in MATLAB. After the application of the ROI command, the tumor may be segmented. This can be achieved by executing the algorithm in MATLAB[4].

All the calculations and techniques can be summarized in following steps for the tumor detection:

Greyscale Imaging: MRI images are magnetic resonance scan of a patient taken with the help of a computer. Generally a greyscale looks like black

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> and white images [25]. The illusion of gray shading in a halftone image is obtained by rendering the image as a grid of black dots on a white background (or vice-versa), with sizes of individual dots determining the apparent lightness of the gray in their vicinity. In the case of transmitted light (for example, the image on a computer display), the brightness levels of the red (R), green (G) and blue (B) components are each represented as a number from decimal 0 to 255, or binary 00000000 to 11111111. For every pixel in a red-green-blue (RGB) greyscale image, R = G = B. The lightness of the gray is directly proportional to the number representing the brightness levels of the primary colours. Black is represented by R = G = B = 0 or R = G = B = 00000000, and white is represented by R = G = B = 255 or R = G = B = 111111111. Because there are 8 bits in the binary representation of the gray level, this imaging method is called 8-bit greyscale [18].

> The darkest possible shade is black, which is the total absence of transmitted or reflected light. The lightest possible shade is white, the total transmission or reflection of light at all visible wavelengths. So because of the above reason first we convert our MRI image to be pre-processed in greyscale image [1].

- II. High Pass Filter: A high pass filter is method for sharpening methods. An image is sharpened when contrast is enhanced between adjoining areas with variation in brightness or darkness. The kernel array of the high pass filter is designed to increase the brightness of the centre pixel relative to neighbouring pixels. The kernel array contains a positive value surrounded by negative values [8].
- III. Medial filter: Median filter is very widely used in digital image processing because, under certain conditions, it preserves edge while removing noise.
  [13] The task of filtering is performed by the median filter by use of window that is a pattern of neighbours. The window pattern slides, entry by entry, over the entire signals. The middle value of the window is decided by the median value of all the entries at a time.
- IV. Threshold segmentation: This method is based on clip-level (or threshold value) to turn a gray-scale image into a binary image by selecting a threshold value. Segmentation is the process of partitioning a digital image into multiple segments, called superpixels. [18] Image segmentation is typically used to locate objects and boundaries in images. By thresholding we assign a specific level to each pixel in image according to the similarity in visual characteristics [5].

Watershed segmentation: A grey-scale may be seen as a topographic relief, where the grey level of a pixel is interpreted as its altitude in the relief. In image processing, different watershed lines may be computed. Mayor's flooding Watershed algorithm is used to compute the watershed segmentation [14]. This algorithm works on a grey scale image. During the successive flooding of the grey value relief, watersheds with adjacent catchment basins are constructed. This flooding process is performed on the gradient image, i.e. the basins should emerge along the edges. Normally this will lead to an oversegmentation of the image, especially for noisy image material, e.g. medical CT data. Either the image must be preprocessed or the regions must be merged on the basis of a similarity criterion afterwards. [28]

V.

VI. Morphological **Operations:** Morphological operation is defined as the relative ordering of all the pixel values, not on their numerical values, and therefore are especially suited for the processing of binary images. Morphological techniques probe an image with a small shape or template called a structuring element. The structuring element is positioned at all possible locations in the image and it is compared with the corresponding neighbourhood of pixels. Some operations test whether the element "fits" within the neighbourhood, while others test whether it "hits" or intersects the neighbourhood.

A morphological operation on a binary image creates a new binary image in which the pixel has a non-zero value only if the test is successful at that location in the input image. [31]

# **3 RESULT AND CONCLUSSION**

The discussed methodology can be made clearer by considering an MRI image of tumor affected brain, shown in figure 3, and then performing all the operations in a sequence.

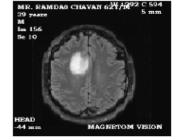


Figure 3 MRI image of tumor affected brain

We get greyscale image of the MRI image as shown in figure 4.

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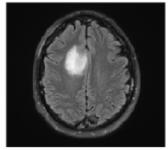


Figure 4 Greyscale image

On performing high pass filtering on the greyscale image of figure 4 the HPF output is obtained as shown in figure 5.



### Figure 5 HPF output

There may be some noise in the greyscale image of the brain, agter removing the noise the enhanced MRI image is figure 6.

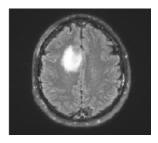


Figure 6 Enhanced MRI image

The MRI greyscale image is converted into a binary image by applying threshold segmentation and given in figure 7 [18].

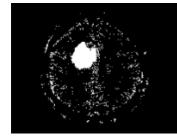


Figure 7 Threshold segmented image of input image

For more accurate and precise result, watershed segmentation is done on the threshold segmented image and we get the output image as shown in figure 8.

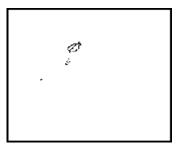


Figure 8 Watershed segmented image of input image

After the watershed segmentation the relative ordering of the pixel value of the image is done by morphological operation and the output in each step is shown in figure 9.

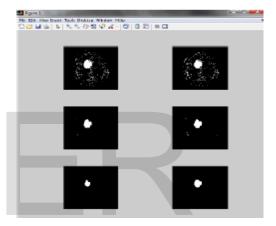


Figure 9 Morphological operations

Finally, on observing the difference in brightness or darkness in the MRI image, the brain tumor can be detected as given in figure 10.

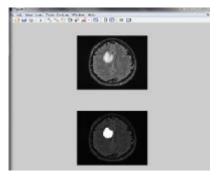


Figure 10 Final Extracted brain tumor from MRI image

### **4 FUTURE DIRECTION**

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IJSER © 2014 http://www.ijser.org In future this programme can be done more advanced so that major tumor can be classified according to its type [16]. Advancement in programming can help the segmentation completely automatic. By plotting the graph the growth of the tumor can be analysed [9].

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