

A SIMPLE IMAGE PROCESSING APPROACH TO ABNORMAL SLICES DETECTION FROM MRI TUMOR VOLUMES

T.Kalaiselvi, P.Nagaraja and P.Sriramakrishnan

Department of Computer Science and Applications,
Gandhigram Rural Institute – Deemed University, Gandhigram, Tamil Nadu, India

ABSTRACT

This paper proposed a method for brain tumor detection from the magnetic resonance imaging (MRI) of human head scans. The proposed work explained the tumor detection process by means of image processing transformations and thresholding technique. The MRI images are preprocessed by transformation techniques and thus enhance the tumor region. Then the images are checked for abnormality using fuzzy symmetric measure (FSM). If abnormal, then Otsu's thresholding is used to extract the tumor region. Experiments with the proposed method were done on 17 datasets. Various evaluation parameters were used to validate the proposed method. The predictive accuracy (PA) and dice coefficient (DC) values of proposed method reached maximum.

KEYWORDS

Minima Transform, Thresholding, Transforms, Tumor.

1. INTRODUCTION

Brain tumor is cluster of abnormal and an uncontrolled growth of cells in the brain [1]. Brain tumors can be classified according to their origin or degree of aggressiveness. Primary brain tumors arise in the brain, while metastatic brain tumors frequently originate from other parts of the body [2]. Magnetic resonance imaging (MRI) plays important role in many medical imaging applications. MRI provides prosperous information about the human soft tissue anatomy as well as helps to diagnosis of brain tumor [3]. In recent years, MRI has become an important modality for neurological image diagnosis. The combination of different sequences of MRI techniques is used to diagnose tumor [4]. The sequences include T1-weighted, T1-weighted with contrast enhancement (T1c), T2-weighted and fluid attenuated inversion recovery (FLAIR). Nowadays, brain tumor detection for MRI is difficult task for medical applications [5].

In recent years, many approaches have been developed for brain tumor detection. Jayachandran and Dhanasekaran proposed a hybrid algorithm for detection brain tumor in MRI images using statistical features and Fuzzy Support Vector Machine (FSVM) classifier. The proposed technique consists of four stages namely, Noise reduction, Feature extraction, Feature reduction and Classification. Stage I, anisotropic filter is applied for noise reduction and extracting the features. Stage II, obtains the texture features related to MRI images. Stage III, the features of magnetic resonance images have been reduced using principles component analysis to the most essential features. Stage IV, the supervisor classifier based FSVM has been used to classify subjects as normal and abnormal brain MR images [6]. Somasundaram and Kalaiselvi proposed an automatic method to analyze the MRI head scans and detect abnormality in brain due to tumors. This method consist four stages: brain extraction algorithm, transformation, fuzzy segmentation and fuzzy symmetric analysis. This method used two measures: false alarm (FA)

and missed alarm (MA) to quantify the performance of the method. The mean FA no more detected, however the MA was detecting minimum quantitative value [7]. Logeswari and karnan proposed an enhanced implementation of brain tumor detection using segmentation based on soft computing. The proposed method used two phased for detection tumor. In first phase MRI image of brain is collected. After that using preprocessing technique image is converted into standard form. Second phase for image segmentation using hierarchical self organizing map (HSOM) method is applied on image [8].

Roy and Bandyopadhyay proposed an interactive segmentation method that enables users can quickly and efficiently segment tumors in MRI of brain. In addition to area of the region and edge information the proposed method uses a type of prior information also its detecting the tumor region on exactly from MRI [9]. Padole and Chaudhari proposed an efficient method for brain tumor detection. Combination of two standard algorithm, mean shift and normalized cut is performed to detect the brain tumor surface area in MRI. Segmentation of brain, detects tumor and also its physical dimension and its segmentation accuracy is discussed [10]. Anandgaonkar and Sable proposed a survey on different segmentation techniques applied to MR images for locating tumor. It also includes a proposed method for the same using Fuzzy C-Means algorithm and an algorithm to find area of tumor which is useful to decide type of brain tumor whether it is benign or malignant [11].

The proposed work is an automatic method for extraction of the complete tumor region which overcomes the above said problems and works efficiently for FLAIR and T2-weighted images. The proposed method introduces enhancing process using top and bottom hat transformations and minima transform. Initially MRI head scans are enhanced by using top and bottom-hat process. Extended minima transform is used to separate tumor region from the enhanced image. In resultant segmentation, the fuzzy symmetric measure (FSM) is used to check the abnormality detection. Finally applied Otsu's thresholding in abnormal slice, it extracts the tumor region. The proposed method gives better results for detecting the tumor regions from FLAIR and T2-weighted images.

This paper is organized as follows. The image processing transformations are explained in section 2, the Otsu's thresholding is explained in section 3, the proposed method is explained in section 4, the results and discussion are given in section 5 and the conclusion is given in section 6.

2. IMAGE PROCESSING TRANSFORMATIONS

2.1. Top-hat and Bottom-hat Transforms

The top-hat transform is defined as the difference between the input image and its opening by some structuring element [12]. Structuring element (SE) of proposed method is given by,

$$SE = ('disk', 15) \tag{1}$$

The application of these transforms is in removing objects from an image by using an SE in the opening and closing that does not fit the objects to be removed. The difference then yields an image with only the removed objects.

Then, the top-hat transform of f is given by:

$$\tau_w(f) = f - (f \circ h) \tag{2}$$

where \circ denotes the opening operation. The top-hat transform is an operation that extracts small elements and details from given images. The top-hat transform performs light objects on a dark background.

The bottom-hat transform is defined dually as the difference between the closing and the input image. Then, the bottom-hat transform of f is given by:

$$T_b(f) = (f \bullet b) - f \quad (3)$$

where \bullet is the closing operation. Bottom-hat performs morphological bottom-hat filtering on the grayscale or binary input image. The bottom-hat performs for dark objects on a light background.

2.2. Extended Minima Transform

The extended minima transform, which is the regional minima of the H-minima transform. Extended minima used 8-connected neighborhoods for 2-D images and 26-connected neighborhoods for 3-D image. Performing the H-minima transform on the inverse distance image can effectively decrease over segmentation is reduced to some extent after applying morphological filters. The H-minima transform [13] is performed by:

$$H_h(f) = R_f(f + h) \quad (4)$$

where h represents the given depth. R represents the reconstruction and erosion operators, respectively.

3. Otsu's Method

This method is called as optimum threshold method and provides satisfied results in MRI brain images [14]. Otsu's thresholding involves all possible threshold values and calculate the pixel levels in each side of the threshold. This threshold value separates the foreground or background of pixels. This algorithm compute the image to be threshold contains the two classes of pixels. We can use the within class variance, it is the weighted sum of the variances of each foreground and background [15].

$$\sigma_{within}^2(T) = W_B(T)\sigma_{bg}^2(T) + W_F(T)\sigma_{fg}^2(T) \quad (5)$$

Where,

$$W_B(T) = \sum_{i=0}^{T-1} p(i) \quad W_F(T) = \sum_{i=T}^{N-1} p(i) \quad (6)$$

p_i – is the probability of occurring of pixel value x_i .

The mean of foreground and background pixels is,

$$\mu_B(T) = \frac{1}{W_B(T)} \sum_{i=1}^T iP(i) \quad (7)$$

$$\mu_F(T) = \frac{1}{W_F(T)} \sum_{i=1}^T iP(i) \tag{8}$$

The variance of foreground and background pixels is,

$$\sigma_{bg}^2(T) = \frac{1}{W_B(T)} \sum_{i=1}^T (i - \mu_B(T))^2 P(i) \tag{9}$$

$$\sigma_{fg}^2(T) = \frac{1}{W_F(T)} \sum_{i=1}^T (i - \mu_F(T))^2 P(i) \tag{10}$$

$\sigma_{bg}^2(T)$ -variance of the pixels in the background.

$\sigma_{fg}^2(T)$ -variance of the pixels in the foreground.

4. PROPOSED METHOD

This proposed method is a fully automated brain tumor detection method. The enhanced MRI images are produced by top, bottom-hat, minima transformations and segmented by Otsu's thresholding to extract the tumor portion from both T2-weighted and FLAIR images. The flow chart of tumor extraction methods are shown in Figure 1.

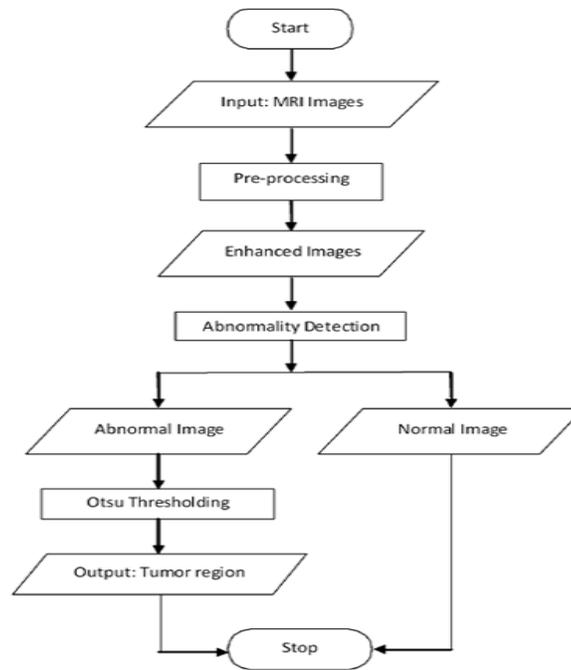


Figure 1. Flow chart for proposed method

4.1. Pre-processing

The proposed method initially has done a pre-processing technique on MRI images. The flow diagram of preprocessing method is given in Figure 2.

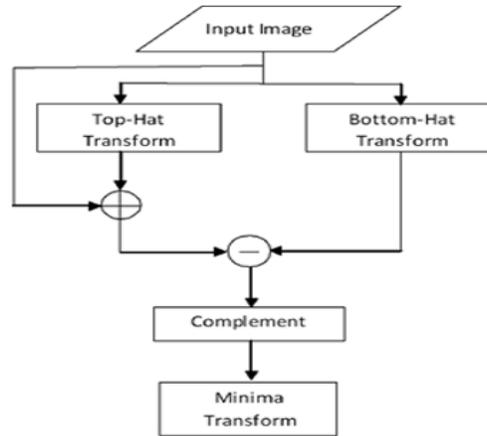


Figure 2. Flow chart for preprocessing

The datasets are processed by top-hat and bottom-hat transformation for enhancing the original image. The preprocessing images are shown in Figure 2. The original image is shown in column 1. The top-hat transformation as shown in column 2 is used to maximize the contrast between the objects and the gaps that separate them from each other. The top-hat transform is defined dually as the difference between the opening and the input image. Bottom-hat performs morphological bottom-hat filtering on the grayscale or binary input image and performs dark objects on a light background. The bottom-hat transform is defined dually as the difference between the closing and the input image. The bottom-hat transform image is shown in column 3. Then add the top-hat image to the original image, and then subtracts the bottom-hat image. This stage tumor area get sharpened region and it is shown in column 4

Then complement the function is used to enhance the image based on the intensity valleys as represented in column 5. The minima transformation is used to detect the intensity valleys deeper than a particular threshold with that function. It removes local peaks which are lower than h intensity values from the background. Based on the analysis done during our experiments, h is set to 8. The enhanced tumor regions extracted by using extended minima transform is shown in column 6. In column 6, the normal volume enhanced images are given in row 1.

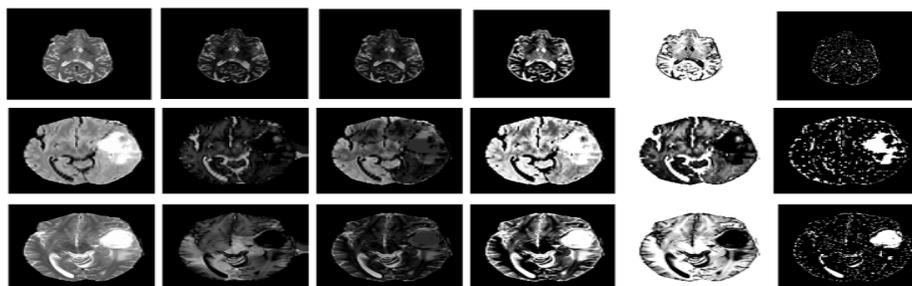


Figure 3. Preprocessing Stage, original images for Normal, FLAIR and T2-weighted

Figure 3. Preprocessing Stage, original images for Normal, FLAIR and T2-weighted are in column 1, top-hat images are in column 2, bottom-hat images are in column 3, enhanced images are in column 4, complement images are in column 5 and minima transformation images are in column 6.

4.2. Abnormality Detection

Abnormal tissues in the enhanced tumor region image can be detected by measuring the symmetry of enhanced tumor image. Here the symmetry is computed by the fuzzy symmetric measure (FSM) [16] given by,

$$FSM = \frac{1}{1 + \left(\frac{n_L - n_R}{200}\right)^2} \quad (11)$$

where n_L and n_R are the number of foreground (white) pixels in the left and right half of the given image present at either side of the central vertical line of slice. The symmetry values calculated from normal enhanced image are generally much larger than 0.1, and the values for abnormal enhanced image are much smaller than 0.1 [7] [17].

4.3. Brain Tumor Detection

Then the process is carried out by abnormal slice. The tumor extraction method is used by Otsu's thresholding technique. It separates the tumor region from background. In the resultant detection, tumors are extract from the abnormal slice and as a largest connected region as shown in Figure 4.

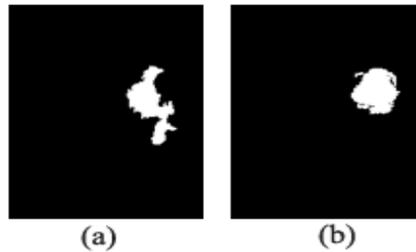


Figure 4. (a) FLAIR tumor region, (b) T2-weighted tumor region.

5. RESULTS AND DISCUSSION

Our algorithm was implemented in MATLAB2009 on a PC with Intel Pentium Core Duo 1.6GHz processor and 512MB RAM. Our proposed method used 17 datasets from Brain Tumor Image Repository (BTIR) maintained by our research group [18]. The first two datasets (N01–N02) consisting of 2 normal volumes and V01–V15 are abnormal volumes with glioma brain tumor. In tumor portion, two types of outputs from FLAIR and T2-weighted were added and its represents to the complete tumor region is as shown in Figure 5. The qualitative validation in the form of visual inspection is done with some of the sample MRI brain images shown in Figure 6. In Figure 6, the original FLAIR images are given in column 1, the original T2-weighted MRI images are given in column 2, the corresponding ground truth images are given in column 3, and the results of proposed method are given in column 4. This proposed work gives good results for glioma images.



Figure 5. Complete tumor region.

For quantitative validation, the performance is checked against three parameters. They are predictive accuracy (PA), dice coefficient (DC). The segmented images could have an error rate and an instance may either fail to identify an abnormality or identify an abnormality when there is none. The parameter PA is used to describe the error rate by the terms true and false positive (TP, FP) and true and false negative (TN, FN).

$$PA (\%) = \frac{TP+TN}{TP+TN+FP+FN} * 100 \quad (11)$$

where,

- TP = the test result is positive in the abnormal cases correctly classified.
- TN= the test result is negative in the normal cases correctly classified.
- FP = the test result is positive in the normal cases classified abnormal.
- FN= the test result is negative in the abnormal cases classified normal.

The parameter DC is used to compare the similarity between manual segmentation and the result of the proposed work. The value for DC ranges from 0 to 1 where 0 for no agreement and 1 for exact agreement.

The DC is given by:

$$DC(A, B) = \frac{2|A \cap B|}{|A|+|B|} \quad (12)$$

where A represents the ground truth image and B represents the proposed result image.

Table 1 is shown the abnormal detection by evaluated parameters FA and MA. The missed alarm is 2.3 % from the proposed method and no more FA in entire datasets. The performance analysis of proposed method is based on the parameters PA and DC values given in Table 2. The graph shows the quantitative representation of PA and DC of proposed method with line representation. In Figure 7, X axis represents volume id considered for experiment and Y axis represents the PA and DC values proposed method. The PA and DC of proposed method are 98% and 75% in glioma images. Our proposed method gives good results for glioma images.

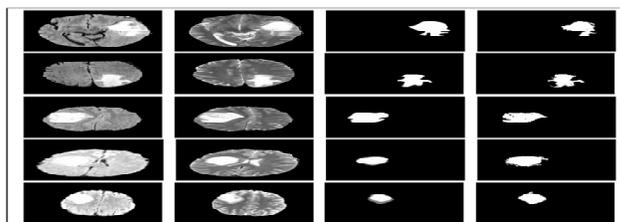


Figure 6. FLAIR images are in column 1, T2-weighted images are in column 2, corresponding ground truth images are in column 3 and the results of proposed method are in column 4.

Table 1. FA and MA values for proposed method

No	Vol. id	Total Slice	Abnormal Slice		FA		MA	
			Actual	Detected	Slices	%	Slices	%
1	N01	54	-	-	0	0	0	0
-	-	43	-	-	0	0	0	0
3	V01	176	46-119	50 - 119	0	0	46-49	2
4	V02	176	44-110	47-106	0	0	44-46, 107-110	4
5	V03	176	62-146	67-141	0	0	62-66, 142-146	6
6	V04	176	81-150	81-145	0	0	146-150	3
7	V05	176	68-131	69-131	0	0	68	0.6
8	V06	176	87-161	87-160	0	0	161	0.6
9	V07	176	78-137	81-135	0	0	78-80, 136,137	3
10	V08	176	55-133	60-133	0	0	55-59	3
11	V09	176	75-147	77-145	0	0	75,76, 146,147	2
12	V10	176	77-117	80-115	0	0	77-79, 116,117	3
13	V11	230	118-167	118-166	0	0	167	0.6
14	V12	165	78-119	78 - 119	0	0	0	0.4
15	V13	220	81-132	85-127	0	0	81-84, 127-132	5
16	V14	230	94-120	96-114	0	0	95-96, 115-120	2
17	V15	163	61-101	61-101	0	0	0	0
Average Performance					0	0		2.3

Table 2. PA and DC values of the proposed method

S.No	Vol. id	PA (%) (Complete Tumor)	DC (%) (Complete Tumor)
1	V01	98	84
2	V02	97	73
3	V03	97	87
4	V04	98	85
5	V05	98	60
6	V06	98	77
7	V07	98	62
8	V08	98	81
9	V09	96	69
10	V10	99	61
11	V11	99	92
12	V12	97	75
13	V13	98	73
14	V14	98	74
15	V15	99	77
	Mean	97.9	75.3

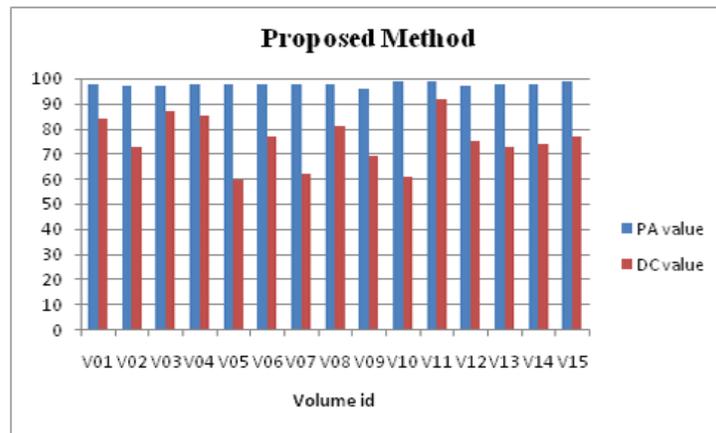


Figure 7. PA and DC measurement Graph for the proposed method

6. CONCLUSIONS

The proposed work developed a brain tumor detection method using top-hat, bottom-hat, minima transformations and Otsu's thresholding. The experimental method is done on 17 datasets. This work detected the abnormal portion of the brain which is present in MRI images. The advantage of this proposed method is it takes minimum missed alarm while detecting the tumor in MRI images. It gives 75% similarity index while compared with ground truth dataset.

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Authors

Kalaiselvi T. currently working as an Assistant Professor in Department of Computer Science and Applications, Gandhigram Rural Institute - Deemed University, Dindigul, India. She received her Bachelor of Science (BSc) degree in Mathematics and Physics in the year 1994 and Master of Computer Applications (MCA) degree in the year 1997 from Avinashilingem University, Coimbatore. She received her Ph.D (Full-time) degree from Gandhigram Rural Institute in the year 2010. In the year 2008, she received a project from Department of Science and Technology (DST), Government of India under Scheme for Young Scientists and Professionals (SYSP) by Science for Equity, Empowerment and Development (SEED) Division for three years (2008-2011). Her research focuses on Brain Image Processing and brain tumor or lesion detection from MR Head Scans to enrich the Computer Aided Diagnostic process, Telemedicine and Tele radiology services. She is Academic Community Member (ACM) in International Congress for Global Science and Technology (ICGST), Life Member (LM) in Indian Society for Technical Education (ISTE) and Lifetime Member (LM) in Telemedicine Society of India (TSI).



Nagaraja P. is a Research Scholar (Full-time) in the Department of Computer Science and Applications, Gandhigram Rural Institute - Deemed University, Dindigul, India. He received his Bachelor of Science (B.Sc) degree in Physics in the year 2008 and Master of Computer Applications (MCA) degree in the year 2011 from Gandhigram Rural Institute - Deemed University. He is currently pursuing Ph.D. degree in Gandhigram Rural Institute- Deemed University. His research focuses on Brain Tissue Segmentation in MRI Head scans.



P. Sriramakrishnan received his Bachelor of Science (B.Sc.) degree in 2011 from Bharathidasan University, Trichy, Tamilnadu, India. He received Master of Computer Application (M.C.A) degree in 2014 from Gandhigram Rural Institute-Deemed University, Dindigul, Tamilnadu, India. He worked as Software Developer in Dhvani Research and Development Pvt. Ltd, Indian Institute of Technology Madras Research Park, Chennai during May 2014 – March 2015. He is currently pursuing Ph.D. degree in Gandhigram Rural Institute- Deemed University. His research focuses on Medical Image Processing and Parallel Computing. He has qualified UGC-NET for Lectureship in June 2015.

