

A SIMPLE APPROACH FOR RELATIVELY AUTOMATED HIPPOCAMPUS SEGMENTATION FROM SAGITTAL VIEW OF BRAIN MRI

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ABSTRACT

In this paper, we present a relatively automated method to segment the hippocampus in 11 weighted magnetic resonance images that can be acquired in the routine clinical setting. This paper describes a simple approach for segmenting the hippocampus automatically from sagittal view of brain MRI. Large datasets of structural MR images are collected to quantitatively analyze the relationships between brain anatomy, disease progression, treatment regimens, and genetic influences upon brain structure. This method segments the hippocampus without any human intervention for few slices present mid position in the total volume. Experimental results using this method show a good agreement with the manual segmented gold standard. These results may support the clinical studies of memory and neurodegenerative disease.

KEYWORDS

Hippocampus, Alzheimer's disease, neurodegenerative, MRI, segmentation, sagittal view.

1. INTRODUCTION

Segmentation of medical images is the task of partitioning the data into contiguous regions representing individual anatomical objects. It is a prerequisite for further investigations in many computer-assisted medical applications, e.g. individual therapy planning and evaluation, diagnosis, simulation and image guided surgery. The hippocampus is a complex brain region primarily responsible for memory and any of its associated neurodegenerative diseases such as Alzheimer's disease. Volume and size of the hippocampus are of great importance in the analysis of neurodegenerative diseases. Hippocampus segmentation and its subfield segmentation are playing important roles in the treatment of neurodegenerative diseases[1].

Chupin et al. [2],[3] used a Markovian deformation process with a deformable constraint based on prior knowledge, which is calculated with manual intervention. A user must manually define bounding boxes and create seed points to make algorithm to work and thus making the entire process as semi-automated.

A study by Ashton et al [4] use elastic deformable model with seed points and constraints to segment the hippocampus. The seed points and the constraint were produced from boundaries that had previously been traced manually. These methodologies speed up processing time of segmentation considerably compared to manual segmentation, allowing them to process a large number of scans. The necessity of including human involvement still increases processing time, and it provides a subjective element to segmentation as in manual segmentation.

Wei Wei Lee et al. [5], proposed a semi-automatic hybrid approach for segmentation of Hippocampus. This method combined low-level image processing techniques such as thresholding, hole filling (based on adjacent voxel connectivity), distance transformation, with high level image processing techniques, and termed it as Geometric Deformable Model (GDM) in a sequential pipeline. The operation of the GDM was based on constraint modeling and cost function minimization. The GDM incorporated 5 constraints which were integrated together to form a local cost function (potential function) associated with each vertex in a 3D model.

An assessment for the performance of standard image registration techniques for MRI-based automated segmentation of the hippocampus was presented by O. T. Carmichael et al. [6]. The study was based on elderly subjects with Alzheimer's Disease as well as mild cognitive impairment (MCI). They have collected structural MR images from Alzheimer's Disease Research Center at the University of Pittsburgh. The subjects were of 54 years and gender-matched healthy individuals, with probable AD, and another set with MCI. The hippocampi from the subject images were segmented by the automated segmentation methods using cohort atlases such as AIR [7], SPM [8], FSL [9], and the fully-deformable method of Chen [10]. The segmented results were aligned to the Harvard atlas [11], MNI atlas [12], and also with manually-labeled subject images [10].

The segmentation of hippocampus in five patients with mesial temporal sclerosis is described by Robert E. Hogan et al. [13]. Using magnetic resonance (MR) imaging, they verified the precision as well as reproducibility of hippocampal segmentations. They used the deformation based segmentation method. The results produced by them had 92.8%, overall percentage overlap with that of automated segmentations.

In this paper we present an automated tool to segment the hippocampus from high sagittal view of brain MRI. Our method segments the hippocampus in few slices of the entire volume. The remaining part of the paper is organized as follows. In section II, we present the methods and the materials used. In section III, the results and discussion are given. Finally in section IV, the conclusion is given.

2. MATERIALS AND METHODS

Materials Used

The data has been collected from the Whole Brain Atlas [14]. The slices are T1 weighted and in sagittal orientation. The hippocampus appears in few slices in the provided atlas. No gold standard is provided for this atlas and it contains only labels for subcortical structures. We have done the manual segmentation with the help of the physicians Dr. C. Kathirvel M.S., M.Ch.(Neuro) and Dr. K. Veni M.D., D.M.(Neuro), Madras Medical College, Chennai..

Methods Used

The proposed method can segment the hippocampus from the multiple slices for the data set available in WBA [14]. The hippocampus appears only in few slices among 100 slices present in this volume. Figure 1 shows the mid-slice of the brain MRI in the sagittal view obtained from the Whole Brain Atlas [14]. Our method is relatively automated, that it is capable of validating the input slices automatically. The validation process involves checking whether the input slice is suitable for segmentation so that the user can provide any slice with or without HC. Our proposed method validates the input slice by analyzing the image characteristics mean intensity and standard deviation, and then selects the suitable image for segmentation automatically [15-17]. Figure 2 gives the flowchart of our proposed method.

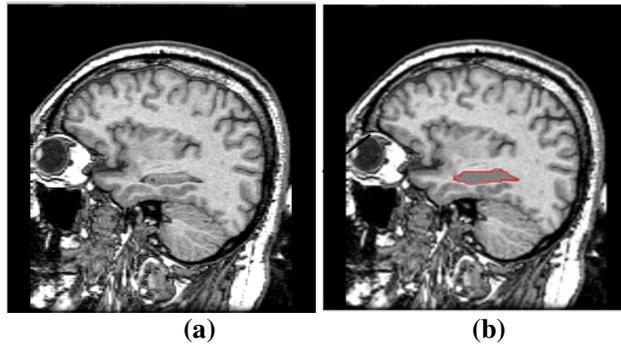


Figure 1 (a) Mid-slice of brain MRI in sagittal view (b) Hippocampus filled with dark gray color and marked with an arrow

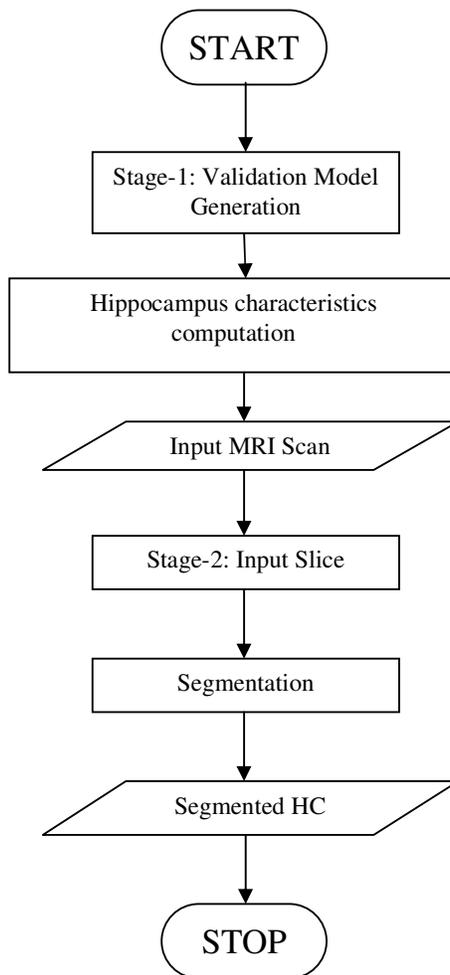


Figure 2 Flowchart of Proposed Method

These values are computed from the slices containing HC available in WBA [89]. The range of values for the valid slices is given in Table 1.

Table 1. Characteristics of Image with Hippocampus in WBA

Image Characteristic	Characteristic Value	
	Minimum	Maximum
Mean Intensity(\bar{x})	81.1062	88.01
Standard deviation (σ)	79.1111	88.2

When the input slice is provided by the user, the specified image characteristics (\bar{x} and σ) are computed. The mean value \bar{x} of the slice is computed as :

$$\bar{x} = \frac{\sum_{x=1}^m \sum_{y=1}^n f(x,y)}{m \times n}$$

where $f(x,y)$ is the intensity of the pixel $A(i,j)$, m is the number of rows and n is the number of columns. The standard deviation of the slice is computed as:

$$S = \left(\frac{1}{n} \sum_{i=1}^n (x_i - \bar{x})^2 \right)^{\frac{1}{2}} \quad \text{where,} \quad \bar{x} = \frac{1}{n \cdot \sum_{i=1}^n x_i}$$

where, x_i is the intensity of the pixel at i^{th} position.

These values are compared with the values mentioned in Table 1. If the values fall within the specified range, then it is a valid slice and our method proceeds with segmentation process, the input slice is discarded otherwise.

In the next step, the valid slice is enhanced to make the hippocampus prominent and distinguishable with the neighboring region resulting in segmentation of multiple slices. Here two-phase enhancement is implemented in order to make the hippocampus prominent from the neighboring region for multiple slices. The median filter is not suitable for this data set to segment multiple slices. Hence, trimmed mean filter is used for enhancement in addition to the hat operation, which makes the hippocampus boundary distinguished from neighboring pixels, in multiple slices, compared to the first method. The enhanced image is converted to binary image using the block mean threshold as explained in method-1. Binary erosion is used to differentiate the edges of HC of the binary image. VH_{s1} separated from the neighborhood pixels and is of the form:

$$H_{s1} = VH_s \ominus D_{s\epsilon}$$

where, $D_{s\epsilon}$ is a disc shaped structuring element with 17 as radius. The edge detection is done for the eroded image. The Laplacian Gaussian method is used here which has been selected by trial. The other edge detection techniques are unable to detect the edges for the multiple slices in this

data set. The middle block contains many connected components other than HC. We use a labeling process as explained in method-1, to assign different labels to each component. The HC is retrieved using the related label which is found by iteration. The HC thus extracted may have lost some of the pixels at the boundary because of the erosion. To recover these pixels, binary dilation is used on H_{s1} , with the same structuring element, which is defined as:

$$H_{s2} = H_{s1} \oplus D_{se}$$

A mask of HC is obtained after this process. The HC region is extracted using this mask by mapping it with that of the original input slice (VH_s) as defined by:

$$HC(i,j) = \begin{cases} VH_s(i,j), & \text{if } H_{s2}(i,j) = 1 \\ 0, & \text{otherwise} \end{cases}$$

where, H_{s2} is the mask, containing white pixels as the HC region and other regions appear as black.

3. RESULTS AND DISCUSSION

The computed values of J, D, S, Sp, PA, FPR and FNR are given in Table 2.

Table 2. The computed values of J, D, S, Sp, PA, FPR and FNR for ARAHS

Slice	Jaccard	Dice	Sensitivity	Specificity	Predictive accuracy	FPR	FNR
WBA-sagittal #95	0.9997	0.9998	0.9998	0.8772	99.968	0.0001	0.0002
WBA-sagittal #94	0.9994	0.9997	0.9995	0.9524	99.942	0.0000	0.0005
WBA-sagittal #93	0.9986	0.9993	0.9997	0.6507	99.8596	0.0011	0.0003
WBA-sagittal #92	0.9977	0.9988	1	0.5664	99.7742	0.0023	0.0000
WBA-sagittal #91	0.997	0.9985	1	0.5523	99.6994	0.0030	0.0000
WBA-sagittal #90	0.9985	0.9992	0.9998	0.7276	99.855	0.0013	0.0002
WBA-sagittal #89	0.9991	0.9995	0.9994	0.928	99.9084	0.0003	0.0006
AVERAGE	0.9986	0.9985	0.9997	0.7507	99.8581	0.0012	0.0003

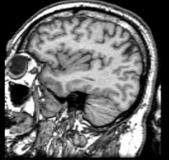
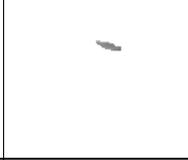
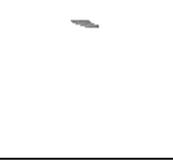
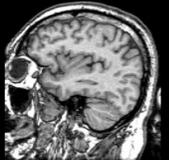
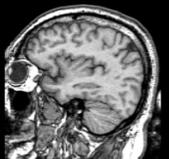
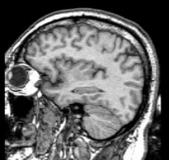
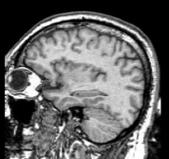
Slice	Input slice	Manual segmentation	Proposed method segmentation
Sagittal # 95			
Sagittal # 94			
Sagittal # 93			
Sagittal # 92			
Sagittal # 91			

Figure 2. Segmented results of ARAHS: first column shows the input slice, second column shows the manually segmented hippocampus and third column contains the segmented hippocampus by proposed method

From Table 2, we observe that the average of Jaccard index values is **0.9986** and the Dice coefficient is **0.9985**. This implies that our segmented results are better. The sensitivity value is closer to 1 and predictive-accuracy is very close to 100. The FPR and FNR values are nearly equal to zero. Hence our proposed method segments the hippocampus nearly equal to the gold standard. The segmented results are shown in Figure 2.

4. CONCLUSION

Our proposed method uses only simple techniques to extract the hippocampus from the sagittal view of the brain MRI. This method is developed in a motive of employing simple image processing techniques to segment the hippocampus from MRI head scans in different orientations. The strengths of these methods are their simplicity, automation and reduction of computation time.

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