Optimized Fuzzy Logic Based Segmentation for Abnormal MRI Brain Images Analysis

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Abstract

In this paper an optimized fuzzy logic based segmentation for abnormal MRI brain images analysis is presented. A conventional fuzzy c-means (FCM) technique does not use the spatial information in the image. In this research, we use a FCM algorithm that incorporates spatial information into the membership function for clustering. The FCM algorithm that incorporates spatial information into the membership function is used for clustering, while a conventional FCM algorithm does not fully utilize the spatial information in the image. The advantage of the technique is less sensitive to noise than the others. Originality of this research is focused in application of the technique on a normal and a glioma MRI brain images, and analysis of the area of abnormal mass from segmented images. The results show that the method effectively segmented MRI brain images, and the segmented normal and glioma MRI brain images can be analyzed for diagnosis purpose. The area of abnormal mass is identified from 7.15 to 19.41 cm².

Keywords: Adaptive image segmentation, FCM clustering, abnormal MRI brain image, fuzzy membership function.

1. Introduction

Image segmentation is one of the most important step to extract information in image processing. Segmentation has wide application in medicine area. The main objective of segmentation of medical image is to partition the image into mutually exclusive and exhausted regions such that each region of interest is spatially contiguous and the pixels within the region are homogeneous with respect to a predefined criterion. Magnetic Resonance Imaging (MRI) is the state-of-the-art medical imaging technology which allows cross sectional view of the body with unprecedented tissue contrast. MRI provides a digital representation of tissue characteristic that can be obtained in any tissue plane. The images produced by an MRI scanner are best described as slices through the brain. MRI has the added advantage of being able to produce images which slice through the brain in both horizontal and vertical planes. The objective of segmenting different types of soft-tissue in MRI brain images is to label complex structures with complicated shapes, as white matter, grey matter, CSF and other types of tissues in neurological conditions.

A variety of segmentation methods have been developed to satisfy increasing requirement of image segmentation over past several years. FCM (Fuzzy c-means) is unsupervised technique that has been successfully applied to future analysis, clustering, and classifier designs in the fields such as astronomy, geology, medical imaging, target recognition, and image segmentation. An image can be represented in various feature spaces, and the FCM method classifies the image by grouping similar data points in the feature space into clusters.

During the past many researchers in the field of medical imaging and soft computing have made significant survey in the field of image segmentation. There has been considerable interest recently in the use of segmentation methods based on fuzzy logic, which retain more information from the original medical image than hard segmentation methods (e.g. Bezdek et al. [1], Udupa et al. [2], Pham [3], Masoole and Moosavi [4]). The fuzzy Cmeans (FCM) clustering, in particular, can be used to obtain a accurate segmentation via fuzzy pixel classification. Unlike hard clustering methods which force pixels to belong exclusively to one class, FCM allows pixels to belong to multiple classes with varying degrees of membership functions. This approach allows additional flexibility in many applications and has recently been used in processing of medical images [5]. For example, in their segmentation of MRI brain images, Pham et al. [5] thresholded the FCM memberships in order to extract pixels which a high confidence of correct classification. Xu et al. [6] used deformable surfaces that converged to the peaks of the memberships. The FCM method, however, does not address the intensity inhomogeneity

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artifact that occurs in nearly all MRI [7]-[8]. Originality of this research is the method applied on a normal MRI brain image and a glioma MRI brain images, and analyze the area of abnormal mass from segmented images.

2. Fundamental Theory

2.1 Magnetic Resonance Imaging

Magnetic Resonance Imaging (MRI) is an imaging used primarily in medical settings to produce high quality medical images of the soft tissues. MRI is an imaging technique used primarily in medical settings to produce high quality images of the inside of the human body. In this section we give a brief description of the principles of MRI which are referred to [9]. In MRI, the image is a map of the local transverse magnetization of the hydrogen nuclei. This transverse magnetization in turn depends on several intrinsic properties of the tissue. MRI is based on the principles of nuclear magnetic resonance (NMR). The NMR phenomenon relies on the fundamental property that protons and neutrons that make up a nucleus possess an intrinsic angular momentum called spin. When protons and neutrons combine to form nucleus, they combine with oppositely oriented spins. Thus, nuclei with an even number of protons and neutrons have no net spin, whereas nuclei with an odd number of protons or neutrons possess a net spin. Hydrogen nuclei have an NMR signal since its nucleus is made up of only a single proton and possess a net spin. The human body is primarily fat and water, which have many hydrogen atoms. Medical MRI primarily images the NMR signal from the hydrogen nuclei in the body tissues.

The net spin of the nucleus around its axis gives it an angular moment. Since the proton is a positive charge, a current loop perpendicular to the rotation axis is also created, and as a result the proton generates a magnetic field. The joint effect of the angular moment and the self generated magnetic field gives the proton a magnetic dipole moment parallel to the rotation axis. Under normal condition, one will not experience any net magnetic field from the volume since the magnetic dipole moments are oriented randomly and on average equalize one another.

When placed in a magnetic field, a proton with its magnetic dipole moment processes around the field axis. The frequency of this precession, v_0 , is the resonant frequency of NMR and is called the Larmor frequency. The precession frequency is directly proportional to the strength of the magnetic field, i.e.

$$v_0 = \mathbf{g}\mathbf{B}_0 \tag{1}$$

where B_0 is the main magnetic field strength, and g is a constant called gyromagnetic ratio which is different for each nucleus (42.56 MHz/Tesla for protons).

Given a specimen, the application of a magnetic field B_0 would create a net equilibrium magnetization M_0 per cubic centimeter, which is aligned to the B_0 field. The M_0 is the net result of summing up the magnetic fields due to each of the H nuclei and is directly proportional to the local proton density (or spin density). However, M_0 is many orders of magnitude weaker than B_0 and is not directly observable. By tipping M_0 away from the B_0 field axis with an appropriate RF pulse having a frequency equals to the Larmor frequency, a longitudinal magnetization component ML and a transverse magnetization component MT is produced. When the RF pulse is turned off, the longitudinal magnetization component ML recovers to M_0 with a relaxation time T1, and the transverse magnetization component MT dephases and decays to zero with a relaxation time T2 1. During relaxation, the protons lose energy by emitting their own RF signal with the amplitude proportional to MT. which is referred to as the free-induction decay (FID) response signal. T2 indicates the time constant required from a given tissue type to decay for the FID response signal. The FID response signal is measured by an RF coil placed around the object being imaged.

In MR imaging, the RF pulse is repeated at a predetermined rate. The *repetition time*, *TR*, is the period of the RF pulse sequence is. The FID response signals can be observed at various times within the *TR* interval. *echo delay time*, *TE* is the time between which the RF pulse is applied and the response signal is measured. The *TE* is the time when the spin echo occurs due to the refocusing effects of the 180 degree refocusing pulse applied after a delay of *TE*/2 from the RF pulse. The *TR* and *TE* control how strongly the local tissue relaxation times, *T1* and *T2*, affect the signal. By adjusting *TR* and *TE* the acquired MR image can be made to contrast different tissue types.

2.2 Image Segmentation

The aim of medical image segmentation on human graphical interaction is to define regions, using methods such as manual slice editing, region painting and interactive thresholding. Rajapakse [13] had classified the different methods of image segmentation as four categories. (1) Thresholding region growing and edge based techniques. (2) The maximum-likelihood-classifier (MLC). These methods are basically depend on the prior model and its parameters. Vannier et al. [14] reported satisfactory preliminary results with Bayesian MLC. Ozkan et al. [15] researched about the MLC and the neural



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network classifier which showed the superiority of the neural network.

Some new methods of image segmentation that could be classified as statistical methods have been introduced in the past few years. Hansen [16] used a probabilistic supervised relaxation technique for segmenting 3D medical images. The method introduced the use of cues to guide the medical image segmentation. Those cues marked by the user have the mean and standard deviation as description parameters. (3) The neural networks methods one example of which is the work of Ahmed et al. [14] who used a two stages neural network system for CT/MRI image segmentation. The first stage is a self-organized principal component analysis (SOPCA) network and the second stage consists of a self-organizing feature map (SOFM). The results obtained compare favorably with the classical and statistical methods. (4) The Fuzzy Clustering methods. In [18] a comparison between the fuzzy logic and artificial neural network techniques in segmenting magnetic resonance images debated for the need of unsupervised technique in segmentation which was provided using the unsupervised fuzzy c-mean algorithm.

3. Fuzzy C-Means Clustering

The FCM method assigns pixels to each cluster by using fuzzy membership functions. Let $X=(x_1, x_2, .., x_N)$ denotes an image with N pixels to be partitioned into c clusters, where x_i represents multispectral (features) data. The algorithm is an iterative optimization that minimizes the cost function defined as follows [16]:

$$J = \sum_{j=1}^{N} \sum_{i=1}^{c} u_{ij}^{m} ||x_{j} - v_{i}||^{2}$$
(2)

where u_{ij} represents the membership function of pixel x_j in the ith cluster, v_i is the ith cluster center, and m is a constant. The parameter m controls the fuzziness of the resulting partition, and m = 2 is used in this study.

The cost function is minimized when pixel close to the centroid of their clusters are assigned high membership values, and low membership values are assigned to pixels with data far from the centroid. The fuzzy membership function represents the probability that a pixel belongs to a specific class. In the FCM algorithm, the probability is dependent solely on the distance between the pixel and each individual cluster center in the feature domain. The membership functions and cluster centers are updated by the following:

$$u_{ij} = \frac{1}{\sum_{k=1}^{c} \left(\frac{\|x_j - v_i\|}{\|x_j - v_k\|}\right)^{2/(m-1)}}$$
(3)

and

$$v_{i} = \frac{\sum_{j=1}^{N} u_{ij}^{m} x_{j}}{\sum_{j=1}^{N} u_{ij}^{m}}$$
(4)

where $u_{ij} \in [0, 1]$.

Starting with an initial guess for each cluster center, the FCM converges to a solution for v_i representing the local minimum or a saddle point of the cost function. Convergence can be detected by comparing the changes in the membership function or the cluster center at two successive iteration steps.

One of the important characteristics of an image is that neighboring pixels possess similar feature values, and the probability that they belong to the same cluster is great. This spatial relationship is important, but it is not utilized in a konvensional FCM algorithm. To exploit the spatial information, a spatial function is defined as.

$$h_{ij} = \sum_{k \in NB(x_j)} u_{ik}$$
(5)

In this formula, $NB(x_j)$ represents a square window centered on pixel x_j in the spatial domain. A 3 x 3 window was used throughout this work. Just like the membership function, the spatial function h_{ij} represents the probability that pixel x_j belongs to *i*th cluster. The spatial function of a pixel for a cluster is large if the majority of its neighborhood belongs to the same cluster. The spatial function in incorporated into membership function as follows:

$$u'_{ij} = \frac{u^{p}_{ij}h^{q}_{ij}}{\sum_{k=1}^{c}u^{p}_{kj}h^{q}_{kj}}$$
(6)

In this formula, p and q are parameters to control the relative importance of both functions. In a homogenous region, the spatial functions fortify the original membership, and the clustering result remains unchanged. However, for a noisy pixel, this formula reduces the weighting of a noisy cluster by the labels of its

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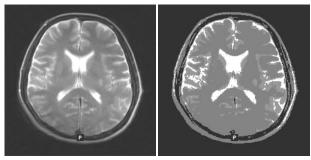
neighboring pixels. As a result, misclassified pixels from noisy regions or spurious blobs can easily be corrected. The spatial FCM with parameter p and q is denoted $sFCM_{p,q}$. Note that $sFCM_{1,0}$ is identical to the conventional FCM.

The clustering is a two-pass process at each iteration. The first pass is the same as that in standard FCM to calculate the membership function in the spectral domain. In the second pass, the membership information of each pixel is mapped to the spatial domain, and the spatial function is computed from that. The FCM iteration proceeds with the new membership that is incorporated with the spatial function. The iteration is stopped when the maximum difference between two cluster centers at two successive iterations is less than a threshold (ε =0,02). After the convergence, defuzzification is applied to assign each pixel to a specific cluster for which the membership is maximal

4. Experimental Results

4.1 Normal MRI Brain Image

In this research, MRI image is segmented using FCM method incorporated into the spatial information. Figure 1(a) shows the 256x256 grayscale original T2-weighted MRI1 brain image. Fig. 1(b) shows the result of the FCM incorporated into the spatial information with parameters p=1 and q=2.



(a) Original MRI1 Brain image

(b) Segmented MRI1 Brain image

Figure 1. MRI1 Brain image

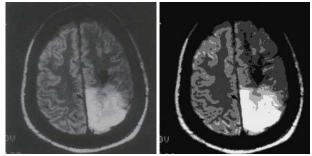
As can be seen in the segmented images in Fig. 2, lesion or abnormal mass is not identified, and the ventricular system is not extensive and it is a median. So, the image is normal brain MRI image.

4.2 Glioma MRI Brain Images

Figure 2(a) shows the 256x256 original MRI2 image [17]. As can be seen in the segmented image in Fig. 2(b), there

is extensive edema extending anteriorly and posteriorly and involving the basal ganglia.

The shift and mass effect on the ventricle have resulted in compromise of the foramen of Monro, and there is evidence of active hydrocephalus, as shown by both the ventricular enlargement and a homogeneous increase through the periventricular region. An additional area of increased signal, separated from the right hemisphere in the left side, suggest that there may even be a metastatic focus or evidence of distant extension of tumor into the left hemisphere. All these features suggest a very high Grade IV glioma or glioblastoma.



(a) Original MRI2 Brain image (b) Segmented MRI2 Brain image

Figure 2. MRI2 Brain image

4.3 Analysis of the Area of Abnormal Mass

In this study, we also apply the FCM method to segment four 256x256 MRI brain images with abnormal mass (i.e. MRI3, MRI4, MRI5, MRI6, MRI7, and MRI8), as shown in Figure 3 [17].

Application of the FCM method in segmentation of the images is in order to analysis of the area of abnormal mass Figure 4 shows the 256x256 segmented MRI3-MRI8 images.

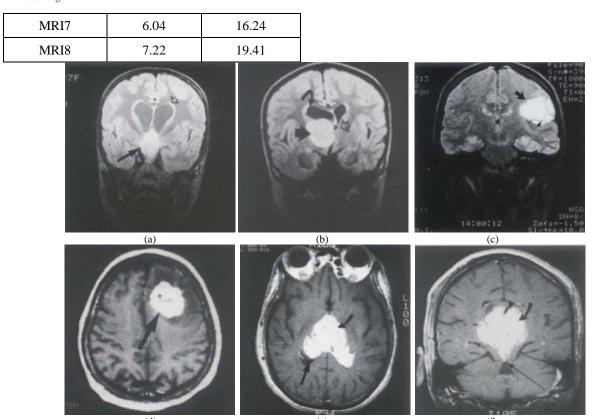
Table 1 shows area of abnormal mass of segmented MRI images. The results are MRI images identified tumors of 2.66% to 7.22% or 7.15 to 19.41 cm². In the brain tumor, glioma, the bigger area of tumor the higher grade of glioma.

Table 1. Area of abnormal mass

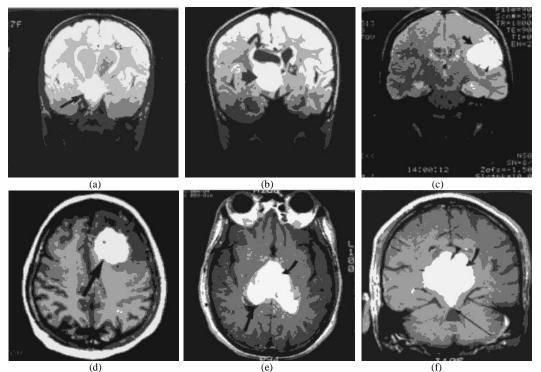
Images	Areas in %	Areas in cm ²
MRI3	2.66	7.15
MRI4	3.05	8.20
MRI5	3.57	9.60
MRI6	3.61	9.70



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(d) (e) (f) Figure 3. Original abnormal mass MRI brain images: (a) MRI3, (b) MRI4, (c) MRI5, (d) MRI6, (e) MRI7, (f) MRI8.



(d) (e) (f) Figure 4.(a) Segmented MRI3, (b) segmented MRI4, (c) segmented MRI5, (d) segmented MRI6, (e) segmented MRI7, (f) segmented MRI8.

5. Conclusions

In this paper we apply an extended FCM method that incorporates the spatial information into the membership function to improve the results of MRI brain image segmentation. The membership functions of the neighbors centered on a pixel of MRI brain image in the spatial domain are enumerated to obtain the cluster distribution statistics. These statistics are transformed into a weighting function and incorporated into the membership function. This neighboring effect reduces the number of spurious blobs and biases the solution toward piecewise homogeneous labeling. The technique was used to analyze a normal MRI brain image and glioma MRI brain images. We applied the technique on a normal MRI brain image and on a glioma MRI brain image. The results show that the method effectively segmented Magnetic Resonance Imaging (MRI) brain images with spatial information, and the segmented normal and glioma MRI brain images can be analyzed for diagnosis purpose. The results are MRI images identified tumors of 7.15 to 19.41 cm².

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