

Hybrid Method for Detection Tumor using Genetic Algorithm and Swarm Optimization after Wavelet Domain Filtering Then using Marr-Hilserth

Hind Rustum Mohammed and Lamyaa Fahem Katran

Department of Computer Science, Faculty of Computer Science and Mathematics,
University of Kufa, Kufa, Iraq

Abstract: Brain tumors consist of several types of abnormal growth within the skull and the Central Nervous System (CNS). They result from abnormal and uncontrolled cell division. Brain tumors are dangerous and when left untreated can result in loss of life. Some tumors, however, do not lead to death, especially, the lipomas which are inherently unharmed. The potential risk associated with tumors depends on several factors such as the type of the tumor, its size and location as well as the condition of the tumor. This study proposes the use of pre-processing phase wave image filter size box algorithm for brain tumor imaging as this algorithm provides improved image quality devoid of noise. The suggested method contains three essential steps, the first step involves the use of a Genetic algorithm which has a zero tolerance for probability and flexibility and can also find near-optimal solutions. In the second step an entrained Particle Swarm Optimization (PSO) technique was used to mechanically determine the mid-clustering the randomly collected data set. The third stage involved the merging of one and two regions of segmentation in the data set using marshalled to detect a tumor in the brain. The noise in the image was filtered using the Laplacian-Gaussian technique before edge detection. The Laplacian-Gaussian technique involves the merging of a Gaussian filtering technique with the Laplacian technique for edge detection. The technique involves three main steps-filtering, improvement and discovery before finally calculating the tumor area using the proposed algorithm.

Key words: Magnetic Resonance Imaging (MRI), wavelet domain filtering square magnitude Genetic algorithm, particle swarm optimization, Marr-Hilserth, seed image region, tumor area

INTRODUCTION

Brain tumor consists of two main types, the benign (non-cancerous) and the malignant (cancerous) tumors. The cancerous brain tumors can develop from any part of the body and slowly spread out to other parts of the human body and the brain too. For this, they are referred to as cancerous or metastatic tumors (Mohammed and Katran, 2018; Singh *et al.*, 2004). A non-cancerous or benign brain tumor consists of a mass of relatively slowly growing cells that settle in any part of the brain. They do not spread from one region to another and if surgically removed do not often relapse. Benign tumors that are not entirely removed during a surgical procedure can grow back but will not affect the other parts of the brain. This behavior of both types of tumor requires close monitoring using scans or treatment with <https://www.nhs.uk/conditions/radiotherapy/pages/introduction.aspx> either chemotherapy or radiotherapy. Benign brain tumors are of different types based on the type of brain

cells they affect. Some of the common benign brain tumors include the gliomas which affects the glial tissue that provides support to the nerve fibres and cells, the meningioma that affect the meninges that cover the brain surface, the acoustic neuromas <https://www.nhs.uk/conditions/acoustic/neuroma/pages/introduction.aspx> (also known as vestibular schwannomas) which affect the acoustic nerves, the craniopharyngiomas that affect the base of the brain. They are the most diagnosed form of brain tumor in children, teenagers and adolescents. Others include the haemangioblastoma as which affect the blood vessels of the brain and the pituitary adenomas which affect the pituitary gland. Brain cancer consists of the primary brain tumors which develop in the brain and does not metastasize to other body parts, the secondary tumors which develop from tumors from other parts of the body and metastasis a size to the brain. There are several types of brain tumor (more than 40 major types), grouped into 2 major types. The first type is the benign or slow-growing types which are unlikely to metastasize. Common types of

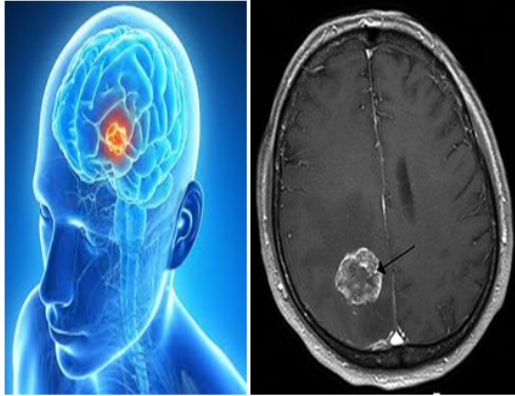


Fig. 1: Image of tumor in brain of human

benign brain tumors include meningiomas, pituitary tumors and neuromas. The second type is the malignant or cancerous brain tumors which can spread from one part of the brain to the other including to the spinal cord. Some examples of the malignant brain tumors include tracy tumors, globalist, oligomer drogliom as and mixed gliomas (Fig. 1).

MATERIALS AND METHODS

Study outline: The pre-processing step (image noise removal) was performed using a wavelet domain filtering square magnitude image software. Removal of noisy is necessary from the data in order to proceed with further data analysis. In literature, many methods are mentioned for removal of noise. It is generally classified into two categories: denoising in the original signal or image domain (e.g., time or space) and denoising in the transform domain (e.g., Fourier or WT). Before using a merged GA and PSO algorithm for feature selection. Finally, the Marr-Hildreth was used for tumor boundary and area calculation (Fig. 2).

Image processing and feature extraction

Image pre-processing: The image was pre-processed using wavelet domain filtering square magnitude image processor for noise removal. This method was preferred in this study because the orthogonal wavelet transforms the power of the preferred image signal into a small number of coefficients. Hence, the transformed image consists of a small number of coefficients with high SNR that was kept and many coefficients with low SNR which were discarded. Having discarded the noisy coefficients, the image was reconstructed using the inverse wavelet transform. With this method, the image can be filtered from the noise (Nowak, 1999; Donoho, 1993). The wavelet domain filtering square image algorithm is as follows

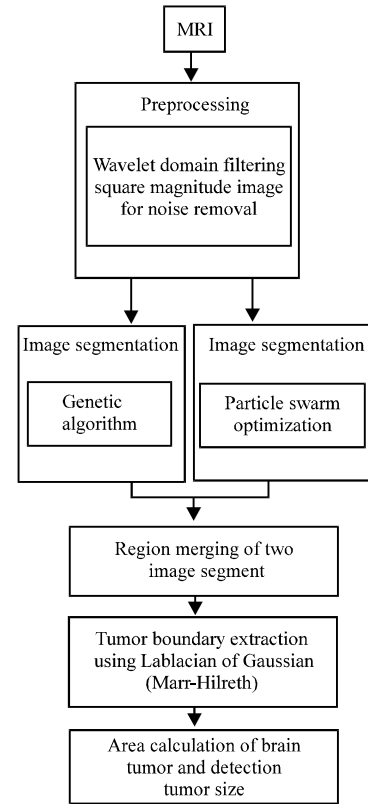


Fig. 2: Block diagram of a purposed system

(Nowak, 1999) compute the J-scale DWT and DWST of the squared magnitude image of $\sigma^2 I$. Form estimates of the variances of each wavelet coefficients:

$$d_i, d_i = \sum_m W_i[m] s[m], d_i = \sigma_i d_i$$

$$\sigma_i^2 = 4\sigma^2 \text{MAX} \left[\left(\sum_m w^2[m] x^2[m] \right) - \sigma^2, \sigma^2 \right], \sigma^2 \text{ is } 4\sigma^4$$

According to is used todistinguish this d_i note that wavelet ccoefficient from the wavelet coefficient corresponding to the magnitude image $x[m]$. Underlying discretetime wavelet basic w_i^k function at scale K and location I . Filter the wavelet coefficients according to:

$$\frac{d_i^2}{d_1^2} - \frac{T^2 \sigma_i}{d_1^2}$$

Remove the basic from the scaling coefficients by subtracting C_i from each:

$$C = 2^{(j+1)\sigma^2}$$

Obtain an estimate of S^2 by computing the inverse DWT of the filtered wavelet and scaling coefficients. Use the pixel-by-pixel square root of the result to get an estimate of s . It should be noted that d_1 was used to differentiate this wavelet coefficient from that was used to differentiate this wavelet coefficient from that of d_1 which corresponds to the magnitude image:

$$X[m]d_1 = \sum_m w_l[m]s[m], d_1 = \delta_1 d_1, w_l^k$$

De note the underlying discrete time wavelet basic function at scale k and location.

Feature extraction: This process identifies the feature characteristics that shows the predictable tumor in the brain through the mass of the output image from the image processing software. There are two stages in this process, the first stage is the segmentation of the image mass using GA. The segmentation technique can be classified into classical algorithms which are mainly based on statistical or mathematical methods, optimization-based techniques and artificial intelligence-based techniques (Dagar and Dahiya, 2016). The classical algorithms comprise characteristic histogram threshold, edge/boundary detection, region extraction or region increasing, recreation, semantic and syntactic move toward. The segmentation techniques are applied AI methods that use artificial neural networks. The GA is mainly used in image segmentation for the modification of the existing parameters in the segmentation algorithm and for segmentation at pixel level (Holland, 1992). GA can determine the optimal number of segmentation result regions and can choose some features such as the heuristic thresholds or the size of the analysis window (Back and Kursawe, 1995; Farmer and Shugars, 2006). The GA is robust and can be applied in different fields due to its basic functionality. Other algorithms in the same class with the GA include Evolutionary Strategy (ES), Genetic Programming (GP) and Evolutionary Programming (EP). A major difference between GA and other conventional optimization methods its ability to use a population of points at the same time in contrast to the single point approach used by the conventional optimization methods. AGA presents the genetic representation of the solution domain as well as a fitness function for the evaluation of the solution domain.

The second stage of the feature selection process is determining the mass of the image using PSO. Digital image segmentation involves the unravelling of a digital image into its various sections. In the image dispensation system it is a complex but necessary task. Image

segmentation is performed to present the image in an obvious way. It is usually used to dissipate an image into regions that ideally correspond to different real-world substances. Image segmentation is a risky step towards image and content psychoanalysis. Its result is a collection of segments that converge to form the real image. There are several problems of real globe image segmentation including feature minimization, overall deviation reduction, classifier error rate reduction and connectivity maximization. The PSO was developed by Kenney and Eberhart (Bhattacharyya, 2011; Gaurav and Bansal, 2013). An evolutionary calculation technique after merge by merge area using Marr-Hildreth to the output of merge two phase of segmentation using region merging algorithm.

Genetic algorithm: The consists of the choosing the initial individual population, evaluating the fitness of the individuals in the population and repeat this process until the termination criteria (time limit, sufficient fitness achieved) is met. The subsequent steps include selecting the best-fit individuals for reproduction, breeding new individuals through mutation and crossover operations, evaluating the fitness of the new individuals and replacing the least-fit individuals with the best-fit new individuals.

Image segmentation using GA: As per (Farmer and Shugars, 2006) there are two types of Genetic algorithms for image segmentation-parameter selection which involves the parameters of an existing image segmentation technique is modified using GA to improve its output and pixel-level segmentation which involves the use of GA to perform region labelling. Most methods of image segmentation have parameters that demands to be optimized, hence, the first method is often used (Bhattacharyya, 2011).

Practical Swarm Optimization (PSO): The PSO is a class of swarm-based intelligence techniques used to handle optimization issues. The PSO mimics the flocking behaviour of bird where a group of birds randomly search an area for food and in this area there is just a piece of food being searched. The birds have no idea of the position of the food but they know about its location in the search space. In this situation, the ideal way to locate the food is to either forage or flock. The location of each particle in the swarm is updated following two “best” values-best and g-best with the p-best, each particle keeps track of its coordinates in the search space with respect to the best solution (fitness) it has so far achieved and hence, the name p-best or personal best. The g-best

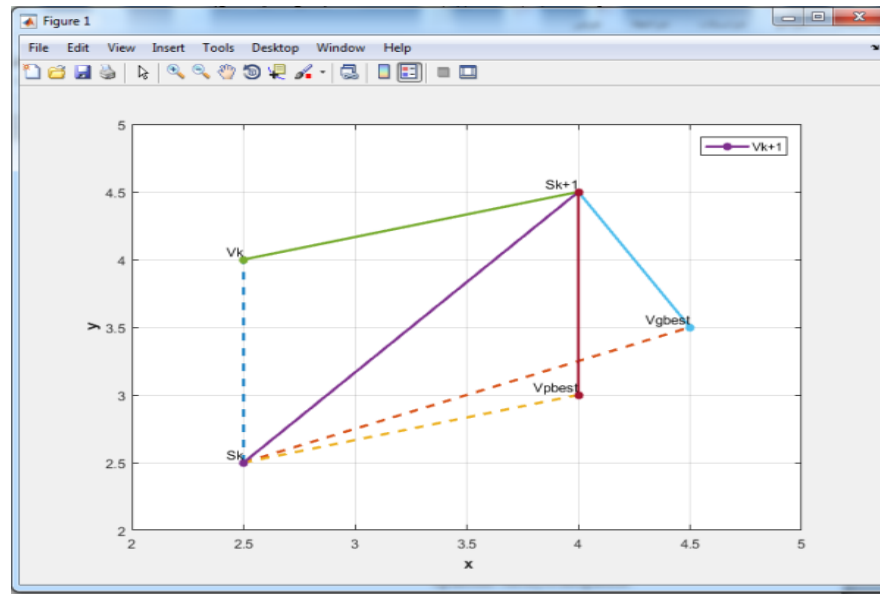


Fig. 3: Concept of modification of a searching point by PSO

is tracked by the PSO as the best value, so far obtained by any particle within the neighbourhood of the particle and hence, the name g-best or global best. The position of each particle is updated using their current positions and velocities the distance between the p-best and the current position as well as the distance between g-best and the current position. Having established the two best values, the velocity and position of the particle are updated using the following Eq. 2 (Bhattacharyya, 2011):

$$v[] = v[] + 1 * \text{rand}() * (pbest[] - \text{ppresent}[]) + c_2 * \text{rand}() * (gbest[] - \text{ppresent}[]) \quad (2)$$

$$(1) \text{present}[] = \text{present}[] + v[]$$

Where:

S^K = Current Searching point
 S^{K+1} = Modified Searching point
 V^K = Modified Velocity
 V^{pbest} = Velocity based on pbest
 V^{gbest} = Velocity based on gbest

where, $v[]$ is the velocity of the particle, $\text{present}[]$ is the current particle (solution). $\text{rand}()$ is a random number between (0; 1). $c_{1,2}$ are learning factors usually $c_1 = c_2 = 2$. The concept of modification of a searching point by PSO algorithm is depicted in Fig. 3 (Bhattacharyya, 2011).

In Fig. 3, it shows the concept of modification of PSO. Each particle moves from the current position to the next one according to the present values. Generally, the fitness function is same the objective functions. The local

best of other particles in the population should be changed if the present fitness function value is better than the previous. The algorithm of the PSO is as follows (Gaurav and Bansal, 2013):

Algorithm 1: PSO; Gaurav and Bansal (2013)

Step 1: Let's represent the number of particles in the swarm with each particle having a position X_i and velocity v_i in the search space. Let's represent the best-achieved position of particle i and g be the best for the entire swarm. Basic PSO algorithm is executed thus

Step 2: For each particle $i = 1, \dots, S$, perform the following: 1 use a uniformly distributed random vector $x_i \sim U(blo, bup)$ to initialize the position of the particle where blo and bup represent the lower and upper limits of the solution space

- 2 Initialize the best-known position of the particle to its initial position: $p_i = x_i$ If $(f(p_i) < f(g))$
- 3-Update the best-known position of the swarm p_i
- 4-Initialize the velocity of the particle

Step 3: Repeat the following steps until the termination criteria (such as the number of iterations performed or have found a solution with adequate objective function value) is met

For each particle $i = 1, \dots, S$ do

For each dimension $d = 1, \dots, n$ do

Pick random numbers: $rp, rg \sim U(0, 1)$

Update the velocity of the particle

Update the position of the particle: $x_i = x_i + v_i$

If $(f(x_i) < f(p_i))$ do

Update the most excellent known position of the particle: $p_i = x_i$

If $(f(p_i) < f(g))$, update the best-identified position of the swarm: $g = p_i$

Step 4: g is now the best found solution. The parameters, ϕp and ϕg are chosen by the practitioner controlling the legacy and behavior of the PSO

Region merging: An appropriate seed selection is a significant consideration during the segmentation process. The selected seed pixels must (Kansal and Jain, 2015). Be highly similar to its neighbouring pixels. At least one seed pixel must be generated for each expected image region. There should be a connection between the seeds

of different image regions. In the proposed approach, the occurrence frequency of the seeds after merging was considered during seed selection. For each image, the seeds were computed using the following algorithm 2.

Algorithm 2; PSO base algorithm:

Step 1: Compute the image grey levels and sort them in ascending order
 Step 2: Compute the frequency of the soft image grey levels
 Step 3: Assign the first pixel as the first seed point
 Step 4: Merge the pixels to get an improved seed point. If the merging conditions are met, replace the seed point by the mean of the merged pixels. Sum up the frequencies of the pixel and the seed point, assign the position of the seed point to be equal to the initial seed point, otherwise, assign the pixel as a new seed point. Repeat step 4 until all the pixels are unmerged
 Step 5: Sort the final seed points based on their occurrence frequency in descending order. Thus, the final seed points achieved are used for image segmentation in the seeded region growing algorithm

Marr-Hildreth: Edge detection is a basic image processing and computer vision technique. It detects the areas in a digital image where there are sharp changes in the image brightness or where there are discontinuities (Tandan *et al.*, 2014). It is used for noise filtration before edge discovery. The Marr-Hildreth method is a combination of the Gaussian filtering and Laplacian edge detection techniques.

Definition of the components: Laplacian is a measure of an image's second spatial derivative and is a useful tool in detecting abrupt changes in images. Gaussian smoothing is performed in image edging before Laplacian to annul the noise effect. The operations are linear and can be interchanged. Gaussian smoothing is a unique form of weighted smoothing where the smoothing kernel coefficients are derived from a Gaussian distribution. The 2-DLaplacian of Gaussian function centred on zero and with Gaussian standard deviation has the form a variation of the value of the standard deviation controls the amount of smoothing.

Calculate area of tumor: Apply the algorithm of find area (Suman *et al.*, 2017) of tumor image as shown in Fig. 4 and 5.

Algorithm 3; Run area of the tumor:

Input: Tumor portion of the image
 Output: Area of the tumor
 Step 1: Read the input grayscale image
 Step 2: Resize this image into $n \times n$ image matrix
 Step 3: Compute numbers of rows and ccolumn in pixels by $[r_2 \ c_2] = \text{size}(J)$
 Step 4: Initialize a variable $a = 0$
 Step 5: For $I = 1:1:r_2$
 Step 6: For $j = 1:1:c_2$
 Step 7: If $I(I, j) == 255$
 Step 8: $a = a + 0$
 Step 9: Else do
 Step 10: $a = a + 1$
 Step 11: End
 Step 12: End
 Step 13: End
 Step 14: Display the area a

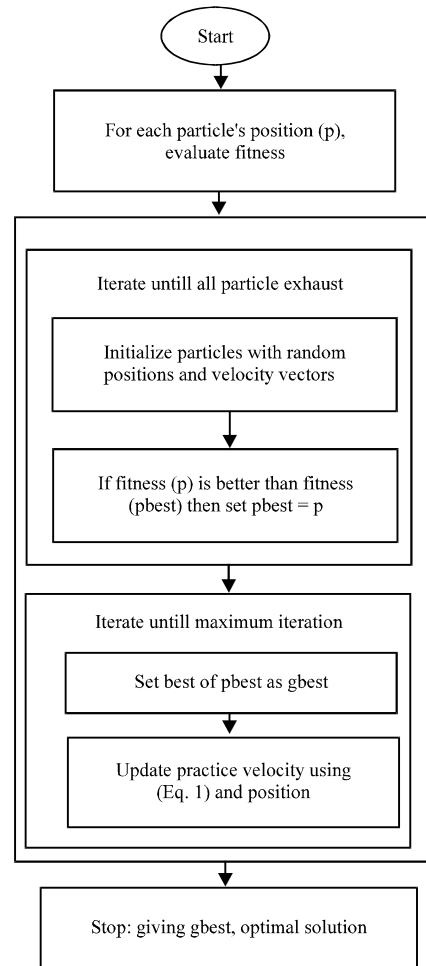


Fig. 4: Flowchart of the PSO base algorithm

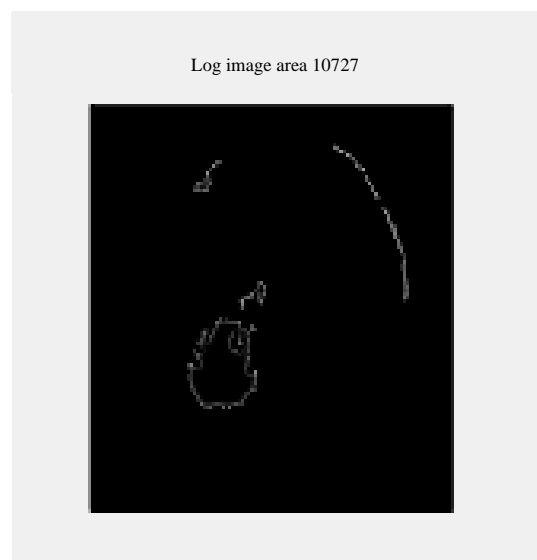


Fig. 5: Run area of the tumor

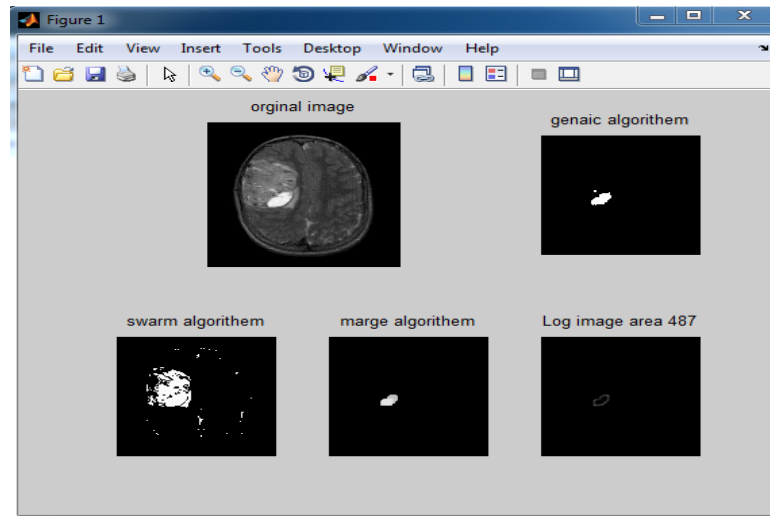


Fig. 6: Run of all work

TP	TN	FP	FN	Total
3	25	2	0	30
Metrics	Values	LAW		
Accuracy	0.93333	(TP+TN)/Total		
Misclassification error	0.06	(FP+FN)/Total		
Sensitive P	1	TP/TP+FN		
Sensitive N	0.9259	TN/TN+FP		
Precision (PPV)	0.6	TP/TP+FP		
NPV	1	TN/TN+FN		

Fig. 7: Calculate the metrics for run program

RESULTS AND DISCUSSION

Images set: The presentation of brain tumor was assessed using a merged GA and PSO. The dataset was acquired through magnetic resonance imaging where this search using 30 image for testing the detect tumor in magnetic resonance imaging.

Evaluation performance factors: The of the proposed technique's performance was evaluated using a set of examination data. Based on this, the output image for detecting a tumor in the brain is shown in Fig. 6 calculate metrics to perform the program of this thesis by calculating accuracy that shows in Fig. 7 to compare with future or old thesis to see a good performance of the thesis. PPV is positive predictive value NPV is Negative

Predictive Value. TP is genuinely positive, TN is a real negative, FP is False Positive and FN is a False Negative.

CONCLUSION

Genetic algorithm has many compensations in getting the optimized reply. It was shown to be the majority influential optimization method in a large space. The Genetic algorithm allows to carry out healthy. The outcome of the optimizations depend on the genetic material indoctrination system and participation of genetic employee as well as on the strength function. However, the fineness of image segmentation can be better by choosing the parameters in an optimized method. While PSO is the usual method of calculating and give a number of habits to decide real-world harms more professionally and rapidly with correctness.

RECOMMENDATION

In feature, research can use one of the previous methods of segment remember in study with other develop method of the segment and deploy to detect a tumor in the brain.

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