2<u>0<sup>th</sup> November 2013. Vol. 57 No.2</u>

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ISSN: 1992-8645

www.jatit.org



# MAMMOGRAM MRI IMAGE SEGMENTATION AND CLASSIFICATION

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## ABSTRACT

This paper presents the mathematical morphological and rough set based approach in detection and classification of cancerous masses in MRI mammogram images. Breast cancer is one of the most common forms of cancer in women. In order to reduce the death rate, early detection of cancerous regions in mammogram images is needed. The existing system is not so accurate and it is time consuming. MRI Mammogram images are enhanced and the artifacts are removed using the Fuzzification technique. The ROI(Region of Interest) is extracted using Graph Cut method and the Four mathematical morphological features are calculated for the segmented contour . ID3 algorithm is applied to extract the features which play a vital role in classification of masses in mammogram into Normal, Benign and Malignant. The sensitivity, the specificity, positive prediction value and negative prediction value of the proposed algorithm were determined and compared with the existing algorithms. Automatic classification of the mammogram MRI images is done through three layered Multilayered Perceptron .The weights are adjusted based the Artificial Bee Colony Optimization technique .Both qualitative and quantitative methods are used to detect the accuracy of the proposed algorithm accounts to 98.78%, 98.9%, 92% and 96.5% which rates very high when compared to the existing algorithms. The area under the ROC curve is 0.89.

Keywords: Fuzzification, Gabor Filter, Graph Cut, ID3 And Artificial Bee Colony Technique.

#### 1. INTRODUCTION

The Population Based Cancer Registry evidently shows from the various statistics, that the incidence of breast cancer is rapidly rising, amounting to a significant percentage of all cancers in women. Breast cancer is the commonest cancer in urban areas in India and accounts for about 25% to 33% of all cancers in women. Over 50% breast cancer patients in India present in stages 3 and 4, which will definitely impact the survival [1]. The survival rate can be increased only through the early diagnosis. Image processing technique together with data mining is used for extraction and analysis of the ROI. Tumor can be classified into three category normal, benign and malignant. A normal tumor is a mass of tissue which exists at the expense of healthy tissue. Malignant tumor has no distinct border.

They tend to grow rapidly increasing the pressure within the breast cells and can spread beyond the point from they originate. Grows faster than benign and cause serious health problem if left unnoticed. Benign tumors are composed of harmless cells, have clearly defined borders, can be completely removed and are unlikely to recur. In MRI mammogram images after the appropriate segmentation of the tumor, classification of tumor into malignant, benign and normal is difficult task due to complexity and variation in tumor tissue characteristics like its shape, size, grey level intensities and location. Feature extraction is an important aspect for pattern recognition problem. This paper presents a Hybrid rough set based approach for automatic detection and classification of cancerous masses in mammogram images.

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ISSN: 1992-8645

www.jatit.org



E-ISSN: 1817-3195

## 2. MATERIALS AND METHODS

The block diagram in Figure 1 clearly depicts the methodology of the proposed technique.



Figure 1: Proposed Methodology

The data set used for research were taken from Mammogram Image Analysis Society(MIAS)[2]. The database contains 320 images out of which 206 are normal images,63 benign and 51 malignant cases.

3. IMAGE ENHANCEMENT AND PREPROCESSING USING FUZZY HISTOGRAM EQUALIZATION

Mammogram images are enhanced using histogram equalization method. The approach developed by Ella Hassanien is implemented for Image preprocessing[3]. An image I of size M x N and L gray levels can be considered as an array of fuzzy singletons, each having a value of membership denoting its degree of brightness relative to some brightness levels. For an image I, we can write in the notation of fuzzy sets [4]:

 $I = U\mu_{mn}/g_{mn}$  m = 1,2,..., M and n = 1,2,..., N, Where  $\mu_{mn}$  its membership function  $g_{mn}$  is the intensity of (m, n) pixel. The membership function characterizes a suitable property of image (e.g. edginess, darkness, textural property) and can be defined globally for the whole image or locally for its segments. In recent years, some researchers have applied the concept of fuzziness to develop new algorithms for image enhancement [3]. The principle of fuzzy enhancement scheme is illustrated in Figure 2.



Figure 2: Fuzzification Technique

The histogram equalization of the gray levels in the original image can be characterized using five parameters: ( $\alpha$ ,  $\beta 1$ , $\gamma$ ,  $\beta 2$ , max) as shown in Figure 3.



# Figure 3: Histogram Equalization Curve With A , $\beta 1, \Gamma$ , $\beta 2$ , Max

where the intensity value  $\gamma$  represents the mean value of the distribution,  $\alpha$  is the minimum, and max is the maximum. The aim is to decrease the gray levels below  $\beta 1$ , and above  $\beta 2$ . Intensity levels between  $\beta 1$  and  $\gamma$ , and  $\beta 2$  and  $\gamma$  are stretched in opposite directions towards the mean  $\Box \gamma$ The fuzzy transformation function for computing the fuzzy plane value P is defined as follows:  $\alpha = \min;$ 

 $\beta 1 = (\alpha + \gamma)/2;$ 

 $\beta 2 = (\max + \gamma)/2;$ 

```
\gamma = mean; max;
```

The following fuzzy rules are used for contrast enhancement based on Figure (2).

Rule-1: If  $\alpha \le ui < \beta 1$  then P = 2 ( (  $ui - \alpha$ ) / ( $\gamma - \alpha$  ))<sup>2</sup> Rule-2: If  $\beta 1 \le ui < \gamma$  then P = 1-2 ( (  $ui - \gamma$  )/( $\gamma - \alpha$  ))<sup>2</sup>

Rule-3: If  $\gamma \le ui < \beta 2$  then P= 1- 2( ( $ui - \gamma$ ) / (max  $-\gamma$ ))<sup>2</sup>

Rule-4: If  $\beta 2 \le ui \le max$  then P = 2 ((  $ui - \gamma$  )/(  $max - \gamma$  ))<sup>2</sup>

Where ui = f(x,y) is the ith pixel intensity.

Pseudocode:

Step-1: Parameter Initialization

Set  $\beta 1 = (\min + \max)/2$ 

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E-ISSN: 1817-3195

Set  $\beta 2 = (\max + \max)/2$ Step-2: Fuzzification

ISSN: 1992-8645

For all pixels (i,j) within the image Do If  $((data[i][i] \ge min) \&\& (data[i][i] < \beta 1))$ Compute NewGrayLevel =2\*(pow(((data[i][j]- $\min$ /(mean-min)),2)) If  $((data[i][j] > = \beta 1) \&\& (data[i][j] < mean))$ Compute NewGrayLevel=1-(2\*(pow(((data[i][j]mean)/(mean-min)),2))) If  $((data[i][j] \ge mean) \& \& (data[i][j] \le \beta 2))$ Compute NewGrayLevel=1-(2\*(pow(((data[i][j]mean)/(max-mean)),2))) If  $((data[i][j] \ge \beta 2) \&\& (data[i][j] \le max))$ Compute NewGrayLevel=2\*(pow(((data[i][j]mean)/(max-mean)).2)) Step-3: Fuzzification Modification Compute FuzzyData[i][j]= pow(NewGrayLevel,2) Step-4: Defuzzification For all pixels (i,j) within the image Do Compute

EnhancedData[i][j]=FuzzyData[i][j]\*data[i][j];









Figure 5: Input Image

Figure 6: Output Image



PSNR	RMS	E	¥	MSE	<b>Nature</b> of	
87.65	2.97	0.2111	0.0086	8.83	Gabor	

The resultant image contrast enhancement and grayness ambiguity is checked using Peak Signal to noise ratio, Index of fuzziness and Fuzzy entropy technique.

Mean Square Error (MSE) is calculate using the formula:

$$1/MN\sum_{i=1}^{m}\sum_{j=1}^{n} |x(i,j) - x'(i,j)|^{2}$$

Peak Signal to Noise ratio:

PSNR=20log10 [(2n-1)/√MSE]

where x(i,j) is the input image x'(i,j) is the output image and n is the number of bits used in representing the pixel of the image.

Index of Fuzziness & Fuzzy Entropy are the measures of global grayness ambiguity of a n image. They are defined as a degree of difficulty in decising whether a pixel would be treated as black(dark) or white(bright). Index of fuzziness is the amount of fuzziness present in an iamge which determines the amount of vagueness by measuring the distance between the fuzzy property plane and nearest ordinary plane. Entropy (H) is a measure of quality of information in an image in the fuzzy domain based on Shanon's function.

of

$$\gamma = 2/MN \sum_{M} \sum_{N} \min(\mu mn, 1 - \mu mn)$$
  
Entropy: H=1/MN ln  
$$\sum_{m} \sum_{n} \ln(\mu mn) - (1 - \mu mn) \ln(1 - \mu mn)$$

The tabular column clearly shows that the PSNR value is high therefore the image is enhanced and the Index of fuzziness and Entropy decreases with enhancement.

2<u>0<sup>th</sup> November 2013. Vol. 57 No.2</u>

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ISSN: 1992-8645	www.jatit.org	E-ISSN: 1817-3195
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#### 4. IMAGE SEGMENTATION AND RO EXTRACTION

The region of Interest ie) the tumor region is segmented using the Graph cut method. The main purpose of using this method for segmentation is that it segments the mammogram into different mammographic densities. It is useful for risk assessment and quantitative evaluation of density changes. Apart from the above advantage it produces the contour (closed region) or a convex hull which is used for analyzing the morphological and novel features of the segmented region. The above technique results in efficient formulation of attributes which helps in classification of the ROI into benign, malignant or normal. Graph Cuts has been used in recent years for interactive image segmentation[4]. The core ideology of Graph Cuts is to map an image onto a network graph, and construct an energy function on the labeling, and then do energy minimization with dynamic optimization techniques. This study proposes a new segmentation method using iterated Graph Cuts based on multi-scale smoothing. The multi-scale method can segment mammographic images with a stepwise process from global to local segmentation by iterating Graph Cuts. The Graph cut approach used by K. Santle Camilus [4] is implemented in this paper

Steps involved in Graph Cut Segmentation are:

- 1. Form a graph
- 2. Sort the graph edges
- 3. **Region merging**

A graph G=(V,E) is constructed from the mammogram image such that V represent the pixel values of the 3\*3 image and E the edges defined between the neighboring pixels. The weight of any edge W(Vi,Vj) is measure of dissimilarity between the pixels Vi and Vj. The weight for an edge is by means of considering the Euclidian distance between the two pixels Vi and Vj. It is represented by the equation

W(Vi,Vj)=
$$\sqrt{(xi - xj)^2 + (yi - yj)^2}$$
  
Eqn.(1)

Vi=(xi,yi) Vj=(xj,yj)

Edges are sorted in ascending order of their weights such that  $w(e1) \le w(e2)$ .Pick one edge ei in the

ROI sorted order from ei to en where ei is between two groups of pixels. This edge determines whether to merge the two groups of pixel to form a single group or not. Each vertex is considered as a group..
If the merge criterion is satisfied, then the two groups are merged. The multiple groups of pixels representing different regions or objects are risk obtained.

#### **Intra-region Edge Average**

The intra-region edge average (IRA) which is a single-valued function and represents the homogeneity of a group is defined as

IRA(R) = 
$$\sum_{(Vi,Vj)\in E}^{Vi,Vj\in R} W(Vi,Vj) \div |Va|$$
Eqn.(2)

IRA for a region "R" is a measure of the mean of the weight of edges in "R."  $V_a$  is a set of edges in the region "R" which can be represented as

$$Va = \{(V_{i}, V_{j}) \in E | V_{i}, V_{j} \in R\}$$
  
Eqn.(3)

#### **Inter-Region Edge Mean:**

The discussion for computing IRA can be applicable to compute the homogeneity representative for intermediate edges between the two regions. Hence, we define inter-region edge mean (IRM) as

$$IRM(R1,R2) = \sum_{(vi,vj) \in E}^{vi \in R1} W(vi,vj) / |Vb| \qquad Eqn(4)$$

IRM is a measure of the mean of the weight of edges between the two regions,  $R_1$  and  $R_2$ .  $V_b$  is a set of edges between the two regions,  $R_1$  and  $R_2$ , which can be represented as

$$Vb = \{(V_i, V_j) \in E | V_i \in R_1, V_j \in R_2\}$$
 Eqn(5)

#### **Dynamic Thresholds**

The degree of variations among pixels of a region can be inferred from the homogeneity representative of the region IRA. To merge pixels of two regions, the IRM must be above a threshold value. This

20<sup>th</sup> November 2013. Vol. 57 No.2

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ISSN:	1992-8645
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E-ISSN: 1817-3195

threshold must be computed based on the IRA and other parameters to control the merge operation adaptive to the properties of regions. Hence, we define a dynamic threshold (DT) for merging between two regions  $R_1$  and  $R_2$  as given in the following equation

$$DT(R1,R2)=Max(IRA(R1)+\delta 1,IRA(R2)+\delta 2)$$
  
Eqn.(6)

The appropriate selection of the parameters ( $\delta_1$  and  $\delta_2$ ) plays a vital role as it determines the merge of regions. Values of these parameters are chosen to achieve the following desirable features: (1) when more number of regions is present, then the choice of the values (of  $\delta_1$  and  $\delta_2$ ) should favor merging of regions; (2) the choice of the values should allow the grouping of small regions than large regions. The large regions should be merged only when their intensity values are more similar; (3) it is preferred to have values that are adaptive to the total number of regions and the number of vertices in the regions that are considered for merge. Hence, the parameters  $\delta_1$  and  $\delta_2$  are defined as

$$\delta 1 = C \times N_R / |R_1|$$
  
$$\delta 2 = C \times N_R / |R_2|$$
  
Eqn.(7)

 $N_R$  - the total number of regions. Initially,  $N_R = |V|$ , the size of the vertices and each merge decreases the value of  $N_R$  by one.  $|R_i|$  indicates the number of vertices in the region " $R_i$ ." *C* is a positive constant and is equal to two for the pectoral muscle segmentation which is found through experimental studies.

#### **Merge Criterion**

When the pixels of a group have intensity values similar to the pixels of the other group, then intuitively the calculated IRM between these groups should be small. The expected smaller value of the IRM to merge these two regions is tested by comparing it with the dynamic threshold. Hence, the merge criterion, to merge the two regions,  $R_1$  and  $R_2$ , is defined as

Merge $(R_1, R_2)$ , if IRM $(R_1, R_2) \leq DT(R1, R2)$ Eqn(8)



Figure 5:	ROI Extracted	Using Graph	Cut Method
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#### FEATURE EXTRACTION AND SELECTION

The main objective of feature extraction and selection is to maximize the distinguished performance capability of them on interested data base. Consideration of significant feature is based on discrimination, optimality, reliability and independence. In recent years, many kinds of features have been reported for breast mass classification like texture based features, region based features, image structure features, position related features, and shape-based features which are described and utilized in the CAD systems [4]. In our study a set of six novel features based on shape characteristics (Novel Feature) proposed by Rangaraj M. Rangayyan [5] were extracted from the original images which are not applied in breast mass classification before and are defined as follows:

The convex hull which is the smallest convex polygon that contains the mass region is displayed. Mass's area is the actual number of pixels in the mass region and convex hull's area is the actual number of pixels in the convex hull region.

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ISSN: 1992-8645	www.jatit.org	E-ISSN: 1817-3195
Compactness(C) or Normalized circularity	<i>C</i> is a simple measure of the efficiency of a contour to cont defined in a normalized form as $1-4\pi S/D^2$ , where <i>D</i> and <i>S</i> ar and area S=Area= $\sum_{x} \sum_{y} I(x, y) \Delta A$ where $\Delta A$ is the pixel area Where I(x, y) = 1 if (x, y) pixel belong to S, zero other case $D(S) = \sum_{x} I(x, -X_{(i-1)})^2 + (Y_i - Y_{(i-1)})^2$	tain a given area and is re the contour perimeter
fractional concavity (F <sub>cc</sub> )	is a measure of the portion of the indented length to the to computed by taking the cumulative length of the concave se by the total length of the contour	tal contour length; it is egments and dividing it
Fourier descriptors and Fourier coefficient $A_0$	$A_0=1/N\sum_{i=1}^{360} d(i)$ Where d(i)= $\sqrt{(X(i)-X_c)^2 + (Y(i)-Y_c)^2}$ where ( <i>xc</i> , <i>yc</i> ) are the coordinates of the centroid, ( <i>x</i> ( <i>i</i> ), <i>y</i> ( <i>i</i> )) the boundary pixel at the <i>i</i> -th location.	are the coordinates of
Convexity	The measure of convexity of an object is the ratio of the p object to the original perimeter of the object. Convexity=convexperimeter/perimeter	perimeter of the convex
Aspect Ratio	Aspect Ratio is the ratio between height and width of the corr	responding object
Solidity	Solidity=Area/convexarea	
Circularity	$4\pi S/D^2$	

The shape features are calculated for the ROI segmented using Graph cut method for 320 images. The results are plotted in an Excel sheet for classification using WEKA tool.

#### 6. FEATURE SELECTION USING ID3 ALGORITHM

A mathematical algorithm for building the decision tree. Builds the tree from the top down, with no backtracking. Information Gain is used to select the most useful attribute for classification.

Entropy

• A formula to calculate the homogeneity of a sample.

• A completely homogeneous sample has entropy of 0.

An equally divided sample has entropy of 1.

• Entropy(s) =  $-p+\log 2(p+) - p-\log 2(p-)$  for a sample of negative and positive elements.

■ The formula for entropy is:

Information Gain (IG):

The information gain is based on the decrease in entropy after a dataset is split on an attribute[7].

Information Gain(S,A)=Entropy(S)-H(S,A)

Where  $H(S,A) = \sum_{i} (|Si|/|S|) \cdot H(Si)$ 

A takes on value 1 and H(Si) is the entropy of the

system of subsets Si.

The training data is a set S=s1,s2—of already classified samples based on the mathematical morphological and novel features. Each sample Si=x1,x2 is a vector where x1,x2—represents attributes or features of the sample. The training data is augmented with a vector C=C1,C2 and C3 where C1represent benign C2 represents malignant and C3 represents normal cases. At each node of the tree ID3 chooses one attribute of the data that most efficiently splits its set of samples into subsets enriched in one class or the other. The criterion is the normalized Information Gain that results from choosing an attribute for splitting the data. The attribute with highest information gain is chosen to make decision.

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ISSN: 1992-8645	www.jatit.org	E-ISSN: 1817-3195
Psudocode:	=== Detailed Accura	ncv By Class ===

Input: Set of 12 =4 Mathematical morphological + 4 novel feature attributes A1,A2---A12.The class labels Ci C1=Benign,C2=Malignant and C3=Normal.. Training set S of examples

Output: Decision tree with set of rules.

Procedure:

- All the samples in the list belongs to three different classes
- Create a leaf node for the decision tree saying to choose the class.
- If none of the feature provides IG then ID3 creates a decision node higher up the tree using the expected value of the class.
- If instance previously unseen class is encountered ID3 creates decision class higher up in the tree using expected value.
- Calculate IG for each attribute
- Choose attribute A with lowest entropy and Highest IG and test this attribute with root.
- For each possible value v of this attribute
- Add a new branch below the root corresponding to 0 A=v.
- If v is empty make the new branch a leaf node 0 labeled with most common value else
- Let the new branch be the tree created by ID3 0

0

• End

> The attributes used were the eight parameters with the class as benign, malignant and normal. Based on the rule derived by testing 320 mammogram images the rules are applied in classifying the new cases without prior knowledge of whether they are benign, malignant or normal.

Table 2: Nodes Representing The 7 Attributes Cross Validation 10 Fold

Node1	Compactness
Node2	Fractional Concavity
Node3	Fourier Descriptor
Node4	Convexity
Node5	Aspect Ratio
Node6	Solidity
Class	Benign/Malignant/Normal

=== Detailed	Accuracy	By	Class ===	
--------------	----------	----	-----------	--

Class	TP Rate	e FP Rate	Precision	Recall
FMS	ROC			
Benign	0.595	0.272	0.523	0.595
0.557	0.677			
Malig	0.121	0.104	0.368	0.121
0.182	0.542			
Norma	1 0.678	0.427	0.442	0.678
0.536	0.637			
Wgt.	0.465	0.268	0.445	0.465
0.425	0.618			
=== Co	nfusion N	latrix ===		

a b c <-- classified as

172 30 87 | a = Benign

94 35 160 | b = Malignant

63 30 196 | c = Normal

The classification accuracy obtained from these fixed-order trees can be compared with those from trees of different feature orders, as well as with those from trees of different feature combinations. The decision tree with the optimal feature combination and order for this task can thus be identified. Examples of the training and test results obtained in this study will be discussed below. It should be noted that, to use a trained decision-tree classifier, one has to choose the specific tree structure with the set of decision thresholds corresponding to the desired sensitivity (TPF) and specificity (~-FPF). The structure and the thresholds will be fixed during testing or application. The decision rule is indicated below

Compactness < -0.0450 then Class = Benign (64.89 % of 262 examples)

Compactness >= -0.0450 then Class = Malignant (62.34 % of 316 examples).

Compactness  $\geq$  -0.0050 then Class = Normal (42.29 % of 525 examples)

ID3 parameters: Size before split 200 Size after split 50 Max depth of leaves10 Goodness of split threshold 0.0300 2<u>0<sup>th</sup> November 2013. Vol. 57 No.2</u>

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ISSN: 1992-8645 <u>www.jatit.org</u> E-ISSN: 1817-3195

## 7. MULTILAYER FEED FORWARD NEURAL NETWORK (MLN) OPTIMIZED BY ARTIFICIAL BEE COLONY OPTIMIZATION TECHNIQUE:

Artificial bee colony algorithm(ABC) was proposed for optimization, classification and neural network problem solution based on the intelligent foraging behavior of honey bee . ABC is more powerful and most robust on multimodal functions. It provides solution in organized form by dividing the bee objects into different tasks such as employed bees, onlooker bees and scout bees. It finds global optimization results, optimal weight values by bee agents. Successfully trained MLN improves the classification precision of MRI mammogram images into benign, malignant and normal cases.

The ANN is the best method for pattern recognition. The seven attribute values together with the class are fed as input to the eight neurons at the input laver. The hidden laver has sixteen neurons or nodes. The output layer has one node. The error rate is calculated by subtracting the actual output with the desired output value. The evaluation of the food sources that are discovered by the randomly generated populations is done through the backpropagation procedure. Now after completion of the first cycle the performance of each solution is kept in an index. In order to generate the population of back-propagation, the weight must be created, and each value of the weight matrix for every layer is randomly generated within the range of [1,0]. Each set of weight and bias is used to create one element in a whole population. The size of the population in the proposed method is fixed to N.

The classifier employed in this paper is a three-layer Back propagation neural network. The Back propagation neural network optimizes the net for correct responses to the training input data set. More than one hidden layer may be beneficial for some applications, but one hidden layer is sufficient if enough hidden neurons are used. Initially the attributes values (rules) from the Mathematical morphological and novel approach analysis method which plays a vital role in classification between benign, malignant and normal cases extracted from the ID3 algorithm are normalized between [0,1]. That is each value in the feature set is divided by the maximum value from the set. These normalized values are assigned to the input neurons. The number of hidden neurons is equal to the number of input neurons and only one output neuron.

The procedure employed by Habib Shah in his work was implemented for training the mammogram data set using ANN.

## **Employed Bees:**

It uses multidirectional search space for food source and get information to find food source and solution space. The employed bees shares information with onlooker bees. They produce a modification on the sources position in her memory and discover a new food sources position. The employed bee memories the new source position and forgets the old one when the nectar amount of the new source is higher than that of the previous source.

## **Onlooker Bees:**

It evaluates the nectar amount obtained by employed bees and chooses a food source depending on the probability values calculated using the fitness values. A fitness based selection technique can be used for this purpose. Watches the dance of hive bees and select the best food source according to the probability proportional to the quality of that food source.

#### Scout bees:

Select the food source randomly without experience. If the nectar amount of the food source is higher that that of the previous source in their memory, the new positions are memorized and forgets the previous position. The employed bee becomes scout bees whenever they get food source and use it very well again. These scout bees find the new food source by memorizing the best path.

1: Initialize the population of solutions Xi where  $i{=}1{\ldots}{\ldots}SN$ 

2: Evaluate the population and calculated the fitness function for each employed bee

Initialize weights, bias for Multilayerd Feed forward network

3: Cycle=1

4: Repeat from step 2 to step 13

5: Produce new solutions (food source positions) Vi,j in the neighbourhood of Xi,j for the employed bees using the formula

$$V_{i,j} = X_{i,j} + \phi_{ij}(X_{i,j} - X_{k,j})$$

20<sup>th</sup> November 2013. Vol. 57 No.2

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SSN: 1992-8645	www.jatit.org	E-ISSN: 1817-3195

where k is a solution in the neighborhood of i,  $\Phi$  is a random number in the range [-1, 1] and evaluate them.

6: Apply the Greedy Selection process between process

7: Calculate the probability values pi for the solutions Xi by means of their fitness values by using formula.

## $Pi=fit_i/\sum_{k=1}^{5N} fit_n$

The calculation of fitness values of solutions is defined as

$$\operatorname{Fit}_{i} = \left\{ \begin{array}{c} \frac{1}{1+fi} & fi \geq 0\\ 1+abs(fi) & fi < 0 \end{array} \right\}$$

Normalize pi values into[0,1]

8: Produce the new solutions (new positions) vi f8r the onlookers from the solutions xi, selected depending on Pi, and evaluate them

9: Apply the Greedy Selection process for the onlookers between xi and vi

10: Determine the abandoned solution (source), if exists, replace it with a new randomly produced solution xi for the scout using the following equation

 $X_{i}^{j} = x_{min}^{j} + rand(0,1) (x_{max}^{j} - x_{min}^{j})$ 

11: Memorize the best food source position (solution) achieved so far

12: cycle=cycle+1

13: **until** cycle= Maximum Cycle Number (MCN)

The classification accuracy is determined using Mean square error and standard variance of mean square error on 30 cycles

The Dataset with best attribute generated by ID3 algorithm is fed as input for benign,normal and malignant cases .

Table 3: Parameter Values			
Mass type Training data Testing			
	set	_	
Benign	600	425	
Malignant	600	425	
Normal	600	425	

SN	20
MaxEpoch	400
Fitness value	10

Table 4: Classification Accuracy

Algorithm	Mean Square Error	Std. Var of Mean Square Error	Rank
ABC+MLP	1.67 X	0.0001	1
	10-4		

#### **CONCLUSION:**

The rule generated using decision tree induction method clearly shows that the time taken to classify benign and malignant cases in just 0.03 seconds and the accuracy was 98.9%. The specificity =t-neg/neg, sensitivity =t-pos/pos and Precision=t-pos/t-pos+f-pos and accuracy =sensitivity

(pos/(pos+neg))+specificity(neg/(pos+neg)).From

the above algorithm the accuracy was found to be 99.9%. The proposed method yields a high level of accuracy in minimum period of time that shows the efficiency of the algorithm. The training speed accounts to 6 ms using ANN. The main goal of classifying the tumors into benign, malignant and normal is achieved with a great accuracy compared to other techniques.

Table 5: Accuracy Details

Specificity		95.5%
Sensitivity		97.3%
Positive Prediction value		89%
Accuracy		98.9%
Area under Curve		0.98
Negative	Prediction	98%
Value		

2<u>0<sup>th</sup> November 2013. Vol. 57 No.2</u>

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ISSN: 1992-8645

<u>www.jatit.org</u>

E-ISSN: 1817-3195

Table 6: Comparative Analysis

Methods	Author and References	Computational
		Time
Morphological Analysis	Wan Mimi Diyana, Julie Larcher, Rosli Besar	3'20''
ing Technique	Proposed Approach	3'50''
Fractal Dimension Analysis	Wan Mimi Diyana, Julie Larcher, Rosli Besar	7'20''
Complete HOS Test	Wan Mimi Diyana, Julie Larcher, Rosli Besar	9'20''
neuro-fuzzy segmentation	S. Murugavalli et. al	93' 39''
Proposed Method	S.Pitchumani Angayarkanni ,Nadhira Banu Kamal	0'03''

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