

A Hybrid Swarm Optimization approach for Feature set reduction in Digital Mammograms

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Abstract: In this paper a CAD (Computer Aided Diagnosis) system is proposed to optimize the feature set using hybrid of Particle Swarm Optimization (PSO) and Genetic Algorithm (GA) technique called Genetical Swarm Optimization (GSO) in Digital Mammogram. Even though PSO is a good optimization technique, it may be trapped in local minima and may prematurely converge. So, the genetic operators are used in PSO to overcome the difficulties. Feature selection plays a major role in diagnosis of mammogram. Gray Level Co-occurrence Matrix (GLCM) texture features are extracted from the mammogram. All the extracted features do not help in detection of abnormality in a mammogram, so it is intended to reduce the feature set to improve classification accuracy. In this work, experiments are conducted on MiniMIAS database and Support Vector Machine (SVM) classifies the mammograms into normal and abnormal mammograms. Performance of GSO is compared with GA and PSO by means of Receiver Operating Characteristic (ROC) curve. Results show that, the GSO convergence is better than both PSO and GA; GSO based SVM (GSO-SVM) classifier exhibits superior performance with an accuracy of 94% which is approximately 1% higher than GA based SVM (GA-SVM) and PSO based SVM (PSO-SVM) classification.

Keywords: Genetic Algorithm, Genetical Swarm Optimization, Particle Swarm Optimization, Support Vector Machine.

1 Introduction

Mammography is the only effective screening method for detection of breast cancer in early stage. Due to wrong interpretation of the radiologist or because of the limitation of human visualization system certain errors like false negative errors may arise. To overcome such limitations of mammography, the researchers developed Computer Aided Diagnosis (CAD) systems to automatically detect and diagnose the abnormalities in digital mammograms.

Genetic Algorithm (GA) is a common optimization technique, introduced by John Holland [1]. It belongs to the family of evolutionary algorithm and

originated from Darwin's theory of natural selection and evolution. Particle Swarm Optimization (PSO) is introduced by Kennedy and Eberhart, is also an optimization technique originated from the idea of swarm intelligence and evolutionary computation. PSO is inspired by the ability of flocks of birds to find rich sources of food, and avoid predators by implementing an "information sharing" mechanism [2].

Both PSO and GA are population based optimization techniques to find a solution to a given objective function but have their own strength and weakness. PSO is a global optimization technique which is known for its speed of convergence, easy to implement and only few parameters to adjust but it

has the drawback that it may quickly cause a particle to stagnate and also prematurely converge on suboptimal solution. In PSO only the best particle share the information with others. In the classical PSO, all parents are directly replaced by their offspring without analyzing whether they can lead a better performance than their parents or not. So, most of the particles move to the worse positions in most of the cases which may lead to early convergence on local minima. In GA, chromosomes of close similarity can converge quickly in fewer generations than PSO.

Some researchers have compared GA and PSO [3]–[6] for various applications. Alessandro Gandelli et al. [7] proposed a hybrid technique that combines PSO and GA called Genetical Swarm Optimization (GSO), to design a planar reflect array antenna, in order to optimize the geometrical features of its elements. It is proved that GSO is reliable and effective technique for wider application in electromagnetic. Alfassio Grimaldi et al. [8] proposed a hybrid evolutionary algorithm called GSO, for the optimization of large-domain electromagnetic problems. The research shows that, GSO performs a quick global search without getting trapped in local minima and performs a quick global search in the synthesis of linear and planar arrays. Davide Caputo et al. [9] presents GSO to optimize the communication energy consumption in a wireless network by selecting the optimal multihop routing schemes. Karnan et al. used hybrid of GA and PSO to detect the nipple position in digital mammogram [10].

In this paper, a hybrid technique that combines the selection strategy of GA and information sharing of PSO to reduce the mammogram feature set is proposed. The roulette wheel selection strategy of GA helps PSO to select the particles for the next generation. This approach helps the PSO to reduce the probability of trapping into local minima. The mutation operator of GA helps to avoid the premature convergence of a particle.

The rest of the paper is organized as follows: Section 2 describes the methodology of the proposed system which includes the Expectation Maximization (EM) segmentation, feature extraction, feature selection by GA, PSO and the proposed hybrid technique and the

Support Vector Machine (SVM) classification.. Section 3 describes the experimental results in detail. Section 4 describes the conclusion and the future enhancements.

2 Methodology

The mammographic images from MiniMIAS database is used in this research. The original mammogram is segmented by Expectation Maximization (EM) algorithm then the 78 Gray Level Cooccurrence Matrix (GLCM) textural features are extracted. The extracted feature set is reduced by GA, PSO and GSO technique. SVM classifier classifies the normal mammogram from an abnormal mammogram. The performance of GA based SVM, PSO based SVM and GSO based SVM is compared by Receiver Operating Characteristic curve (ROC). The block diagram of the proposed system is shown in fig. 1.

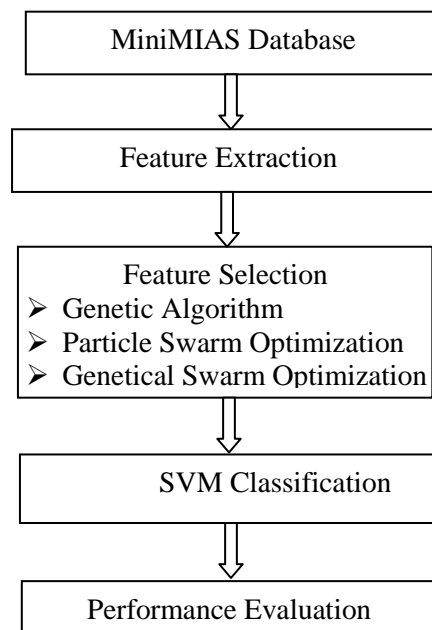


Fig. 1 Block diagram of the proposed system

2.1 EM Segmentation

The Expectation Maximization Algorithm was first introduced by Dempster, Laird, and Rubin [11], [12]. EM is a clustering method to segment background from the breast tissue. EM algorithm is to estimate the parameters of a mixture model and compute maximum likelihood estimates for a given data points

that are generated by a mixture of Gaussian. The algorithm alternates between E step and M step. E step finds the expectation of log likelihood for the given mean vector and variance vector based on the current expected values of mixed weights. M step computes new log likelihood for the μ_k (mean) and σ_k (variance) based on the current expected log-likelihood value.

EM algorithm estimate missing values which is the centers of the clusters. The algorithm optimize the log likelihood of the parameters from a given data set. EM algorithm estimates the probability of elements to be in certain clusters.

Assume a mixture model formed by the image space to be the combination of k Gaussians or k clusters, $\{1, 2, \dots, k\}$, with some prior probabilities w_1, w_2, \dots, w_k of a random point belonging to the associated class. And since each class represents a Gaussian distribution, the probability of each point in image data is given as

$$f(x|\theta) = \sum_{i=1}^k f_i(x|\theta_i) \quad (1)$$

where x is a input image, α_i represents m w_k for k^{th} cluster ($\sum_{i=1}^k \alpha_i = 1$). θ represents collection of parameters ($\mu_1, \mu_2, \dots, \mu_k, \sigma_1, \sigma_2, \dots, \sigma_k$) means and covariance matrix in this case and f_i is a multivariate Gaussian Density function given as,

$$f_i(x|\theta_i) = f_i(x|\mu_i, \sigma_i) \quad (2)$$

$$f_h(x|\mu_i, \sigma_i) = \frac{1}{\sqrt{2\pi} \sigma_i} \exp\left[-\frac{(x-\mu_i)^2}{2\sigma^2}\right] \quad (3)$$

where μ_i stands for mean and σ_i for variance

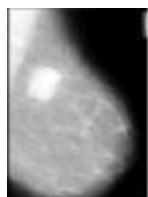


Fig. 2 Original Mammogram



Fig. 3 EM Segmented Mammogram

Fig. 2 is the original mammogram and Fig. 3 is the EM segmented mammogram. From the above figure 3 we can observe that the background muscles are

removed and the region of interest is segmented from the original mammogram.

2.2 Feature Extraction

The purpose of feature extraction is to reduce the original mammogram image into a set of features, by measuring certain properties or features that distinguish one input pattern from another pattern. Gray Level Co-occurrence Matrix (GLCM) is the second order textural measure. GLCM features are extracted in four angles ($0^\circ, 45^\circ, 90^\circ, 135^\circ$) and at four distances ($d=1, 2, 3, 4$). Thirteen features namely energy, correlation, inertia, entropy, inverse difference moment, sum average, sum variance, sum entropy, difference average, difference variance, difference entropy, information measure of correlation 1 and information measure of correlation 2 are extracted at four different angles and four different distance [13]. The thirteen haralick features are listed in table 1. Mean and variance of the thirteen features at four angles are extracted, making a total of 78 features.

Table 1 Haralick's 13 features

1. Energy	8. Sum entropy
2. Correlation	9. Difference average
3. Inertia	10. Difference variance
4. Entropy	11. Difference entropy
5. Inverse difference moment	12. Information measure of correlation 1 and
6. Sum average	13. Information measure of correlation 2
7. Sum variance	

2.3 Feature Selection

Feature selection refers to the problem of dimensionality reduction of data, which initially consists of large number of features. Large number of features led to slow learning of any classifier. Also the classification algorithm gets complex and the cost of classification increases. Using GLCM, 78 features are extracted, but not all the features help in discriminating an abnormal mammogram from a normal mammogram. So, it is necessary to identify and ignore the irrelevant features. The objective of this research is to choose optimal subsets of the original features which contain the information essential for the classification task. For this, three

techniques are proposed namely PSO, GA and a hybrid technique called GSO.

2.3.1 Particle Swarm Optimization Algorithm for feature set optimization

PSO is a population based metaheuristic optimization algorithm, optimizes a problem by having a population of solutions called particles. Each particle has a position represented by a position vector x_i . The feature vector is multiplied with position vector. These particles move around the search space with the velocity vector v_i , searching of the objective function which determines the fitness of the solution. The basic steps of this technique are shown in algorithm 1. In each of the iteration, particles are updated by two best values, called pbest and gbest. Each particle keeps track of its own best position, which is associated with the best fitness it has achieved so far, called pbest. When a particle takes the whole population as its topological neighbor, the global best value is called as gbest.

At each time step t , the individuals best position pbest(t) and the global position gbest(t) are computed for each particle based on this measure. A new velocity of each particle is updated by the equation,

$$v_i[t + 1] = \omega v_i[t] + c1 * r1 * (pbest[t] - x_i[t]) + c2 * r2 * (gbest[t] - x_i[t]) \quad (4)$$

where,

ω is the inertia weight, which controls the impact of the previous velocity,

$v_i[t]$ is the particle velocity,

$x_i[t]$ is the current position of the particle,

$r1, r2$ are random number between $[0, 1]$,

$c1$ and $c2$ are acceleration constants.

On changing the velocity, the particle i searches around its pbest and gbest values. Based on the updated velocities, each particle changes its position according to the following equation.

$$x_i[t + 1] = x_i[t] + v_i[t + 1] \quad (5)$$

During each generation, each particle is accelerated toward the particles previous best position and the global best position. At each iteration a new velocity value is calculated based on its current velocity, the distance from its previous best position, and the

distance from the global best position. The new velocity value is then used to calculate the next position of the particle in the search space. This process is then iterated until a maximum number of iteration has reached or until the target fitness value v_{max} has achieved or until a minimum error has achieved. The PSO algorithm is given below

Algorithm 1 PSO for Feature Optimization

/* s is the dimension of the feature vector

1. Set the constants $c1, c2, \omega$ and s
Randomly initialize particle positions $x_i[]$,
for $i = 1, 2, \dots, s$
Randomly initialize particle velocities $v_i[]$,
for $i = 1, 2, \dots, s$
 2. Set $t = 1$
 3. Evaluate fitness function f_t of each particle
If $f_t \geq f_{pbest}$, then $pbest(t) = x_i[t]$
If $f_t \geq f_{gbest}$, then $gbest(t) = x_i[t]$
 4. Update particle velocity using the equation (4),
and
update particle position vector using equation (5).
 5. Increment i , if $i > s$ then increment t and set $i = 1$.
 6. Repeat steps 3 to 5 until stopping condition is reached.
 7. Sort the particles based on their gbest value and choose the first five feature from that particle.
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The particles are sorted based on their gbest value. Finally the first five feature of the gbest particle is selected as the best feature for the digital mammogram.

2.3.2 Genetic Algorithm for feature set optimization

A common method to select an optimized subset of features is genetic algorithm (GA), which are adaptive heuristic search algorithms based on the principles of Darwinian evolution. The input features are called chromosome. The set of chromosomes are called population. The feature vector is multiplied by the individual population. To each chromosome, calculate the fitness value. The stopping criterion of the GA is the number of generations. Algorithm 2

explains the feature selection mechanism of the mammogram using GA. The GA uses three operators namely: reproduction, crossover and mutation. Reproduction copies the parent chromosomes to generate new population. Crossover mixes genes of two chromosomes to create new generation. Mutation is the random alteration to the position value which simply means changing 0 to 1 and vice versa.

The population is initialized randomly and a new dataset is created. The fitness value is computed for all the chromosomes. Not all the chromosomes are taken to the next generation; the roulette wheel selection strategy is applied to select the parents for the next iteration. A single point cross over followed by mutation is performed to produce new chromosome. Again the fitness is calculated for the chromosomes. This sequence of selection, crossover and mutation processes are repeatedly applied until a best combination of features obtained.

Algorithm 2 GA for Feature Optimization

1. Generate random population with 78 genes in each chromosomes
 2. Evaluate the fitness $f(x)$ of each chromosome x in the population
 3. Create new population by repeating the following steps until optimized feature set is obtained
 - 3.1 Select two chromosomes from the population using the Roulette wheel selection strategy based on their fitness
 - 3.2 Perform crossover of the two parent to form new offspring.
 - 3.3 Do mutation on the newly obtained offspring.
 - 3.4 Replace offspring in the new population
 4. Go to Step 2 and repeat the process until maximum iteration has reached.
 5. Select the first five feature of the highest fit chromosomes.
-

Here the 78 features are represented by a chromosome with 78 genes (bits) corresponding to the number of features. Initial population sizes of 100 chromosomes are randomly generated. Repeatedly apply the genetic operators selection, cross over and mutation in order, until maximum iteration has reached.

2.3.3 Genetical Swarm Optimization for feature set optimization

In the classical PSO, all parents are directly replaced by their offspring without analyzing whether they can lead a better performance than their parents or not. So, most of the particles move to the worse positions in most of the cases. This may lead to early converge on local minima. In this paper, roulette wheel selection in PSO to select the particles for the next generation is introduced. Therefore, the particle's position for the next generation is not only due to the position update but also to the roulette selection strategy. This approach helps the PSO to reduce the probability of trapping into local minima. Select the pbest position and gbest position based on the fitness using the genetic operator, as shown in algorithm 3. Roulette wheel selection strategy selects the best parent, perform the crossover of the selected parent and mutate the offspring. From the new population compute the pbest value and gbest value. Update the position and velocity vector using equation 1 and 2 respectively. Repeat the process until a maximum number of iteration is reached.

Algorithm 3 GSO for Feature Optimization

/*s is the dimension of the feature vector.

1. Set the constants c_1 , c_2 , ω and s . Randomly initialize particle positions $x_i[]$, for $i = 1, 2, \dots, s$. Randomly initialize particle velocities $v_i[]$, for $i = 1, 2, \dots, s$.
 2. Set $t=1$.
 3. Evaluate fitness function f_t of each particle
If $f_t \geq f_{pbest}$ then $pbest(t) = x_i[t]$
If $f_t \geq f_{gbest}$ then $gbest(t) = x_i[t]$
 4. Use roulette wheel strategy to select the parent particle, do crossover and perform mutation on the offspring particle.
 5. Update particle velocity using equation (4), Update particle position vector using equation (5).
 6. Repeat steps 3 to 5 until maximum iteration has reached.
 7. Sort the particle based on their gbest value and choose the first five feature.
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The genetic operators likely selection, cross over and mutation helps the PSO to overcome the difficulty to fall in to local optima and helps the particle to fasten

the convergence of particle. The hybrid technique reduced the 78 featured particle into best five features.

2.4 Support Vector Machine (SVM) Classification

SVM is a classification techniques based on statistical learning theory developed by Vapnik et al., [14], [15]. The major advantage of SVM is that it classifies the samples well in small training samples in high dimensional space. SVM approach is also known as Structural Risk Minimization (SRM) as it produces the largest separation margin. SVM learning algorithm takes two sets of feature vectors, one for normal mammograms and other for abnormal mammograms in a 78-dimensional feature space. It constructs a separating hyperplane in the feature space, which maximizes the margin between the two data sets and find the support vectors that lies on this hyperplane. During testing the sample can be classified as normal class or abnormal class based on the distance of the image to the separating hyperplane.

Hyper planes are defines as $w \cdot x = b$ where w and b are weight and bias parameters respectively. For training data (x_i, y_i) , $i = 1, 2, \dots, n$ are separated by the hyperplanes:

$$w \cdot x + b = +1 \text{ for } y = +1 \quad (6)$$

$$w \cdot x + b = -1 \text{ for } y = -1 \quad (7)$$

where x_i is the feature vector, y_i is the output and can be combined into one set of inequalities:

$$y_i(w \cdot x_i) + b \geq 0 \quad \forall i \quad (8)$$

For a given training set, even though many hyperplanes exist, SVM classifier maximizes the separating margin between the two classes, the SVM classifier is based on the hyperplane that maximizes the separating margin between the two classes. This hyperplane can be found by minimizing the margin given by

$$\text{minimize } \frac{1}{2} \|w\|^2 \quad (10)$$

$$\text{subject to constraints } y_i(w \cdot x_i + b) \quad (11)$$

All the problems are not linearly seperable, in case of non linear-seperable problems the input vector is

mapped to a higher dimension feature space by the non-linear function $\phi(x_i)$ The kernel plays the role of mapping the feature space to higher dimension.

The kernel function is defined as

$$k(x, y) = \sum \phi(x)_i \phi(y)_i \quad (12)$$

The discriminant function of SVM using the Kernel function is

$$f(x) = \sum_{i=1}^n \alpha_i y_i K(x_i, y_i) + b \quad (13)$$

where x_i is the support vector, y_i is the classification output (+1 for benign and -1 for malignant), α_i and b are quadratic programming coefficients. Radial Basis Function Kernel (RBF) is used and it is defined as:

$$K(x_i, x_j) = \exp(-\gamma \|x_i - x_j\|^2), \gamma > 0 \quad (14)$$

where γ , is kernel parameter.

3 Experimental Results and Discussion

3.1 Image Database

In this research, mammograms from the Mammographic Image Analysis Society (MIAS), a Mini Mammographic Database [16] is used. Each mammogram image has a spatial resolution of 1024x1024 pixels. This database is chosen since it contains various types of abnormalities such as calcification, well-defined, circumscribed masses, spiculated masses, ill-defined masses, architectural distortion, asymmetry and normal. Each of these abnormalities has been diagnosed and confirmed by a biopsy.

3.2 Experimental Setup

The experiments implemented in MATLAB. These techniques are experimented on 100 mammogram images with various abnormalities, 50 abnormal images with microcalcification, spiculation, circumscribed and 50 normal mammogram images. The following table (Table 2) shows the parameters used in PSO and GA.

Table 2 Parameters used in PSO and GA

PSO Parameters	GA Parameters
Swarm size: 100	Population size: 100
Max. generations: 200	Max. generations: 200
c1: 2	Selection: Roulette wheel strategy
c2: 2	Crossover: single point cross over
Vmin : 0.4	Crossover rate :0.8
Vmax: 0.9	Mutation rate: 0.2
ω : 0.25	

3.3 Experimental Results

Among the 78 features, the best five feature selected by all three techniques and their sample values are shown in table 3 and table 4 respectively. The five features selected by GSO hybrid techniques is Difference Average, Correlation, Entropy, Sum Variance, Inertia. Correlation is chosen by all the three techniques. The sum variance, inverse difference and information measure of correlation 2 are chosen by at most two techniques.

Table 3 Best five features selected by GA, PSO and GSO

Techniques	Selected GLCM Feature
GA	Difference Average, Correlation, Entropy, Sum Variance, Inertia
PSO	Correlation, Energy, Information measure of correlation2, Inverse Difference, Sum Variance
GSO	Inverse Difference, Sum Variance, Correlation, Difference Variance, Information measure of correlation2

Table 4 Sample of GLCM best features

Image Id.	Category	Inverse Difference	Sum Variance	Correlation	Difference Variance	Information measure of Correlation2
m001	Normal	0.9673	9.7969	0.9719	0.1336	0.9901
m002	Normal	0.9623	9.9808	0.9674	0.1573	0.9877
m003	Normal	0.9667	11.9089	0.9649	0.2045	0.9857
m004	Abnormal	0.9805	13.8280	0.9906	0.0632	0.9781
m005	Abnormal	0.9803	10.7580	0.9898	0.0531	0.9771
m006	Abnormal	0.9798	9.9549	0.9877	0.05952	0.96239

The convergence results of these techniques are shown in fig. 4. PSO from the initial iteration itself scored highest fitness value and converge quickly than the others. In GA and GSO there is no much difference initially stage, but to conclude the PSO and GSO converge in almost same time.

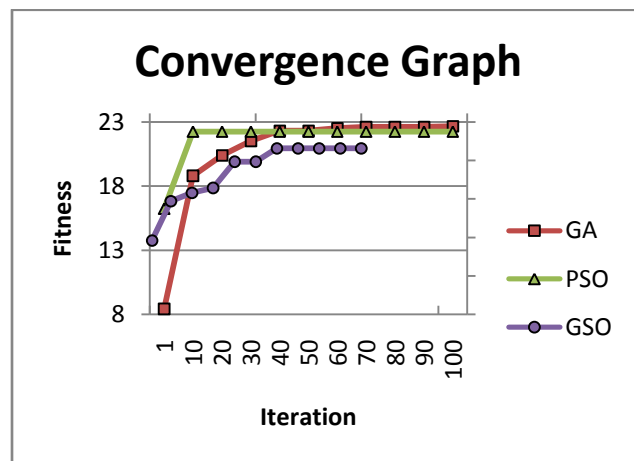


Fig. 4 Convergence graph of the proposed techniques.

The three techniques are classified by means of SVM classifier. The RBF kernel classifies this non linear feature set into normal and abnormal mammogram. The main advantage in using the SVM is that it classifies the data with minimum number of training vectors and achieves good classification accuracy. Fig. 5 shows the output of GSO-SVM classifier. The GSO based SVM classifier achieves the best classification accuracy of 94%, which is approximately 1% higher than PSO and GA based SVM classifier.

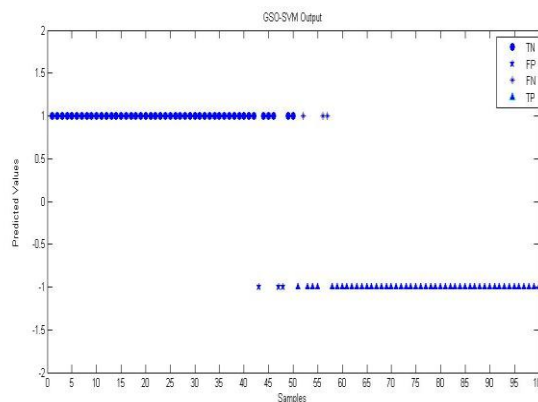


Fig. 5 Result of GSO-SVM

Three mammograms which really contains abnormality is wrongly predicted as normal (FP) and three mammograms which is normal is misclassified as abnormal (FN). The confusion matrices are build from the results of classification and are shown in table 5c. Based on this confusion matrix the sensitivity, specificity and accuracy are calculated. Fig. 6 compares the performance measures of the three techniques in terms of accuracy and an excel graph is plotted.

Table 5a Confusion matrix for GA-SVM

Actual	Predicted	
	Abnormal	Normal
Abnormal	43	5
Normal	7	45

Table 5b Confusion matrix for PSO-SVM

Actual	Predicted	
	Abnormal	Normal
Abnormal	44	4
Normal	6	46

Table 5c Confusion matrix for GSO-SVM

Actual	Predicted	
	Abnormal	Normal
Abnormal	47	3
Normal	3	47

Table 6 Performance of GA-SVM, PSO-SVM and GSO-SVM

Classifier	Accuracy (%)	Sensitivity (%)	Specificity (%)
GA-SVM	88	86	90
PSO-SVM	90	88	92
GSO-SVM	94	94	94

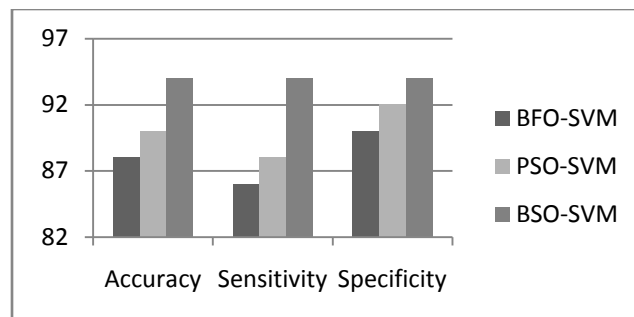


Fig. 6 Performance measure of the proposed techniques

ROC is a graphical tool to plot Sensitivity (TPR-true positive ratio) Vs Specificity (FPR- False negative ratio) for a classifier. The ROC curve is plot against the PSO-SVM, GA-SVM and GSO-SVM classification and the graph is shown in fig. 7.

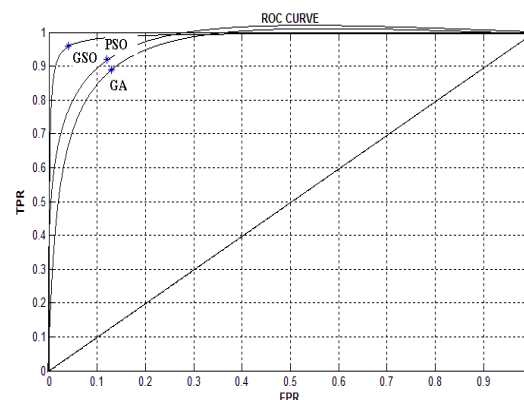


Fig. 7 ROC curves of GA-SVM, PSO-SVM and GSO-SVM

3.4 Discussion

In this experiment, among the 78 features extracted from the mammogram, the best five features of each technique are shown in the table 4. The SVM Classifier is trained with GA based features, PSO based features and GSO based features. 100 images were used in testing, among these 6 mammograms were misclassified and the accuracy of the classifiers are shown in the table 5. It is observed that GSO-SVM is able to classify the mammogram more accurately than GA-SVM and PSO-SVM. Overall classification accuracy of GA-SVM is 88%, PSO-SVM is 90% and GSO-SVM is 94%. As PSO does not need complex operators like crossover, mutation and it requires only primitive and simple mathematical operators also it is faster than GA, but

the convergence of GA is better than PSO. On the whole, GSO shows better accuracy and faster convergence than GA and PSO.

4 Conclusions

In this paper, a CAD to diagnose a digital mammogram as normal or abnormal based on a hybrid technique called GSO was proposed. The hybrid GSO technique performance is compared with GA and PSO. The GSO selects the Inverse Difference, Sum Variance, Correlation, Difference Variance, Information measure of correlation² as the best five features. The experimental results indicate that the GSO converges earlier than PSO and GA algorithms. A classification accuracy of 94% is obtained by GSO based SVM which is approximately 1% increase over PSO based SVM and GA based SVM. Additional features including morphological feature, wavelet based feature, history of the patient and age may be included to target 100% accuracy in results.

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