

## Optimal Design of Fuzzy Clustering-based Fuzzy Neural Networks for Pattern Classification

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### Abstract

We introduce a new category of fuzzy neural networks with multiple-output based on fuzzy clustering algorithm, especially, fuzzy c-means clustering algorithm (FCM-based FNNm) for pattern classification in this paper. The premise part of the rules of the proposed networks is realized with the aid of the scatter partition of input space generated by FCM clustering algorithm. The partitioned local spaces describe the fuzzy rules and the number of the partitioned local spaces is equal to the number of clusters. Due to these characteristics, we may alleviate the problem of the curse of dimensionality. The consequence part of the rules is represented by polynomial functions with multiple-output for pattern classification. And the coefficients of the polynomial functions are learned by back propagation algorithm. To optimize the parameters of the proposed FCM-based FNNm we consider real-coded genetic algorithms. The proposed networks are evaluated with the use of numerical experimentation.

**Keywords:** Fuzzy Neural Networks, FCM clustering algorithm, Scatter partition of input space, Optimization, Genetic Algorithms

### 1. Introduction

Fuzzy neural networks (FNNs) [1, 2] have emerged as one of the active areas of research in fuzzy inference systems and neural networks. FNNs are predominantly concerned with the integration of these two fields. Fuzzy inference systems have been studied to model uncertain and/or ambiguous characteristics inherent to experimental data. Fuzzy inference systems are good at explaining decisions but they cannot directly obtain the fuzzy rules. Neural networks are good at classifying patterns but they are not good at explaining how they acquire decisions. These limitations have been the major points and these techniques are combined to overcome the shortcomings. Since its inception, the research of FNNs has been a focal point of various endeavors and has demonstrated many fruitful results in application [3, 4, 5]. Typically, FNNs are represented by fuzzy “if-then” rules while the back propagation (BP) is used to optimize the parameters.

The generation of the fuzzy rules and the adjustment of its membership functions were done by trial and error and/or operator’s experience. The designers find it difficult to develop adequate fuzzy rules and membership functions to reflect the essence of the data. Moreover, some information gets lost or ignored on purpose when human operators articulate their

experience in the form of linguistic rules. As a consequence, there is a need for an optimization environment to construct and/or adjust a collection of linguistic rules.

In this paper, we present the structure of fuzzy neural networks with multiple-output by means of fuzzy c-means clustering algorithm [6] (FCM-based FNNm). The premise part of the rules is realized with the aid of the scatter partition of input space generated by FCM clustering algorithm. The partitioned local spaces describe the fuzzy rules. The consequence part of the rules is represented by polynomial functions with multiple-output for pattern classification. And the coefficients of the polynomial functions are learned by BP algorithm. We also optimize the parameters of the networks using real-coded genetic algorithms [7]. The proposed networks are evaluated through the numeric experimentation.

The paper is organized as follows. Section 2 is concerned with the design of FCM-based FNNm. Section 3 deals with the optimization of FCM-based FNNm. Section 4 presents results of numeric experimentations. Finally Section 5 concludes the paper.  $\square u_{ip}=1, 1 \leq p \leq N$

\*

**2. Design of Fuzzy c-means Clustering-based Fuzzy Neural Networks**

In this section, the form of fuzzy clustering if-then rules along with their development mechanism is discussed. More specifically, we elaborate on the three types of fuzzy inference and present the learning algorithm.

**2.1. Fuzzy c-means Clustering Algorithm**

The premise part of the FCM-based FNNm is developed by means of the Fuzzy C-Means clustering algorithm [6] as the fuzzy clustering. This algorithm is aimed at the formation of  $c$  fuzzy sets (relations) in  $\mathbf{R}^n$ . Consider the set  $\mathbf{X}$ , which consists of  $N$  data points treated as vectors located in some  $n$ -dimensional Euclidean space. In clustering we assign patterns  $\mathbf{x}_p \in \mathbf{X}$  into  $c$  clusters, which are represented by its prototypes  $\mathbf{v}_i \in \mathbf{R}^n$ . The assignment to individual clusters is expressed in terms of the partition matrix  $\mathbf{U} = [u_{ip}]$  where

$$u_{ip} = \frac{1}{\sum_{c=1}^c u_{ip}} \quad (1)$$

The objective function  $Q$  guiding the clustering is expressed as a sum of the distances of individual data from the prototypes  $\mathbf{v}_1, \mathbf{v}_2, \dots,$  and  $\mathbf{v}_c$ ,

$$Q = \sum_{p=1}^N \sum_{i=1}^c u_{ip}^m \|\mathbf{x}_p - \mathbf{v}_i\|^2 \quad (3)$$

$\square$

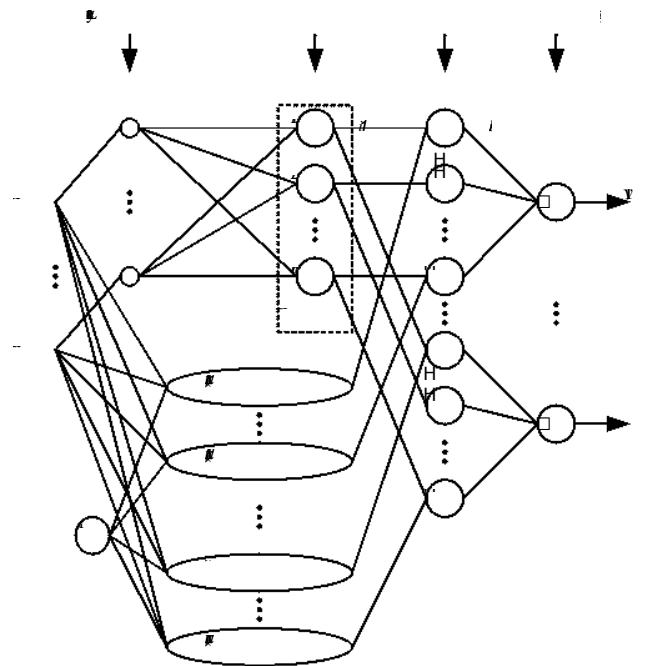
Here  $\|\cdot\|$  denotes the Euclidean distance; ' $m$ ' stands for a fuzzification coefficient,  $m > 1.0$ .

52 The resulting partition matrix is denoted by  $\mathbf{U} = [u_{ip}]$ .

The minimization of  $Q$  is realized through successive iterations by adjusting both the

prototypes and entries of the partition matrix, that is  $\min Q(\mathbf{U}, \mathbf{v}_1, \mathbf{v}_2, \dots, \mathbf{v}_c)$ . The corresponding formulas used in an iterative fashion read as follows.





**Figure 2. The Structure of FCM-based FNNm**

FCM-based FNNm is implied by the fuzzy scatter partition of input space. In this sense, each rule can be viewed as a certain rule of the following format

As far as inference schemes are concerned, we distinguish these types Type 1 (Simplified Inference):

$$f^w = \sum_{k=1}^s \frac{w_{kx}^z}{\sum_{k=1}^s w_{kx}^z}, \quad z = d + 1, \dots, d(d+1)/2.$$

Type 2 (Linear Inference):

$d$

$s$

Type 3 (Modified Quadratic Inference):

(9)

To be more specific, is the  $j$ -th fuzzy rule, while denotes  $j$ -th membership grades using FCM clustering algorithm. , are consequent parameters of the rule and  $s$  is the number of output.

(7)

(8)

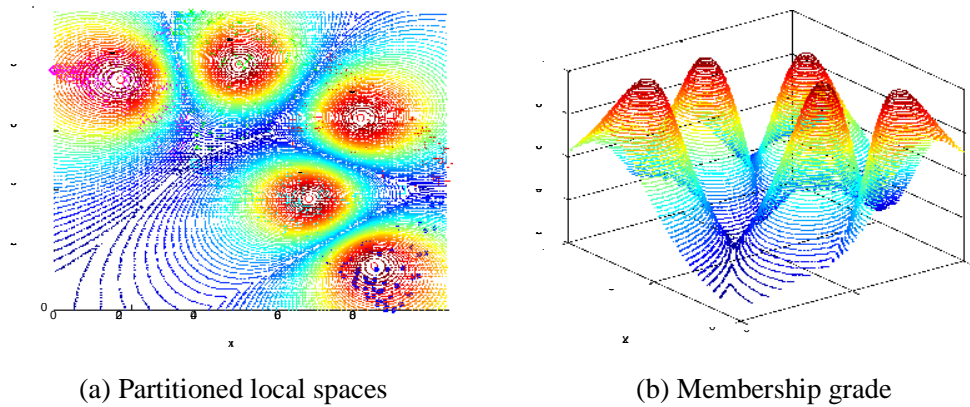
The functionality of each layer is described as follows.

[Layer 1] The nodes in this layer transfer the inputs.

[Layer 2] The nodes here are used to calculate the membership degrees using FCM clustering algorithm. From this algorithm, the firing strengths are as follows.

$$(10)$$

Figure 3 visualizes the example of the fuzzy partitioned spaces of input space with five clusters by means of FCM clustering algorithm.



**Figure 3. Scatter Partition of Input Space**

□

[Layer 3] The nodes in this layer realize a certain inference process.

$$h_j = \frac{1}{n} \sum_{i=1}^n \mu_{ij} \quad (11)$$

$$h_j = \frac{1}{n} \sum_{i=1}^n \mu_{ij} \quad (12)$$

Where, all the entries of  $h_j$  sum up to 1 as indicated by (1).

[Layer 4] The nodes in this layer compute the outputs.

$\hat{\sim}$

### 2.3. Learning Algorithm

The parametric learning of the network is realized by adjusting connections of the neurons and as such it could be realized by running a standard Back-Propagation (BP) algorithm. The performance index is based on the Euclidean distance,

$q$

(13)



$$\hat{y}_{ps}$$

where,  $E_p$  is an error reported for the  $p$ -th data,  $y_{ps}$  is the  $p, s$ -th target output data and  $\hat{y}_{ps}$  stands for the  $p, s$ -th actual output of the network.

As far as learning is concerned, the connections are adjusted in a standard fashion,

$$\Delta w_{ps} = -\eta \frac{\partial E_p}{\partial w_{ps}} \tag{14}$$

where this update formula follows the gradient descent method.

$$\Delta w_{ps} = -\eta \frac{\partial E_p}{\partial w_{ps}} \tag{15}$$

with  $\eta$  being a positive learning rate.

If the type of consequence part is simplified inference then  $\Delta w_{ps}$ . From the chain rules we have the following expression.

$$\Delta w_{ps} = -\eta \frac{\partial E_p}{\partial w_{ps}} \tag{16}$$

$$\Delta w_{ps} = -\eta \frac{\partial E_p}{\partial w_{ps}} \tag{17}$$

$$\Delta w_{ps} = -\eta \frac{\partial E_p}{\partial w_{ps}} \tag{18}$$

$$\Delta w_{ps} = -\eta \frac{\partial E_p}{\partial w_{ps}} \tag{19}$$

Quite commonly to accelerate convergence, a momentum coefficient is being added to the learning expression. Then the complete update formula reads as follows

(20)

(21)

(22)

### **3. Optimization of Networks**

The need to solve optimization problems arises in many fields and is especially dominant in the engineering environment. There are several analytic and numerical optimization techniques, but there are still large classes of functions that are fully addressed by these techniques. Especially, the standard gradient-based optimization techniques that are being used mostly at the present time are augmented by a differential method of solving search problems for optimization processes. Therefore, the optimization of fuzzy models may not be fully supported by the standard gradient-based optimization techniques, because of the nonlinearity of models represented by rules based on linguistic levels. This forces us to explore other optimization techniques such as genetic algorithms (GAs) [7].

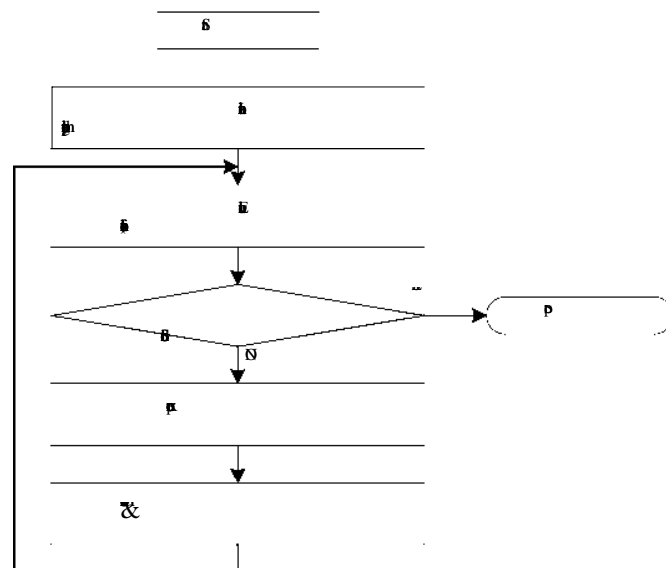
It has been demonstrated that genetic algorithms are useful in a global population-based optimization. GAs are shown to support robust search in complex search spaces. Given their stochastic character, these methods are less likely to get trapped in local minima in comparison to the performance offered by gradient-descent techniques.

GAs start with a randomly generated population of  $l$  chromosomes positioned in solution (parameter) space. The population is evolved repeatedly toward achieving a better overall fitness value. The search in the solution space is completed with the aid of several genetic operators. There are three main generic genetic operators such as reproduction, crossover, and mutation supporting movements in the search space. Let us briefly elaborate on the essence of these operators.

Reproduction is a process in which the mating pool for the next generation becomes selected. Individual strings are copied into the mating pool according to their fitness function values.

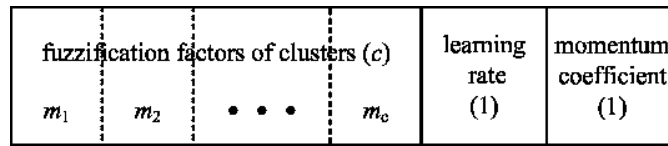
Crossover usually proceeds in two steps. Firstly, members from the mating pool are mated at random. Secondly, each pair of strings undergoes crossover as follows; a position  $l$  along the string is selected uniformly at random from the interval  $[1, l-1]$ , where  $l$  is the length of the string. Swapping all characters between the positions  $k$  and  $l$  creates two new strings.

Mutation is a random alteration of the value of a string position. In real coding, mutation is defined as an alternation at a random value in special boundary. Mutation occurs with a small probability. Those operators, combined with the proper definition of the fitness function, constitute the main body of the genetic computing. A general flowchart of the genetic algorithm is shown in Figure 4.



**Figure 4. A General GA Flowchart**

In order to optimize the parameters of the FCM-based FNNm, we determined the fuzzification coefficients associated with the corresponding clusters, the learning rate, and the momentum coefficient as the parameters. These parameters are genetically optimized across generation. Figure 5 illustrates an arrangement of the content of the chromosomes. Each chromosome is coded using real numbers (instead of binary numbers). This type of coding is helpful from the point of view of effectiveness of the overall search process.



**Figure 5. Data Structure of Chromosomes**

#### 4. Experimental Studies

We discuss three numerical examples in order to evaluate the proposed approach.

For the evaluation of the performance of the network, the random sub-sampling method was applied. In this method,  $K$  data splits of the overall data set were performed. Each split was randomly selected with a fixed number of examples. The random sub-sampling was performed with 5 data splits of the data set ( $K=5$ ). Each split was randomly selected from the training examples and the test examples with the ratio of 7:3.

The classification ratio (CR) is defined as the average of the separate estimates  $E_p$ .

$$CR = \frac{1}{K} \sum_{k=1}^K CR_k$$

Another performance index (PI) is based on the Mean Squared Error (MSE)

$$PI = \frac{1}{K} \sum_{k=1}^K MSE_k$$

We experimented with the proposed network using the parameters outlined in Table 1 and

Table 2 with the weight factor [8].

~~Table 1. Initial Parameters for GAs~~

Parameter	Value
Generation	100
Population size	50
Crossover rate	0.65
Mutation rate	0.1

~~Table 2. Initial Parameters for FCM-based FNNm~~

Parameter	Value
Fuzzification coefficients	$1.0 < m_i \leq 2.5$
Learning rate	$0.0 \leq \eta \leq 0.01$
Moment coefficient	$0.0 \leq \alpha \leq 0.001$

#### 4.1. Iris Dataset

In this section, we use the Iris dataset [9]. The Iris dataset is a collection of 150 Iris flowers of 3 kinds, with four attributes, leaf and petal width and length in *cm*. Three classes are the setosa, versicolor, and virginica.

Table 3 and Figure 6 show the performance of CR and PI for FCM-based FNNm before optimization. Table 4 and Figure 7 present the performance of CR and PI for FCM-based FNNm using genetic optimization. From these tables and figures we know that the optimized FCM-based FNNm is better than before optimization.

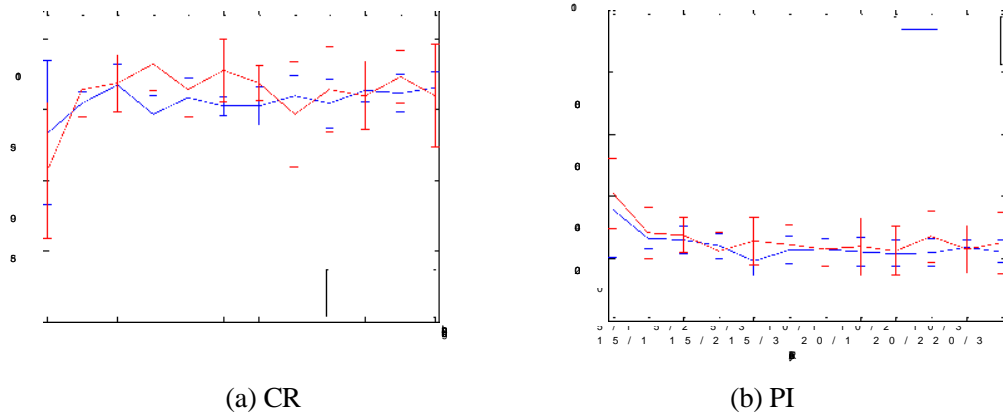
**Table 3. Performance of the FCM-based FNNm for Iris Dataset**

	No. of Clusters	Inference (Type)	CR		PI	
			Training	Testing	Training	Testing
5	1	93.33±5.13	90.67±4.82	0.035±0.02	0.040±0.01	
	2	95.43±0.80	96.44±1.99	0.026±0.00	0.028±0.01	
	3	96.76±1.44	96.89±1.99	0.025±0.00	0.027±0.01	
10	1	94.67±1.28	98.22±1.86	0.024±0.00	0.022±0.01	
	2	95.81±1.44	96.44±1.99	0.019±0.00	0.025±0.01	
	3	95.24±0.67	97.78±2.22	0.022±0.00	0.024±0.01	
15	1	95.24±1.35	96.89±1.22	0.023±0.00	0.023±0.01	
	2	96.00±1.41	94.67±3.72	0.022±0.00	0.023±0.01	
	3	95.43±1.70	96.44±2.98	0.021±0.00	0.022±0.01	
20	1	96.38±0.80	96.00±2.43	0.021±0.00	0.026±0.01	
	2	96.19±1.35	97.33±1.86	0.023±0.00	0.023±0.01	
	3	96.57±1.09	96.00±3.65	0.022±0.00	0.024±0.01	

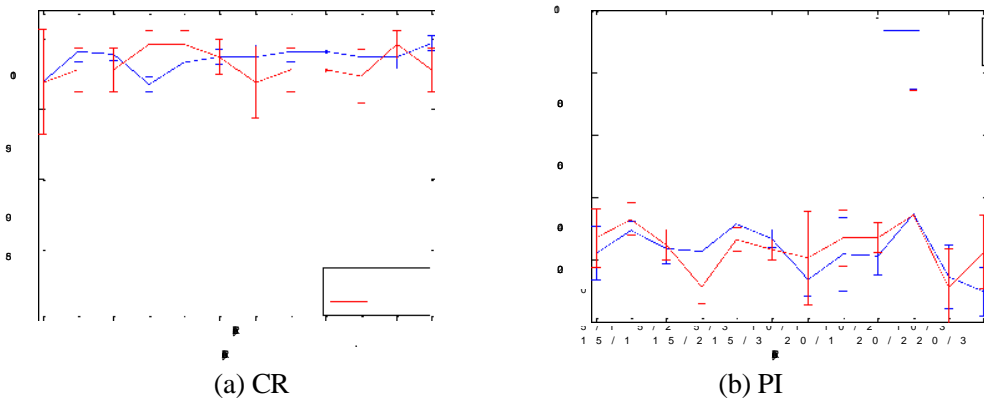
**Table 4. Performance of the Optimized FCM-based FNNm for Iris Dataset**

	No. of Clusters	Inference (Type)	CR		PI	
			Training	Testing	Training	Testing
5	1	96.95±1.56	96.89±3.72	0.022±0.01	0.027±0.01	
	2	99.05±0.67	97.78±1.57	0.029±0.00	0.033±0.01	
	3	98.86±0.43	97.78±1.57	0.023±0.00	0.024±0.00	
10	1	96.76±0.52	99.56±0.99	0.022±0.00	0.011±0.01	
	2	98.29±0.43	99.56±0.99	0.031±0.00	0.026±0.00	
	3	98.67±0.52	98.67±1.22	0.026±0.00	0.023±0.00	
15	1	98.67±0.85	96.89±2.53	0.013±0.01	0.020±0.02	
	2	99.05±0.67	97.78±1.57	0.021±0.01	0.026±0.01	
	3	99.05±0.00	97.78±0.00	0.021±0.01	0.027±0.00	
20	1	98.67±0.85	97.33±1.86	0.034±0.04	0.034±0.04	
	2	98.67±0.85	99.56±0.99	0.014±0.01	0.011±0.01	
	3	99.62±0.52	97.78±1.57	0.009±0.01	0.022±0.01	

From Table 4 and Figure 7 we select the network that has twenty fuzzy rules and linear inference (Type 2) engine. This network exhibits CR=98.67±0.85, PI=0.014±0.01 for training datasets and CR=99.56±0.99, PI=0.011±0.01 for testing datasets.



**Figure 6. Performance of the FCM-based FNNm for Iris Dataset**



**Figure 7. Performance of the Optimized FCM-based FNNm for Iris Dataset**

Table 5 shows the confusion matrix for the selected network. The result indicates some misclassification for Versicolor and Virginica for training datasets and Virginica for testing datasets.

**Table 5. Confusion Matrix for the Selected Network**

(a)	Training datasets		
	Setosa	Versicolor	Virginica
Setosa	100.00±0.00	0.00±0.00	0.00±0.00
Versicolor	0.00±0.00	97.14±2.86	2.86±2.86
Virginica	0.00±0.00	1.14±1.56	98.86±1.56

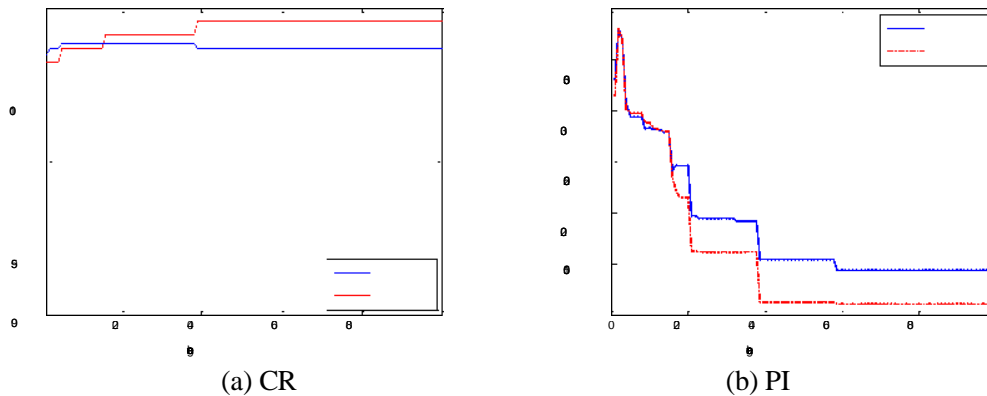
(b)	Testing datasets		
	Setosa	Versicolor	Virginica
Setosa	100.00±0.00	0.00±0.00	0.00±0.00
Versicolor	0.00±0.00	100.00±0.00	0.00±0.00
Virginica	0.00±0.00	1.33±2.98	98.67±2.98

Table 6 shows the optimized parameters of fuzzification coefficients associated with the corresponding clusters, the learning rate, and the momentum coefficient for the selected network using genetic algorithms.

**Table 6. Optimized Parameters for the Selected Network**

$m_1$	$m_2$	$m_3$	$m_4$	$m_5$	$m_6$	$m_7$	$m_8$
1.565712	1.488977	1.382376	1.142042	2.238865	1.916492	2.176308	1.301495
$m_9$	$m_{10}$	$m_{11}$	$m_{12}$	$m_{13}$	$m_{14}$	$m_{15}$	$m_{16}$
1.928584	2.383059	1.208994	2.343709	1.662824	1.412027	1.253087	2.29295
$m_{17}$	$m_{18}$	$m_{19}$	$m_{20}$	$\eta$	$\alpha$		
2.430314	1.711756	2.407883	1.887396	0.009332	0.000449		

Figure 8 presents the optimization procedure for the CR and PI when using twenty rules with Type 2 (Linear Inference) obtained in successive generations of the genetic optimization. These figures depict the average values using the random sub-sampling.



**Figure 8. Optimization Process for the Selected Network**

The performance of the proposed model is compared with the performance of some other models reported in the literature; refer to Table 7. The comparison shows that the proposed model outperforms several previous developed models.

**Table 7. Comparison of Performance with Previous Models**

Model	Classification Ratio (%)
NEFCLASS [10]	96.0
C4.5 [11]	94.0
FID3.1 [12]	96.0
HNFB [13]	98.67
HNFBQ [14]	98.67
HNFB-1 [15]	98.67
Our model	99.56

#### 4.2. WDBC Dataset

In this section, we use the WDBC dataset [16]. Features of WDBC are computed from a digitized image of a fine needle aspirate (FNA) of a breast mass. They describe characteristics of the cell nuclei present in the image. The WDBC dataset is a collection of 569 instances of 2 kinds, with 30 attributes. Two classes are benign and malignant.

Here we use the simplified inference (Type I) and linear inference (Type II) to deal with the high dimensionality. Table 8 and Table 9 summarize the performances of CR and PI for FCM-based FNNm before and after optimization, respectively. Figure 9 and Figure 10 depict the performances of CR and PI in the same case.

**Table 8. Performance of the FCM-based FNNm for WDBC Dataset**

	No. of Inference Clusters	Type	CR				PI	
			Training		Testing		Training	Testing
			CR	PI	CR	PI		
5	1	91.91±1.26	93.10±1.72	0.057±0.00	0.056±0.01			
	2	97.59±0.14	96.96±1.40	0.038±0.00	0.041±0.00			
10	1	92.46±0.36	91.93±0.96	0.057±0.00	0.057±0.01			
	2	96.78±0.45	95.79±1.82	0.039±0.00	0.041±0.01			
15	1	92.11±0.92	92.40±1.43	0.056±0.00	0.057±0.01			
	2	96.23±0.50	94.97±0.98	0.038±0.00	0.043±0.00			
20	1	92.41±0.84	92.87±2.20	0.054±0.00	0.057±0.01			
	2	95.78±0.37	94.74±1.24	0.039±0.00	0.044±0.00			

**Table 9. Performance of the Optimized FCM-based FNNm for WDBC Dataset**

	No. of Inference Clusters	Type	CR				PI	
			Training		Testing		Training	Testing
			CR	PI	CR	PI		
5	1	95.53±0.74	95.32±1.43	0.039±0.01	0.044±0.01			
	2	98.59±0.38	98.13±0.96	0.041±0.01	0.042±0.01			
10	1	96.03±0.45	94.85±0.76	0.035±0.00	0.042±0.00			
	2	98.49±0.18	98.01±1.06	0.045±0.01	0.048±0.01			
15	1	96.18±0.54	95.32±1.24	0.038±0.01	0.044±0.00			
	2	99.05±0.48	97.08±1.31	0.042±0.00	0.053±0.01			
20	1	96.43±0.57	95.32±1.17	0.032±0.00	0.041±0.01			
	2	98.69±0.37	97.54±1.05	0.039±0.01	0.050±0.02			

From Table 9 and Figure 10 we select the network that has five fuzzy rules (clusters) and linear inference (Type 2) engine. This network exhibits CR=98.59±0.38, PI=0.041±0.01 for training datasets and CR=98.13±0.96, PI=0.042±0.01 for testing dataset.

Table 10 shows the confusion matrix for the selected network. The result indicates some misclassification for both Benign and Malignant for training datasets and testing datasets.

Table 11 shows the optimized parameters of fuzzification coefficients associated with the corresponding clusters, the learning rate, and the momentum coefficient for the selected network using genetic algorithms.

Figure 11 presents the optimization procedure for the CR and PI when using five rules with linear inference (Type 2) obtained by genetic optimization.



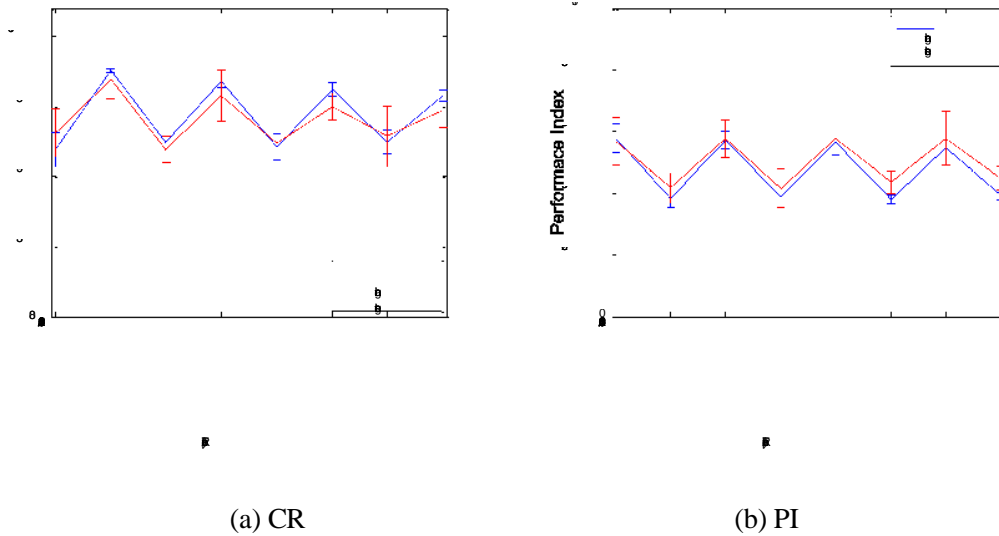


Figure 9. Performance of the FCM-based FNNm for WDBC Dataset

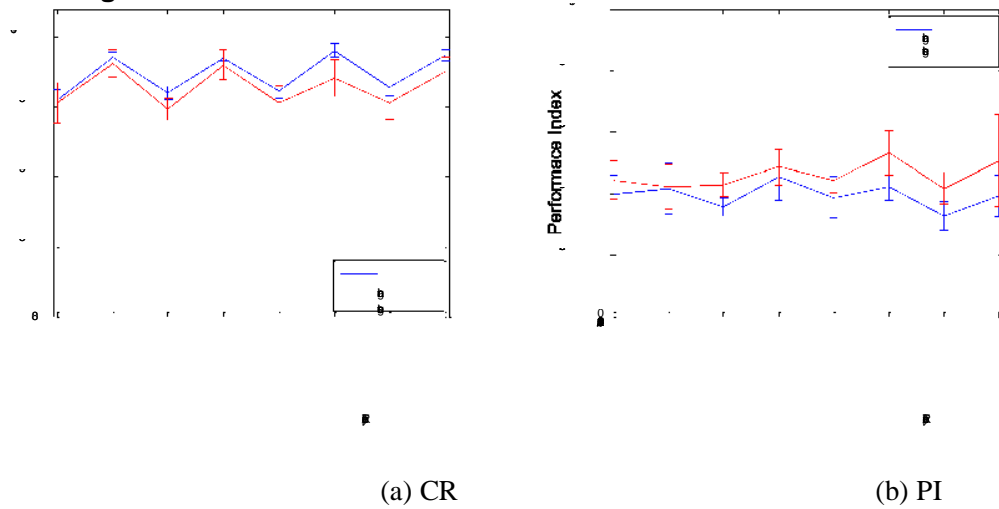


Figure 10. Performance of the Optimized FCM-based FNNm for WDBC Dataset

Table 10. Confusion Matrix for the Selected Network

(a) Training datasets

	Benign	Malignant
Benign	99.52±0.52	0.48±0.52
Malignant	2.97±1.02	97.03±1.02

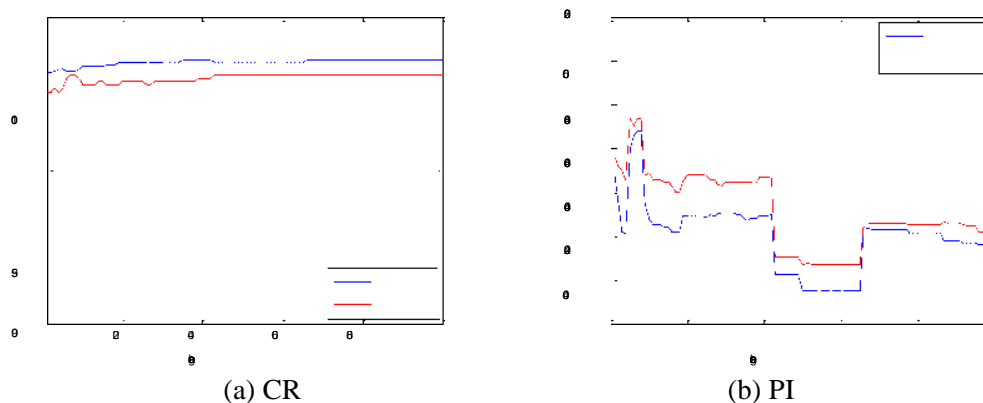
(b) Testing datasets

	Benign	Malignant
Benign	98.88±1.22	1.12±1.22
Malignant	3.13±1.56	96.88±1.56

Table 11. Optimized Parameters for the Selected Network

$m_1$	$m_2$	$m_3$	$m_4$	$m_5$	$\eta$	$\alpha$
1.476586	1.462957	1.88349	2.335035	1.902575	0.007307	
0.000259						

Table 12 shows the performance of the proposed model to compare with the performance of some other models reported in the literature.



**Figure 11. Optimization Process for the Selected Network**

**Table 12. Comparison of Performance with Previous Models**

Model	Classification Ratio (%)
SVM [17]	96.68
Bayes Net [18]	95.81
RVM [19]	97.2
MPANN [20]	98.1
MLP [21]	85.92
DigaNN [22]	97.9
RBF2 [23]	97.13
Our model	98.13

### 4.3. Wine Dataset

In this section, we use the Wine dataset [24]. These data are the results of a chemical analysis of wines grown in the same region in Italy but derived from three different cultivars. The analysis determined the quantities of 13 constituents found in each of the three types of wines.

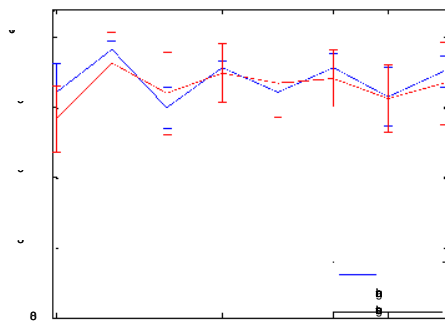
**Table 13. Performance of the FCM-based FNNm for Wine Dataset**

	No. of Clusters	Inference (Type)	CR		PI	
			Training	Testing	Training	Testing
5	1	96.10±2.02	94.18±2.37	0.040±0.01	0.042±0.00	
	2	99.19±0.57	98.18±2.23	0.019±0.00	0.031±0.00	
10	1	94.96±1.45	96.00±2.99	0.038±0.00	0.035±0.01	
	2	97.89±0.45	97.45±2.07	0.026±0.00	0.031±0.01	
15	1	96.10±1.85	96.73±2.37	0.037±0.00	0.036±0.00	
	2	95.77±2.10	95.64±2.44	0.038±0.00	0.039±0.01	
20	1	97.89±0.93	97.09±2.07	0.028±0.00	0.034±0.00	
	2	97.56±1.15	96.73±2.99	0.029±0.00	0.032±0.01	

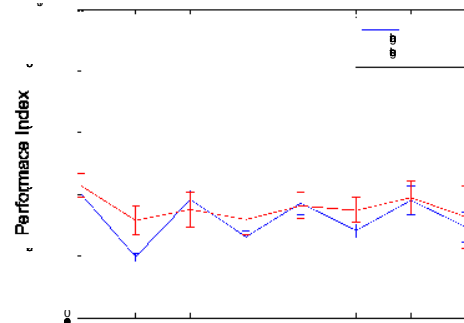
Here we use the simplified inference (Type I) and linear inference (Type II). Table 13 and Figure 12 show the performance of CR and PI for FCM-based FNNm before optimization. Table 14 and Figure 13 present the performance of CR and PI for FCM-based FNNm using genetic optimization.

**Table 14. Performance of the Optimized FCM-based FNNm for Wine Dataset**

	No. of Clusters	Inference (Type)	CR		PI	
			Training	Testing	Training	Testing
5	1	98.86±0.45	98.55±0.81	0.020±0.00	0.024±0.00	
	2	100.00±0.00	100.00±0.00	0.012±0.00	0.018±0.00	
10	1	99.51±0.45	98.91±1.00	0.010±0.00	0.016±0.01	
	2	99.84±0.36	99.64±0.81	0.012±0.00	0.020±0.01	
15	1	99.67±0.45	99.27±1.00	0.007±0.01	0.016±0.00	
	2	99.84±0.36	99.64±0.81	0.008±0.00	0.013±0.01	
20	1	99.84±0.36	99.27±1.00	0.011±0.02	0.019±0.01	
	2	100.00±0.00	99.27±1.00	0.006±0.01	0.017±0.01	

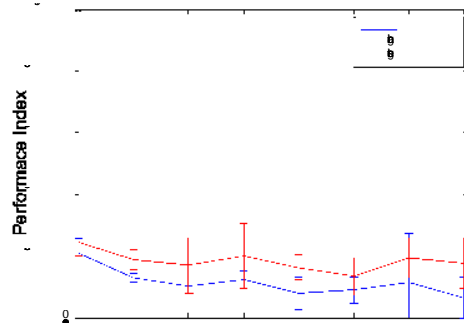
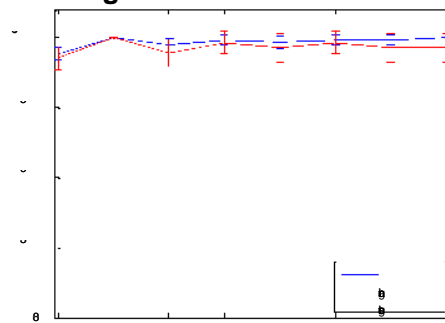


(a) CR



(b) PI

**Figure 12. Performance of the FCM-based FNNm for Wine Dataset**



(a) CR

(b) PI

**Figure 13. Performance of the Optimized FCM-based FNNm for Wine Dataset**

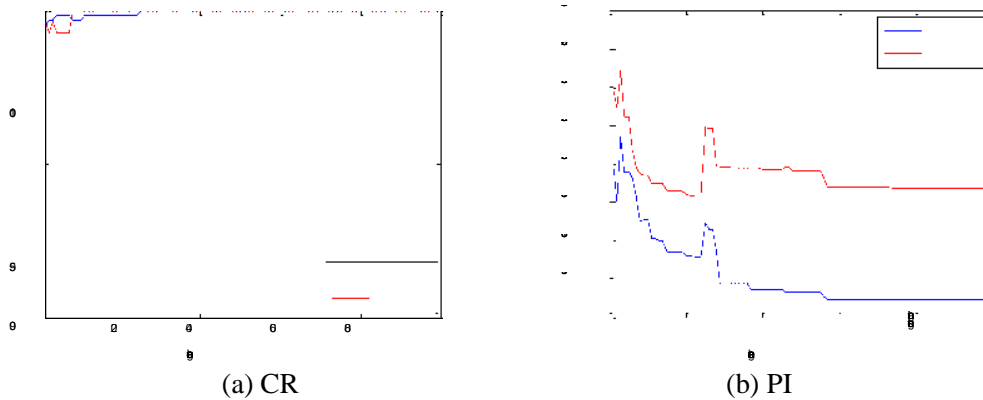
From Table 14 and Figure 13 we select the network that has five fuzzy rules (clusters) and linear inference (Type 2) engine. This network exhibits  $CR=100.00\pm 0.00$ ,  $PI=0.012\pm 0.00$  for training datasets and  $CR=100.00\pm 0.00$ ,  $PI=0.018\pm 0.00$  for testing dataset.

Table 15 shows the optimized parameters of fuzzification coefficients associated with the corresponding clusters, the learning rate, and the momentum coefficient for the selected network using genetic algorithms.

**Table 15. Optimized Parameters for the Selected Network**

$m_1$	$m_2$	$m_3$	$m_4$	$m_5$	$\eta$	$\alpha$
1.145054	2.332643	1.25783	1.195322	1.726746	0.009304	0.000998

Figure 14 presents the optimization procedure for the CR and PI when using five rules with linear inference (Type 2) obtained by genetic optimization.



**Figure 14. Optimization Process for the Selected Network**

0.012

## 5. Conclusions

In this paper, the design of the fuzzy c-means clustering-based fuzzy neural networks has been introduced and its optimization using real-coded genetic algorithms has been discussed for pattern classification.

The input spaces of the proposed networks were divided as the scatter form using FCM clustering algorithm to generate the fuzzy rules. The partitioned spaces describe the fuzzy  
 66 rules and the number of the fuzzy rules is equal to the number of clusters. From this

method, we could alleviate the problem of the curse of dimensionality and have designed fuzzy neural networks compact and easy. Effectively partitioning input space can decrease the number of fuzzy rules and thus increase the learning speed. And genetic algorithms were also used for parametric optimization of the proposed networks.

From the results in the previous section, we were able to design good networks for pattern classification. Through the use of performance we were able to achieve a balance between the approximation and generalization abilities of the resulting network. Finally it could be possible to apply to many fields.

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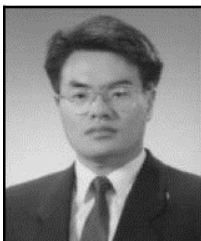
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