

# Classifier Ensemble Framework: a Diversity Based Approach

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**Abstract**—Pattern recognition systems are widely used in a host of different fields. Due to some reasons such as lack of knowledge about a method based on which the best classifier is detected for any arbitrary problem, and thanks to significant improvement in accuracy, researchers turn to ensemble methods in almost every task of pattern recognition. Classification as a major task in pattern recognition, have been subject to this transition. The classifier ensemble which uses a number of base classifiers is considered as meta-classifier to learn any classification problem in pattern recognition. Although some researchers think they are better than single classifiers, they will not be better if some conditions are not met. The most important condition among them is diversity of base classifiers. Generally in design of multiple classifier systems, the more diverse the results of the classifiers, the more appropriate the aggregated result. It has been shown that the necessary diversity for the ensemble can be achieved by manipulation of dataset features, manipulation of data points in dataset, different sub-samplings of dataset, and usage of different classification algorithms. We also propose a new method of creating this diversity. We use Linear Discriminant Analysis to manipulate the data points in dataset. Although the classifier ensemble produced by proposed method may not always outperform all of its base classifiers, it always possesses the diversity needed for creation of an ensemble, and consequently it always outperforms all of its base classifiers on average.

**Index Terms**— Classifier Ensemble, Diversity, Linear Discriminant Analysis.

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## I. INTRODUCTION

Different pattern recognition tasks are employed in different problems. Pattern recognition is considered as a general tool for solving any problem in any field [2], [18-19], [21], [27-35]. Clearly, it is always needed to find a better pattern recognition model. Classification as a major task in pattern recognition is not an exception to this subject. Classification is a task that tries to predict category of some objects. In ensemble method for classification, many classifiers are combined to make a final prediction. Ensemble methods show better performances than a single classifier in general. The final decision is usually made by voting after combining the predictions from set of classifiers.

Most of the classification researches resulted in algorithms that have provided a good performance for specific problem, but they have not enough robustness for other problems. Because of the difficulty that these algorithms are faced to, the recent researches have been directed to the combinational methods that have more power, robustness, resistance, accuracy and generality [15]. Although the accuracy of the classifier ensemble is not always better than the most accurate classifier in ensemble pool, its accuracy is never less than their average accuracy [9]. Classifier ensemble can be considered as a general solution method for pattern recognition problems [14-15]. Inputs of classifier ensemble are predicted class tags of base classifiers and its output is consensus predicted class tags. It is an accepted subject in pattern recognition that finding the best classifier model for solving a given problem is impossible [22-23] and it has some serious drawbacks. The main drawback is

that the best individual classifier for the given classification problem is very difficult to identify, unless deep prior knowledge is available for such a task [3]. It is worthy to noting that the motivations in favor of classifier ensemble strongly resemble those of a “hybrid” intelligent system. The obvious reason for this is that classifier ensemble can be regarded as a special-purpose hybrid intelligent system.

It is believed that “combining the diverse classifiers any of which has better results than random ones, creates a good ensemble”. Diversity is always considered as a crucial concept in classifier ensemble. It is considered as the most effective factor in succeeding an ensemble. The diversity in an ensemble refers to the amount of dissimilarity in the outputs of its components (base classifiers) in deciding for a given sample. Assume an example dataset with two classes. Indeed the diversity concept for an ensemble of two classifiers refers to the probability that they may produce two dissimilar results for an arbitrary input sample. The diversity concept for an ensemble of three classifiers refers to the probability that one of them produces dissimilar result from the two others for an arbitrary input sample. It is worthy to mention that the diversity can converge to 0.5 and 0.66 in the ensembles of two and three classifiers respectively. Although reaching the more diverse ensemble of classifiers is generally handful, it is harmful in boundary limit. It is very important dilemma in classifier ensemble field: the ensemble of accurate/diverse classifiers can be the best. It means that although the more diverse classifiers, the better ensemble, it is provided that the classifiers are better than random.

Classifier ensemble systems can be categorized by the ways they are built. Four dimensions for characterizing ensemble methods have been proposed: combination level, classifier level, feature level, and data level. In this article, we will focus on the classifier level and feature level that deals with the ways the base classifier are created and some new features may be created.

Bagging [1], Random Forest [5] and AdaBoost [6-8] may be the three most widely used techniques to generate homogeneous classifiers. Bagging trains its diverse classifiers by employing a primary learning algorithm to some bootstrapped sub-samples. These sub-samples are randomly extracted out of the train dataset with replacement and have the same

sample size as that of the train dataset. Random Forest [5] is an ensemble approach that uses a decision tree as its primary classifier. It uses bootstrapped sub-sampling to obtain different train datasets like Bagging. Both Bagging and Random Forest utilize simple majority voting mechanism to aggregate their primary classifiers into a consensus classifier. AdaBoost, the most prominent member in boosting family, generates a series of base classifiers by applying a given base learning algorithm to successive derived training sets that are obtained by either resampling or reweighting the original train dataset in the light of a weight distribution maintained over the training set. AdaBoost initially assigns equal weights to each training instance and in subsequent iterations, it adjusts these weights so that the weight of the instances misclassified by the previously trained classifier is increased, whereas that of the correctly predicted ones is decreased. Thus, AdaBoost attempts to produce new classifiers that are able to better predict the “hard” instances for the previous ensemble members. The final classification is obtained from a weighted vote of the base classifiers.

## II. BACKGROUND

A classifier ensemble will be named a generative classifier ensemble if it produces base classifiers during the training of ensemble. In generative classifier ensemble methods, diversity is usually made using two categories of classifier ensembles. One category of these methods obtains diverse individuals by training classifiers on different training set, such as bagging [1], boosting [25], cross validation [20] and using artificial training examples [13]. Another category of methods for creating diversity employs different structures, different initial weighing, different parameters and different base classifiers to obtain ensemble individuals. For example, [24] adapted the training algorithm of the network by introducing a penalty term to encourage individual networks to be decorrelated. Liu and Yao [12] used negative correlation learning to generate negatively correlated individual neural network.

A classifier ensemble will be named a non-generative classifier ensemble if it produces base classifiers first, then during the training of ensemble it only selects a subset of the ensemble. The non-generative classifier ensemble is also named selective classifier ensemble approach

where the diverse components are selected from a number of trained accurate base classifiers. For example, Opitz and Shavlik [17] proposed a generic algorithm to search for a highly diverse set of accurate networks. Lazarevic and Obradoric [11] proposed a pruning algorithm to eliminate redundant classifiers. Navone et al. [16] proposed another selective algorithm based on bias/variance decomposition. GASEN proposed by Zhou et al. [26] and PSO based approach proposed by Fu et al. [4] also were introduced to select the ensemble components.

### Linear Discriminant Analysis:

Linear discriminant analysis (LDA) is an algorithm used in pattern recognition to discover a linear transformation of attributes. It is widely used to discover a linear mapping of dimensions where data points of different classes in the dataset are discriminated from each other. Indeed, the objective function of LDA is to find the best locations for Gaussian distributions of different clusters and the best parameters for those Gaussian distributions in any given dataset. LDA tries to decrease dimensionality while preserving as much of the class discriminatory information as possible. The output of LDA can be considered as a linear classifier. It can also be considered as dimensionality reduction technique.

However it is commonly considered as a dimensionality reduction. LDA is also closely related to Principal Component Analysis (PCA) in which it explores linear combinations of features which perfectly represent the data. As LDA is supervised, it tries to model the difference between the classes of data. PCA is an unsupervised task. So, PCA does not take into account any difference in class.

### III. PROPOSED METHOD

Before presenting our method, some materials must be clarified. Assume our dataset is always denoted by  $D$ . Our dataset contains  $n$  data points and defined in a  $f$ -dimensional feature space. Also assume that our dataset has  $c$  classes. Let us assume  $T$  is target vector. It means that  $T_i$  is category of  $i$ th data point. Also we assume that  $i$ th data point is denoted by  $D_i$ .

**Definition 1** *data point*: a vector of  $f$  continuous values that represents a data object. So  $D_{ij}$  is  $j$ th feature of  $i$ th data point.

**Definition 2** *hard classifier*: a model that receives a data point and returns a categorical

label.

**Definition 3** *soft classifier*: a model that receives a data point and returns a vector of  $c$  continuous values where each value is in range  $[0,1]$ . The  $j$ th output of a *soft classifier* is denoted by  $o_j$  and represents for *support* of the classifier for  $j$ th class. It means that when a *data point* is given to a *soft classifier*, it produces a vector  $\vec{o}$

that each value of that represents amount of classifier *support* for its corresponding class for the *data point*. It is clear that if you are obliged to select only one class tag for a *data point* using a *soft classifier*, the class with maximum support, is the best candidate. The class with maximum probability (definition 5) is named the *most probable tag* (MPT). The class with the second maximum probability is named *runner-up tag* (RUT).

**Definition 4** *support for a class*: A value in range  $[0,1]$  produced by a *soft classifier* on a *data point* indicating how much the classifier believes the *data point* belongs to that class.

**Definition 5** *probability for a class*: A probability value indicating the *data point* belongs to a class. To compute it, assume a *soft classifier* output over a *data point* is a vector  $\vec{o} = [o_1, o_2, \dots, o_c]$ . The probability vector is

$\vec{p} = [p_1, p_2, \dots, p_c]$  and computed based on equation 1.

$$p_i = \frac{P_i \times o_i}{\sum_{j=1}^c (P_j \times o_j)} \quad (1)$$

where  $P_j$  is the prior probability presented in definition 11 and formulated in equation 2.

**Definition 6** *ensemble support for a class*: A value in range  $[0,1]$  produced by a *soft classifier* on a *data point* indicating how much the classifier believes the *data point* belongs to that class.

**Definition 7** *ensemble*: a  $Ens\_Size$  *soft classifiers* that is denoted by  $E$ . It is worthy to mention that  $E_i$  stands for  $i$ th *soft classifier* of the *ensemble*.

**Definition 8** *ensemble support for a class*: A value in range  $[0,1]$  produced by an ensemble of *soft classifiers* on a *data point* indicating how much the *ensemble* believes the *data point*

belongs to that class. It is the averaged *support of soft classifiers* on that *data point for the class*.

**Definition 9** *hard data point* (HDP): A *data point* will be defined as a *hard data point* if (*ensemble*) *probability* difference between MPT and RUT is more than a threshold. The mentioned threshold that is denoted by *hard\_Th* is a parameter of the algorithm.

**Definition 10** *erroneous data point* (EDP): *ith data point* will be defined as an *erroneous data point* if MPT is not equal to  $T_i$ .

The proposed method gets dataset as input, and puts it into three partitions: *training set*, *test set* and *validation set*. The size of *training set* divided by the size of dataset is named training set ratio and denoted by *TrR*. The size of test set divided by the size of dataset is named test set ratio and denoted by *TeR*. The size of *validation set* divided by the size of dataset is named validation set ratio and denoted by *VaR*. Throughout the paper, training set, test set and validation set are denoted by *TrS*, *TeS* and *VaS* respectively. Also in the paper, target vector of *training set*, *test set* and *validation set* are denoted by *TTrS*, *TTeS* and *TVaS* respectively.

**Definition 11** *Prior Probability*: a  $P_i$  where  $i \in \{1, 2, \dots, c\}$  is computed based on equation 2.

$$P_i = \frac{n_i^{TrS}}{n^{TrS}} \quad (2)$$

where  $n_i^{TrS}$  is the number of *data points* of class  $i$  in *TrS* and  $n^{TrS}$  stands as the number of *data points* in *TrS*. The algorithm is depicted in Fig. 1.

Then the data of each class is extracted from the original validation data set. The proposed algorithm assumes that a classifier is first trained on training set, and then this classifier is added to our ensemble. Now using this classifier, we can obtain erroneous data points on validation data set. Using this work we partition validation data points into two classes: erroneous and non-erroneous. At this step, we label validation data points according the two above classes and then using a pairwise classifier we approximate probability of the error occurrence. This pairwise classifier indeed works as an error detector. Next all data, including training, testing and validation are served as input for that classifier, and then their outputs are considered as new features of those data points. At the next step, using linear

discriminant analysis (LDA) we reduce the dimensionality of the above new space to that of previous space [3]. We repeat this process in predefined number of iterations. Repeating the above process as many as the predefined number of data sets and consequently also that number of classifiers.

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Inputs:
TrR, TeR, VaR, D, c, Ens_Size, Erroneous_Th, T, @TrainSoftClassifier
Output:
E, Error
Begin
D = Shuffle(D)
TrS = round(n × TrR)
TeS = round(n × TeR)
VaS = round(n × VaR)
TrD = D1..TrS
TTrD = T1..TrS
TeD = D(TrS+1)..(TrS+VaS)
TTeD = T(TrS+1)..(TrS+VaS)
VaD = D(TrS+VaS+1)..(TrS+VaS+TeS)
TVaD = T(TrS+VaS+1)..(TrS+VaS+TeS)
For i = 1..c
    Pi =  $\frac{n_i^{TrS}}{n^{TrS}}$ 
End
E = ∅
For i = 1 to Ens_Size
    Ei = TrainSoftClassifier(TrD, TTrD)
End
HD_Ind = HardDataDetection(E, P, D, T, Erroneous_Th)
ED_Ind = ErroneousDataDetection(E, P, D, T)
HDTi =  $\begin{cases} 1 & i \in HD\_Ind \\ 0 & otherwise \end{cases}$ , where  $i \in \{1..(TrS + VaS)\}$ 
EDTi =  $\begin{cases} 1 & i \in ED\_Ind \\ 0 & otherwise \end{cases}$ , where  $i \in \{1..(TrS + VaS)\}$ 
HDClassifier = TrainSoftClassifier(D1..(TrS+VaS), HDT)
EDClassifier = TrainSoftClassifier(D1..(TrS+VaS), EDT)
TestHDPredict = Test(HDClassifier, TeD)
TestEDPredict = Test(EDClassifier, TeD)
NewD = D
NewDi,(f+1)} = HDTi, where  $i \in \{1..(TrS + VaS)\}$ 
NewD(TrS+VaS)+i,(f+1)} = HDClassifieri, where  $i \in \{1..TeS\}$ 
NewDi,(f+2)} = EDTi, where  $i \in \{1..(TrS + VaS)\}$ 
NewD(TrS+VaS)+i,(f+1)} = EDClassifieri, where  $i \in \{1..TeS\}$ 
TrD = NewD1..TrS
TeD = NewD(TrS+1)..(TrS+VaS)
VaD = NewD(TrS+VaS+1)..(TrS+VaS+TeS)
For i = 1 to Ens_Size
    Ei = TrainSoftClassifier(TrD, TTrD)
End
ED_Ind = ErroneousDataDetection(E, P, NewD, T)
Error =  $\frac{|ED\_Ind|}{n}$ 
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Fig. 1. The pseudo code of the proposed combinational algorithm

Pseudo code of the proposed algorithm is shown in Fig. 1. It can be said about time order of this algorithm that the method just multiplies a constant multiplicand in the time order of simple algorithm (training a simple classifier). Suppose that the time order of training a simple classifier on a data set with  $n$  data points and  $c$  classes to be  $O(f(n,c))$ , also assume that in the worst case the time order of training pairwise classifier on that data set to be  $O(g(n,c))$  and also  $m$  to be the number of *max\_iteration* (or that predefined

number). Then the time order of this method is  $\Omega(3*m*f(n,c))$ . Consequently the time order of the method will be  $\Omega(m*f(n,c))$ . This shows time order of the algorithm relevant to just a constant factor is reduced, that this waste of time is completely tolerable against important achieved accuracy.

After creating diverse classifiers for our classifier ensemble, the next step is finding a method to fuse their results and make final decision. The part of making final decision is named combiner part. There are many different combiners. Combination method of base classifier decisions depend on their output type. Some traditional methods of classifier fusion which are based on soft/fuzzy outputs are as below:

Majority vote: assume that we have  $k$  classifiers. Classifier ensemble vote to class  $j$  if a little more than half of base classifiers vote to class  $j$ .

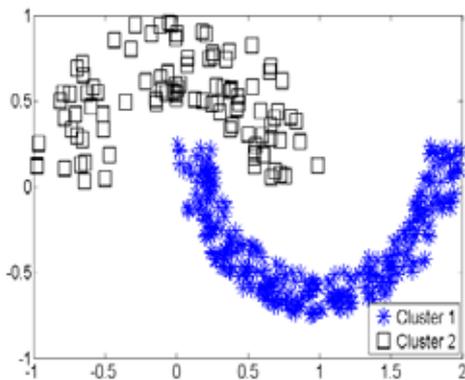


Fig. 2. Half Ring dataset.

Simple average: the average of results of separate classifiers is calculated and then the class that has the most average value is selected as final decision.

Weighted average: it is like simple average except that a weight for each classifier is used for calculating that average.

#### IV. EXPERIMENTAL RESULTS

The metric for evaluating an output of a classifier is accuracy; i.e. the accuracy is taken as the evaluation metric throughout all the paper for reporting performance of classifiers.

The proposed method is examined over 6 different standard datasets and one artificial dataset. These real datasets are available at UCI repository [11]. Brief information about the used datasets is available in Table 1. The details of

HalfRing dataset can be available in [14]. The artificial HalfRing dataset is depicted in Fig. 2. The HalfRing dataset is considered as one of the most challenging dataset for the classification algorithms.

Table 1. Brief information about the used datasets.

#	Dataset Name	# of Class	# of Features	# of Samples	Data distribution per classes
1	HalfRings	2	2	400	300-100
2	Ionosphere	2	34	351	126-225
3	Iris	3	4	150	50-50-50
4	Wine	3	13	178	59-71-48
5	Bupa	2	6	345	145-200
6	BreastCancer	683	9	2	444-239
7	Yeast	1484	8	10	463-5-35-44-51-163-244-429-20-30

The predefined number of max\_iteration in the algorithm is experimentally considered 3 here. Here, train set, test set and validation set are considered to contain 60%, 15% and 25% of entire dataset respectively. The summary of the results are reported in Table 2. All classifiers used in the ensemble are support vector machines (SVM).

Table 2. A summary of seven independent runs of algorithm over "Bupa" dataset

"Bupa"	Iteration 1	Iteration 2	Iteration 3	Ensemble
Run 1	61.77	69.12	48.53	67.65
Run 2	67.65	66.18	73.53	67.65
Run 3	72.06	75.00	70.59	75.00
Run 4	66.18	57.35	64.71	66.18
Run 5	66.18	66.18	67.65	69.12
Run 6	63.24	60.29	66.18	64.71
Run 7	66.18	65.69	65.20	68.14

As it is inferred from Table 2, different iterations have resulted in diverse and usually better accuracies than initial classifier. Of course the ensemble of classifiers is not always better than the best classifier over different iterations, but always it is above the average accuracies and more important is the fact that it almost outperforms initial classifier and anytime it is not worse than the first. Indeed the first classifier (classifier in the iteration 1) is simple classifier that we must compare its results to ensemble results. In the Table 2 each row is one independent run of algorithm, and each column of it is the accuracy obtained using that classifier generated in iteration number corresponds to column number. The ensemble column is the ensemble accuracy of those classifiers generated in iteration number 1-3.

In the second experimentation, the predefined

number of *max\_iteration* in the algorithm is experimentally considered 7 here. Here, train set, test set and validation set are considered to contain 60%, 15% and 25% of entire dataset respectively. The summary of the results are reported in Table 3. All reported results are averaged over 10 distinct runs.

Table 3. Proposed method vs. simple ensemble

	Base Classifier Type	Dataset Name				
		Bupa	Wine	Iris	Breast	Yeast
Proposed Method	MLP	68.48	98.79	95.32	96.49	59.50
	kNN	63.91	79.82	93.25	95.81	58.28
	SVM	68.35	99.01	95.04	96.31	59.82
	DT	70.21	99.58	96.12	96.19	54.78
Simple Ensemble	MLP	67.68	98.16	94.36	96.04	58.11
	kNN	63.86	79.82	93.28	95.95	58.69
	SVM	68.22	98.73	94.88	96.28	59.36
	DT	69.20	99.033	95.14	95.87	54.24

As it can be inferred from Table 3, recognition ratio is improved considerably when DT is the base classifier rather other base classifiers. Because of low number of features and records in Iris, the improvement is more significant on Wine dataset.

Table 3 shows the results of performance of classification accuracy of the proposed method. These results are average of the ten independent runs of the algorithm. In these results, the parameter k in k-Nearest Neighbor algorithm, kNN, is set to one. The MLPs have two hidden layer with 10 and 5 neurons respectively in each of them.

The detailed results of the proposed method comparing with different classification algorithms are presented in Fig 3. To reach these results, 10

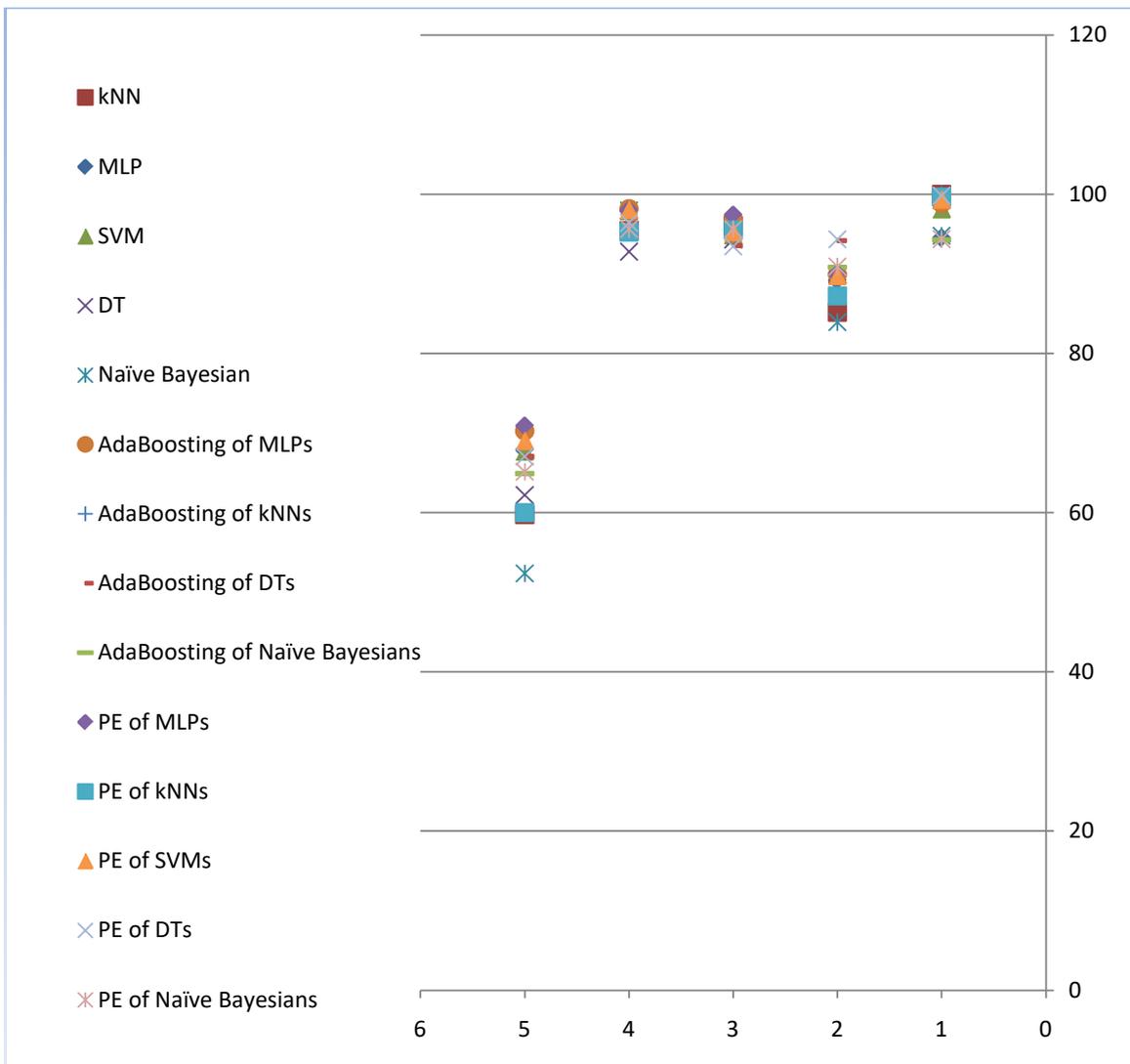


Fig. 3. Performance of different classification methods in terms of accuracy. X axis stands for dataset number.

independent runs of each algorithm are employed and their average accuracy is reported. In each run 66 random percent of dataset is considered as train set and the rest 34% is considered as test set. The results still confirm that the proposed method is promising comparing with other classification algorithms including AdaBoosting. Fig 4 depicts a more detailed comparison between the proposed

method and Adaboosting method. It contains the result of Fig. 3 plus the confidence interval for each method.

Fig. 5 compares the only AdaBoosting method with the proposed method. The results of the Fig 5 show that the proposed method can compete with AdaBoosting. The proposed method can even outperform the AdaBoosting in some cases.

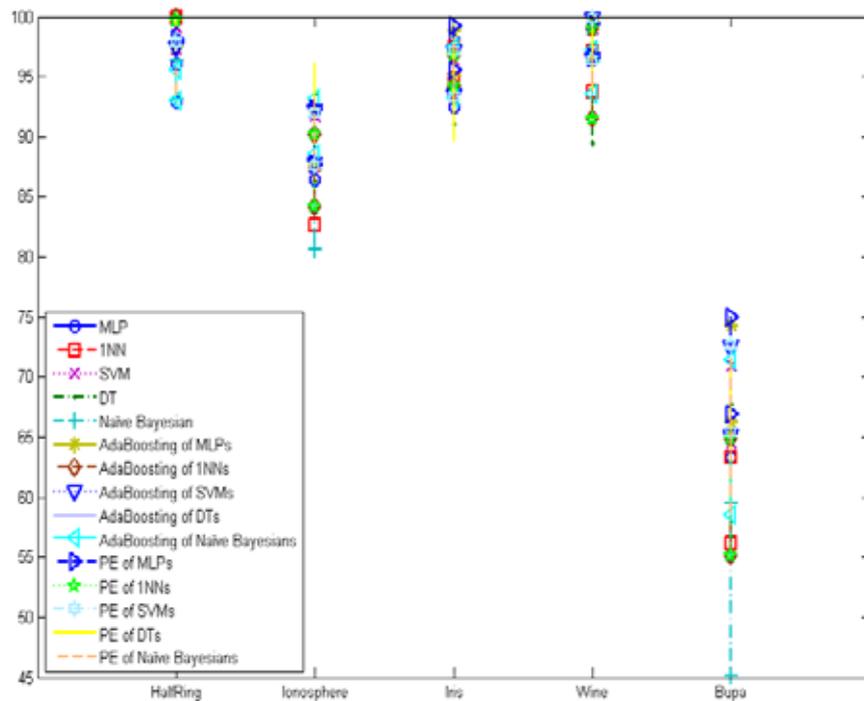


Fig. 4. Performance of different classification methods in terms of accuracy with the confidence interval.

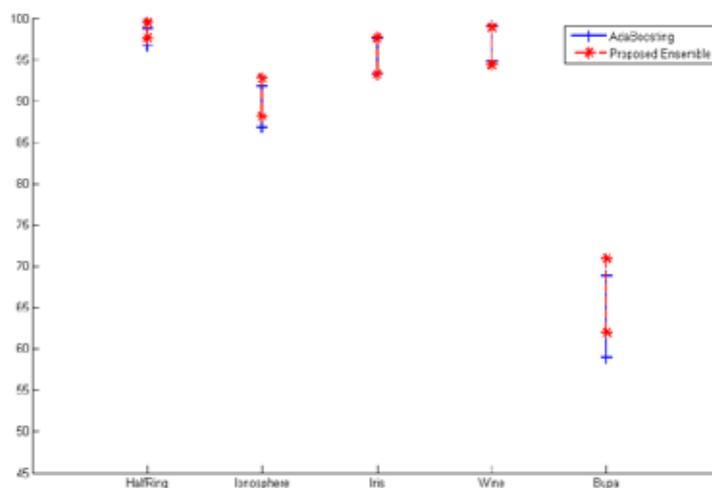


Fig. 5. Performance of the proposed method with AdaBoosting method in terms of accuracy with the confidence interval.

## V. CONCLUSION AND DISCUSSION

Thanks to the good performance of the ensemble methods, they have been employed in various applications. Generally in design of combinational classifier systems, the more diverse the results of the classifiers, the more appropriate the final result. We propose a new method of creating an ensemble. It has been shown that the necessary diversity of an ensemble can be achieved by the proposed algorithm. The method was explained in detail above and the results over some real data sets prove the correctness of our claim. Although the ensemble created by proposed method may not always outperform all of the classifiers existing in ensemble, it always possesses the diversity needed for creation of an ensemble, and consequently it always outperforms the first or the simple classifier. We have also showed that time order of this mechanism is not much more than simple methods. Indeed using manipulation of data set features, we inject the necessary diversity in the ensemble; it means this method is a type of generative methods that manipulates data set in another way different with previous methods such as bagging and boosting.

These results can be due to its emphasis on boundary data points. By emphasizing on different boundary data points, in each iteration we obtain a diverse and well-scattered bag of data and consequently a diverse classifier.

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