

# A Novel Class Imbalance Learning Method using Subset Filtering

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**Abstract**—In many real-world applications, the problem of learning from imbalanced data (the imbalanced learning problem) is a relatively new challenge that has attracted growing attention from both academia and industry. The imbalanced learning problem is concerned with the performance of learning algorithms in the presence of underrepresented data and severe class distribution skews. Due to the inherent complex characteristics of imbalanced data sets, learning from such data requires new understandings, principles, algorithms, and tools to transform vast amounts of raw data efficiently into information and knowledge representation. In this paper, we present a new hybrid subset filtering approach for learning from skewed training data. This algorithm provides a simpler and faster alternative by using C4.5 as base algorithm. We conduct experiments using eleven UCI data sets from various application domains using four base learners, and five evaluation metrics. Experimental results show that our method has higher Area under the ROC Curve, F-measure, precision, TP rate and TN rate values than many existing class imbalance learning methods.

**Index Terms**— Classification, class imbalance, weighted sampling, subset filtering.

A dataset is class imbalanced if the classification categories are not approximately equally represented. The level of imbalance (ratio of size of the majority class to minority class) can be as huge as 1:99 [1]. It is noteworthy that class imbalance is emerging as an important issue in designing classifiers [2], [3], [4]. Furthermore, the class with the lowest number of instances is usually the class of interest from the point of view of the learning task [5]. This problem is of great interest because it turns up in many real-world classification problems, such as remote-sensing [6], pollution detection [7], risk management [8], fraud detection [9], and especially medical diagnosis [10]–[13].

There exist techniques to develop better performing classifiers with imbalanced datasets, which are generally called Class Imbalance Learning (CIL) methods. These methods can be broadly divided into two categories, namely, external methods and internal methods. External methods involve preprocessing of training datasets in order to make them balanced, while internal methods deal with modifications of the learning algorithms in order to reduce their sensitiveness to class imbalance [14]. The main advantage of external methods as previously pointed out, is that they are independent of the underlying classifier. In this paper, we are laying more stress to propose an external CIL method for solving the class imbalance problem.

This paper is organized as follows. Section 2 briefly reviews the Data Balancing problems and its measures. In Section 3, we discuss the proposed method of using the Subset filtering technique for CIL. Section 4 presents the imbalanced datasets used and measures used for validation, while In Section 5, we present the experimental setting and In Section 6 discuss, in detail, the classification results obtained by the proposed method and compare them with the results obtained by different existing methods and finally, in Section 7, we conclude the paper.

## 2. DATA BALANCING

Whenever a class in a classification task is underrepresented (i.e., has a lower prior probability) compared to other classes, we consider

the data as imbalanced [15], [16]. The main problem in imbalanced data is that the majority classes that are represented by large numbers of patterns rule the classifier decision boundaries at the expense of the minority classes that are represented by small numbers of patterns. This leads to high and low accuracies in classifying the majority and minority classes, respectively, which do not necessarily reflect the true difficulty in classifying these classes. Most common solutions to this problem balance the number of patterns in the minority or majority classes.

Either way, balancing the data has been found to alleviate the problem of imbalanced data and enhance accuracy [15], [16], [17]. Data balancing is performed by, e.g., oversampling patterns of minority classes either randomly or from areas close to the decision boundaries. Interestingly, random oversampling is found comparable to more sophisticated oversampling methods [17]. Alternatively, undersampling is performed on majority classes either randomly or from areas far away from the decision boundaries. We note that random undersampling may remove significant patterns and random oversampling may lead to overfitting, so random sampling should be performed with care. We also note that, usually, oversampling of minority classes is more accurate than undersampling of majority classes [17].

Resampling techniques can be categorized into three groups. Undersampling methods, which create a subset of the original data-set by eliminating instances (usually majority class instances); oversampling methods, which create a superset of the original data-set by replicating some instances or creating new instances from existing ones; and finally, hybrids methods that combine both sampling methods. Among these categories, there exist several different proposals; from this point, we only center our attention in those that have been used in under sampling.

- *Random undersampling*: It is a nonheuristic method that aims to balance class distribution through the random elimination of majority class examples. Its major drawback is that it can discard potentially useful data, which could be important for the induction process.

- *Random oversampling*: In the same way as random oversampling, it tries to balance class distribution, but in this case, randomly replicating minority class instances. Several authors agree that this method can increase the likelihood of occurring overfitting, since it makes exact copies of existing instances.
- *Hybrid Methods*: In this hybrid method both undersampling and oversampling will be applied for the datasets so as to make it a balance dataset.

The bottom line is that when studying problems with imbalanced data, using the classifiers produced by standard machine learning algorithms without adjusting the output threshold may well be a critical mistake. This skewness towards minority class (positive) generally causes the generation of a high number of false-negative predictions, which lower the model's performance on the positive class compared with the performance on the negative (majority) class. A comprehensive review of different CIL methods can be found in [18]. The following two sections briefly discuss the external-imbalance and internal-imbalance learning methods.

The external methods are independent from the learning algorithm being used, and they involve preprocessing of the training datasets to balance them before training the classifiers. Different resampling methods, such as random and focused oversampling and undersampling, fall into to this category. In random undersampling, the majority-class examples are removed randomly, until a particular class ratio is met [19]. In random oversampling, the minority-class examples are randomly duplicated, until a particular class ratio is met [18]. Synthetic minority oversampling technique (SMOTE) [20] is an oversampling method, where new synthetic examples are generated in the neighborhood of the existing minority-class examples rather than directly duplicating them. In addition, several informed sampling methods have been introduced in [21]. A clustering-based sampling method has been proposed in [22], while a genetic algorithm based sampling method has been proposed in [23].

### 3. Class Imbalance Learning using Subset Filtering

In this section, we follow a design decomposition approach to systematically analyze the different unbalanced domains. We first briefly introduce the design decomposition methodology adopted for new proposed approach.

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**Algorithm 1** The Proposed Algorithm.

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**1: {Input: A set of minor class examples  $P$ , a set of major class examples  $N_j, j=1, \dots, J$ , and  $T$ , the number of subsets to be sampled from  $N_j$ }**  
**2:  $i \leftarrow 0, T=N/P$ .**  
**3: repeat**  
**4:  $i = i + 1$**   
**5: Randomly sample a subset  $N_i$  from  $N_j, j=1, \dots, J$ .**  
**6: Combine  $P$  and  $N_i$  to form  $NP_i$**   
**6: Apply filter on  $NP_i$**   
**7: Train and Learn A Base Classifier (C4.5) using  $NP_i$ . Obtain the values of AUC, TP, FP, F-Measure**  
**7: until  $i = T$  8: Output: Average Measure;**

The different components of our proposed algorithm are elaborated in the next subsections.

### 3.1 Dataset Sampling

An easy way to sample a dataset is by selecting instances randomly from all classes. However, sampling in this way can break the dataset in an unequal priority way and more number of instances of the same class may be chosen in sampling. To resolve this problem and maintain uniformity in sample, we propose a sampling strategy called weighted component sampling.

Before creating multiple subsets, we will create the number of majority subsets depending upon the number of minority instances.

### 3.2 Identifying number of subsets of majority class

The ratio of majority and minority instances in the unbalanced dataset is used to decide the number of subset of majority instances ( $T$ ) to be created.

$$T = \text{no. of majority inst}(N) / \text{no. of minority inst}(P).$$

### 3.3 Applying filter

Subsets of majority instances are combined with minority subset and multiple balanced subsets are formed. Applying a specific filtering technique at this stage will help to reduce the class imbalance effects. So, Correlation based Feature Subset (CFS) filter is applied at this stage.

### 3.4 Averaging the measures

The subsets of balanced datasets created are used to run multiple times and the resulted values are averaged to find the overall result. In results we have obtained observations for AUC, Precision, F-measure, Sensitivity, Specificity and Accuracy.

## 4. Datasets and measures

We considered four benchmark real-world imbalanced dataset from the UCI machine learning repository [24] to validate our proposed method. Table II summarizes the details of these datasets in the ascending order of the positive-to-negative dataset ratio. This contains the name of the dataset, the total number of examples (Total), attribute, the number of target classes for each dataset, number of minority class examples (#min.), the number of majority class examples (#maj.). These datasets represent a whole variety of domains, complexities, and imbalance ratios. For every data set, we perform a tenfold stratified cross validation. Within each fold, the classification method is repeated ten times considering that the sampling of subsets introduces randomness. The AUC, Precision, F-measure, TP rate and TN Rate of this cross-validation process are averaged from these ten runs. The whole cross-validation process is

repeated for ten times, and the final values from this method are the averages of these ten cross-validation runs.

**Evaluation Criteria:**

To assess the classification results we count the number of true positive (TP), true negative (TN), false positive (FP) (actually negative, but classified as positive) and false negative (FN) (actually positive, but classified as negative) examples. It is now well known that error rate is not an appropriate evaluation criterion when there is class imbalance or unequal costs. In this paper, we use AUC, Precision, F-measure, TP Rate and TN Rate as performance evaluation measures.

Let us define a few well known and widely used measures:

Apart from these simple metrics, it is possible to encounter several more complex evaluation measures that have been used in different practical domains. One of the most popular techniques for the evaluation of classifiers in imbalanced problems is the Receiver Operating Characteristic (ROC) curve, which is a tool for visualizing, organizing and selecting classifiers based on their tradeoffs between benefits (true positives) and costs (false positives).

A quantitative representation of a ROC curve is the area under it, which is known as AUC. When only one run is available from a classifier, the AUC can be computed as the arithmetic mean (macro-average) of TP rate and TN rate:

The Area under Curve (AUC) measure is computed by,

$$AUC = \frac{1 + TP_{RATE} - FP_{RATE}}{2}$$

Or

$$AUC = \frac{TP_{RATE} + TN_{RATE}}{2}$$

On the other hand, in several problems we are especially interested in obtaining high performance on only one class. For example, in the diagnosis of a rare disease, one of the most important things is to know how reliable a positive diagnosis is. For such problems, the precision (or purity) metric is often adopted, which can be defined as the percentage of examples that are correctly labeled as positive:

The Precision measure is computed by,

$$Precision = \frac{TP}{TP + FP}$$

The F-measure Value is computed by,

$$F - measure = \frac{2 \times Precision \times Recall}{Precision + Recall}$$

To deal with class imbalance, sensitivity (or recall) and specificity have usually been adopted to monitor the classification performance on each class separately. Note that sensitivity (also called true positive rate, TP rate) is the percentage of positive examples that are correctly classified, while specificity (also referred to as true negative rate, TN rate) is defined as the proportion of negative examples that are correctly classified:

The True Positive Rate measure is computed by,

$$TruePositiveRate = \frac{TP}{TP + FN}$$

The True Negative Rate measure is computed by,

$$TrueNegativeRate = \frac{TN}{TN + FP}$$

**5. Experimental Settings**

*A. Algorithms and Parameters*

In first place, we need to define a baseline classifier which we use in our proposed algorithm implementation. With this goal, we have used C4.5 decision tree generating algorithm [25]. Furthermore, it has been widely used to deal with imbalanced data-sets [26]–[28], and C4.5 has also been included as one of the top-ten data-mining algorithms [29]. Because of these facts, we have chosen it as the most appropriate base learner. C4.5 learning algorithm constructs the decision tree top-down by the usage of the normalized information gain (difference in entropy) that results from choosing an attribute for splitting the data. The attribute with the highest normalized information gain is the one used to make the decision. To validate the proposed algorithm, we compared it with the traditional C4.5, CART, REP and SMOTE. Eleven real world benchmark data sets taken from the UCI Machine Learning Repository are used throughout the experiments (see Table 1). We performed the implementation using Weka on Windows XP with 2 Duo CPU running on 3.16 GHz PC with 3.25 GB RAM.

*2) Evaluations on Four Real-World Datasets:*

We evaluate the CILSW model on four real-world datasets including Ecolic, Diabetes, Hepatitis and Breast-w datasets. The four datasets are obtained from the University of California at Irvine machine learning repository [24].

We then construct classifiers from the imbalanced data based on the training dataset, and perform evaluation on the test data. We repeat this procedure ten times and use the average of the results as the performance metric. The detailed information about the datasets is described in Table 1.

**Table 1 Summary of benchmark imbalanced datasets**

Datasets	# Ex.	Atts.	Class (.,+)
Ecolic	336	8	(cp, im)
Hepatitis	155	19	(die; live)
Ionosphere	351	34	(b;g)
Labor	56	16	(bad ; good )
Breast	268	9	(recurrence; no-recurrence)
Breast_w	699	9	(benign; malignant)

Diabetes	768	8	(tested-positive; tested-negative)
Vote	435	16	(democrat ;republican )
Sonar	208	61	(Rock, Mine)
Sick	3772	30	(Negative, Sick)

## 6. Experimental Results

We have analysis the performance of our proposed algorithm on class imbalance problem in the following eleven real-world datasets.

(1) **Ecolic Dataset:** This UCI dataset was contributed by Paul Horton. Number of instances in the data set is 101, number of attributes is 7. The number of classes is 8. There are no missing values in this dataset. The results of the tenfold cross validation are shown in Table 2 From Tables 13-17, we can observe the results of Proposed Algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that Proposed algorithm has performed well in the case of AUC improvement, Precision improvement, F-measure improvement and it is comparable in the case of TP Rate and TN Rate. The reason for the better performance of proposed algorithm is due to the multiple class nature of the dataset and the majority and minority ratio of the dataset is very low.

(2)**Diabetes Dataset :**The Pima Indians diabetes data set obtained from the University of California at Irvine (UCI) repository [42] contains 768 samples from two classes with 500 negative samples and 268 positive samples. The positive class is interpreted as “tested positive for diabetes.” There are eight input features for the data samples. The results of the tenfold cross validation are shown in Table 9. From Tables13- 17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given good results on AUC and tie and some underperforming results in the case of remaining measures. The Reason for the performance of proposed algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is very high (i.e. 500:268).

(3) **Hepatitis Dataset:** This data set is used to diagnose whether a hepatitis patient will die or live. Number of instances in the data set is 155, number of attributes is 20, and number of classes is 2 including DIE and LIVE. There are 123 LIVE instances and 32 DIE instances. There are 168 missing values in this data set. The results of the tenfold cross validation are shown in Table 3. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given good results on all the measures. The Reason for the performance of Proposed Algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is very high (i'e.123:32).

(4) **Breast-w Dataset:** This is one of the breast cancer databases at UCI, collected at the University of Wisconsin by W. H.Wolberg. The problem is to predict whether a tissue sample taken from a patient's breast is malignant or benign. There are two classes, ten numerical attributes, and 699 observations. The results of the tenfold cross validation are shown in Table 7. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given moderate results on Breast-w dataset. The Reason for the performance of proposed algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 458:241).

(5) **Breast Cancer Dataset:** This is one of the breast cancer databases at UCI, collected at the University Medical Centre, Institute of Oncology, Ljubljana, Yugoslavia by Ming Tan and Jeff Schlimmer. There are two classes, in which 201 instances of one class and 85 instances of another class. Nine attributes, some of which are linear and some are nominal, and in total 286 observations. There are many missing values in this data set. The results of the tenfold cross validation are shown in Table 6. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given moderate results on Breast-w dataset. The Reason for the performance of proposed algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 201:85).

(6) **Credit-g Dataset:** This UCI dataset was contributed by Hans Hofmann. This UCI dataset is concerned regarding credit card applications. Number of instances in the data set is 1000, number of attributes is 20, out of which 7 are numeric and 13 are nominal. The number of classes is 2. There are no missing values in this dataset. The results of the tenfold cross validation are shown in Table 8. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given underperformed results on Credit-g dataset. The Reason for the moderate performance of proposed algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 700:300).

(7) **Ionosphere Dataset:** This UCI dataset was contributed by Vince Sigillito. This radar data was collected by a system in Goose Bay, Labrador. This system consists of a phased array of 16 high frequency antennas with a total transmitted power on the order of 6.4 kilowatts. Number of instances in the data set is 351, number of attributes is 34. The number of classes is 2. There are no missing attribute values. The results of the tenfold cross validation are shown in Table 4. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given moderate results on Breast-w dataset. The Reason for the performance of proposed algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 225:126).

**Table 1. Tenfold cross validation performance for Ecolic dataset**

System	AUC	Precision	F-measure	TP Rate	TN Rate
<b>C4.5</b>	0.963±0.033	0.935±0.058	0.945±0.040	0.959±0.054	0.948±0.050
<b>CART</b>	0.955±0.032	0.920±0.062	0.944±0.039	0.973±0.041	0.934±0.054
<b>REP</b>	0.950±0.036	0.904±0.071	0.928±0.042	0.959±0.052	0.919±0.071
<b>SMOTE</b>	0.960±0.037	0.935±0.061	0.943±0.041	0.955±0.057	0.948±0.053
<b>Prop. Alg.</b>	0.968±0.038	0.940±0.083	0.943±0.060	0.958±0.077	0.961±0.060

**Table 2. Tenfold cross validation performance for Hepatitis dataset**

System	AUC	Precision	F-measure	TP Rate	TN Rate
<b>C4.5</b>	0.668±0.184	0.510±0.371	0.409±0.272	0.374±0.256	0.900±0.097
<b>CART</b>	0.563±0.126	0.232±0.334	0.179±0.235	0.169±0.236	0.928±0.094
<b>REP</b>	0.619±0.149	0.293±0.386	0.210±0.259	0.187±0.239	0.942±0.093
<b>SMOTE</b>	0.792±0.112	0.709±0.165	0.677±0.138	0.681±0.188	0.837±0.109
<b>Prop. Algor.</b>	0.745±0.186	0.740±0.215	0.705±0.192	0.722±0.248	0.713±0.253

**Table 3 Tenfold cross validation performance for ionosphere dataset**

System	AUC	Precision	F-measure	TP Rate	TN Rate
<b>C4.5</b>	0.891±0.060	0.895±0.084	0.850±0.066	0.821±0.107	0.940±0.055
<b>CART</b>	0.896±0.059	0.868±0.096	0.841±0.070	0.803±0.112	0.921±0.066
<b>REP</b>	0.902±0.054	0.886±0.092	0.848±0.067	0.826±0.104	0.933±0.063
<b>SMOTE</b>	0.904±0.053	0.934±0.049	0.905±0.048	0.881±0.071	0.928±0.057
<b>Prop. Algor.</b>	0.901±0.070	0.928±0.068	0.893±0.072	0.868±0.106	0.921±0.079

**Table 4 Tenfold cross validation performance for labor dataset**

System	AUC	Precision	F-measure	TP Rate	TN Rate
<b>C4.5</b>	0.726±0.224	0.696±0.359	0.636±0.312	0.640±0.349	0.833±0.127
<b>CART</b>	0.750±0.248	0.715±0.355	0.660±0.316	0.665±0.359	0.871±0.151
<b>REP</b>	0.767±0.232	0.698±0.346	0.650±0.299	0.665±0.334	0.765±0.194
<b>SMOTE</b>	0.833±0.127	0.871±0.151	0.793±0.132	0.765±0.194	0.847±0.187
<b>Prop. Algor.</b>	0.856±0.225	0.863±0.246	0.861±0.234	0.890±0.257	0.832±0.267

**Table 5 Tenfold cross validation classification performance for breast\_cancer dataset**

System	AUC	Precision	F-measure	TP Rate	TN Rate
<b>C4.5</b>	0.606±0.087	0.753±0.042	0.838±0.040	0.947±0.060	0.260±0.141
<b>CART</b>	0.587±0.110	0.728±0.038	0.813±0.038	0.926±0.081	0.173±0.164
<b>REP</b>	0.578±0.116	0.721±0.037	0.805±0.042	0.917±0.087	0.151±0.164
<b>SMOTE</b>	0.717±0.084	0.710±0.075	0.730±0.076	0.763±0.117	0.622±0.137
<b>Prop. Algor.</b>	0.596±0.108	0.613±0.074	0.677±0.077	0.767±0.122	0.416±0.164

**Table 6. Tenfold cross validation performance for Breast\_wdataset**

System	AUC	Precision	F-measure	TP Rate	TN Rate
<b>C4.5</b>	0.957±0.034	0.965±0.026	0.962±0.021	0.959±0.033	0.932±0.052
<b>CART</b>	0.950±0.032	0.968±0.026	0.959±0.020	0.952±0.034	0.940±0.051
<b>REP</b>	0.957±0.030	0.965±0.030	0.960±0.021	0.957±0.033	0.931±0.060
<b>SMOTE</b>	0.967±0.025	0.974±0.024	0.960±0.022	0.947±0.035	0.975±0.024
<b>Prop. Algo.</b>	0.956±0.032	0.964±0.039	0.948±0.032	0.935±0.047	0.964±0.039

**Table 7. Tenfold cross validation performance for Credit-g dataset**

	<b>AUC</b>	<b>Precision</b>	<b>F-measure</b>	<b>TPRate</b>	<b>TN Rate</b>
<b>C4.5</b>	0.647±0.062	0.767±0.025	0.805±0.022	0.847±0.036	0.398±0.085
<b>CART</b>	0.716±0.055	0.779±0.030	0.820±0.028	0.869±0.047	0.421±0.102
<b>REP</b>	0.705±0.057	0.765±0.025	0.814±0.026	0.872±0.057	0.371±0.105
<b>SMOTE</b>	0.778±0.041	0.768±0.034	0.787±0.034	0.810±0.058	0.713±0.056
<b>Prop. Algor.</b>	0.718±0.067	0.701±0.59	0.711±0.55	0.728±0.085	0.631±0.099

**Table 8. Tenfold cross validation performance for Pima Diabetes dataset**

<b>System</b>	<b>AUC</b>	<b>Precision</b>	<b>F-measure</b>	<b>TP Rate</b>	<b>TN Rate</b>
<b>C4.5</b>	0.751±0.070	0.797±0.045	0.806±0.044	0.821±0.073	0.603 ±0.111
<b>CART</b>	0.743±0.071	0.782±0.042	0.812±0.040	0.848±0.066	0.554±0.113
<b>REP</b>	0.754±0.060	0.785±0.037	0.809±0.037	0.8384±0.072	0.567±0.105
<b>SMOTE</b>	0.791±0.041	0.781±0.064	0.741±0.046	0.712±0.076	0.807±0.077
<b>Prop Algor</b>	0.795±0.61	0.778±0.075	0.735±0.64	0.706±0.96	0.803±0.88

**Table 9 Tenfold cross validation performance for vote dataset**

<b>System</b>	<b>AUC</b>	<b>Precision</b>	<b>F-measure</b>	<b>TP Rate</b>	<b>TN Rate</b>
<b>C4.5</b>	0.979±0.025	0.971±0.027	0.972±0.021	0.974±0.029	0.953±0.045
<b>CART</b>	0.973±0.027	0.971±0.028	0.966±0.022	0.961±0.037	0.953±0.046
<b>REP</b>	0.957±0.023	0.969±0.035	0.961±0.025	0.955±0.034	0.949±0.059
<b>SMOTE</b>	0.984±0.017	0.977±0.027	0.969±0.021	0.963±0.037	0.981±0.023
<b>Prop. Algor.</b>	0.968±0.031	0.980±0.073	0.946±0.041	0.918±0.071	0.984±0.030

**Table 10 Tenfold cross validation performance for Sonar dataset**

<b>System</b>	<b>AUC</b>	<b>Precision</b>	<b>F-measure</b>	<b>TP Rate</b>	<b>TN Rate</b>
<b>C4.5</b>	0.753±0.113	0.728±0.121	0.716±0.105	0.721±0.140	0.749±0.134
<b>CART</b>	0.721±0.106	0.709±0.118	0.672±0.106	0.652±0.137	0.756±0.121
<b>REP</b>	0.746±0.106	0.733±0.134	0.689±0.136	0.685±0.192	0.762±0.145
<b>SMOTE</b>	0.814±0.090	0.863±0.068	0.861±0.061	0.865±0.090	0.752±0.113
<b>Prop. Algor.</b>	0.772±0.112	0.864±0.096	0.849±0.081	0.847±0.150	0.752±0.182

**Table 11 Tenfold cross validation performance for Sick dataset**

<b>System</b>	<b>AUC</b>	<b>Precision</b>	<b>F-measure</b>	<b>TP Rate</b>	<b>TN Rate</b>
<b>C4.5</b>	0.726±0.224	0.696±0.359	0.636±0.312	0.640±0.349	0.833±0.127
<b>CART</b>	0.750±0.248	0.715±0.355	0.660±0.316	0.665±0.359	0.871±0.151
<b>REP</b>	0.767±0.232	0.698±0.346	0.650±0.299	0.665±0.334	0.765±0.194
<b>SMOTE</b>	0.833±0.127	0.871±0.151	0.793±0.132	0.765±0.194	0.847±0.187
<b>Prop. Algor.</b>	0.935±0.036	0.886±0.052	0.917±0.037	0.953±0.043	0.870±0.065

**Table 13. Summary of results on AUC Vs Prop. Algor.**

System	C4.5	CART	REP	SMOTE
<b>Dataset</b>				
Ecolic	Win	Win	Win	Win
Hepatitis	Win	Win	Win	Loss
Ionosphere	Win	Win	Tie	Loss
Labor	Win	Win	Win	Win
Breast	Loss	Win	Win	Loss
Breast_w	Tie	Win	Tie	Loss
Credit-g	Win	Win	Win	Loss
Diabetes	Win	Win	Win	Loss
Vote	Loss	Loss	Win	Loss
Sonar	Win	Win	Win	Loss
Sick	Win	Win	Win	Win

**Table 14. Summary of results on Precision Vs Prop. Algor.**

System	C4.5	CART	REP	SMOTE
<b>Dataset</b>				
Ecolic	Win	Win	Win	Win
Hepatitis	Win	Win	Win	Win
Ionosphere	Win	Win	Win	Loss
Labor	Win	Win	Win	Loss
Breast	Tie	Loss	Tie	Loss
Breast_w	Loss	Loss	Loss	Loss
Credit-g	Loss	Loss	Loss	Loss
Diabetes	Loss	Loss	Loss	Loss
Vote	Win	Win	Win	Win
Sonar	Win	Win	Win	Tie
Sick	Win	Win	Win	Win

**Table 15. Summary of results on F-Measure Vs Prop. Algor.**

System	C4.5	CART	REP	SMOTE
<b>Dataset</b>				
Ecolic	Tie	Tie	Win	Tie
Hepatitis	Win	Win	Win	Win
Ionosphere	Win	Win	Win	Loss
Labor	Win	Win	Win	Win
Breast	Loss	Loss	Loss	Loss
Breast_w	Loss	Loss	Loss	Loss
Credit-g	Loss	Loss	Loss	Loss
Diabetes	Loss	Loss	Loss	Loss
Vote	Loss	Loss	Loss	Loss
Sonar	Win	Win	Win	Loss
Sick	Win	Win	Win	Win

**Table 16. Summary of results on TP Rate Vs Prop. Algor.**

System	C4.5	CART	REP	SMOTE
<b>Dataset</b>				
Breast	Loss	Loss	Loss	Win
Breast_w	Loss	Loss	Loss	Loss
Credit-g	Loss	Loss	Loss	Loss
Diabetes	Loss	Loss	Loss	Loss
Ecolic	Tie	Loss	Tie	Win
Hepatitis	Win	Win	Win	Win
Ionosphere	Win	Win	Win	Loss
Labor	Win	Win	Win	Win
Vote	Loss	Loss	Loss	Loss
Sonar	Win	Win	Win	Loss

Sick Win Win Win Win

**Table 17. Summary of results on TN Rate Vs Prop. Algor.**

System	C4.5	CART	REP	SMOTE
<b>Dataset</b>				
Ecolic	Win	Win	Win	Win
Hepatitis	Loss	Loss	Loss	Loss
Ionosphere	Loss	Tie	Loss	Win
Labor	Tie	Loss	Win	Loss
Breast	Win	Win	Win	Loss
Breast_w	Win	Win	Win	Loss
Credit-g	Win	Win	Win	Loss
Diabetes	Win	Win	Win	Loss
Vote	Win	Win	Win	Win
Sonar	Win	Loss	Loss	Tie
Sick	Win	Tie	Win	Win

**(8) Labor Dataset:** This UCI dataset was contributed by Stan Matwin. This dataset was used to test 2tier approach with learning from positive and negative examples. Number of instances in the data set is 57, number of attributes is 16, out of which 8 are numeric and 8 are nominal. The number of classes is 2. There are some missing attribute values. The results of the tenfold cross validation are shown in Table 5. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given good results on Labor dataset. One the Reason for the performance of proposed algorithm is due to the small size, the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 37:20).

**(9) Vote Dataset:** This UCI dataset was contributed by This data set is used to predict the result of a vote. It is from 1984 united states congressional voting records database. This data set includes votes for each of the U.S. House of representatives congressmen on the 16 key votes identified by the Congressional Quarterly Almanac (CQA). The CQA lists nine different types of votes: voted for, paired for, and announced for (these three simplified to yea), voted against, paired against, and announced against (these three simplified to nay), voted present, voted present to avoid conflict of interest, and did not vote or otherwise make a position known (these three simplified to an unknown disposition). Number of instances is 435 (267 democrats, 168 republicans), number of attributes is 17, and number of classes is 2. There are 392 missing values in this data set. The results of the tenfold cross validation are shown in Table 7. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given moderate results on Breast-w dataset. The Reason for the performance of proposed algorithm is the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 287:168).

**(10) Sonar Dataset:** This UCI dataset was contributed by Terry Sejnowski. This dataset contains 111 patterns obtained by bouncing sonar signals off a metal cylinder at various angles and under various conditions. This data set can be used in a number of dif-

ferent ways to test learning speed, quality of ultimate learning, ability to generalize, or combinations of these factors. Number of instances in the data set is 208, number of attributes is 60. The number of classes is 2. There are no missing values in dataset. The results of the tenfold cross validation are shown in Table 11. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given good results on Sonar dataset. One the Reason for the performance of proposed algorithm is due to the large size, the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 111:97).

**(11) Sick Dataset:** This UCI dataset was contributed by the Garavan Institute and J. Ross Quinlan, New South Wales Institute, Sydney, Australia. Number of instances in the data set is 3772, number of attributes is 29, out of which 7 are numeric and 22 are nominal. The number of classes is 2. There are some missing values in dataset. The results of the tenfold cross validation are shown in Table 12. From Tables 13-17, we can observe the results of proposed algorithm Vs various algorithms with respect to AUC, Precision, F-measure, TP rate and TN rate. From all the tables we can conclude that proposed algorithm has given excellent results on Sick dataset. One the Reason for the performance of proposed algorithm is due to the very huge size of the dataset, irrelevant attributes present in the dataset, the multi class nature of the dataset and the majority and minority ratio of the dataset is moderately high (i'e: 3541:231).

## 7. Conclusion:

In this paper we present the class imbalance problem paradigm, which exploits the subset filtering strategy in the supervised learning research area, and implement it with C4.5 as its base learner. Experimental results show that our proposed algorithm performed well in the case of multi class imbalance datasets. Furthermore, our proposed algorithm is much less volatile than C4.5. In our future work, we will apply our proposed algorithm to more learning tasks, especially high dimensional feature learning tasks.

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