# Classification and Novel Class Detection in Concept-Drifting Data Streams under Time Constraints

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**Abstract**—Most existing data stream classification techniques ignore one important aspect of stream data: arrival of a novel class. We address this issue and propose a data stream classification technique that integrates a novel class detection mechanism into traditional classifiers, enabling automatic detection of novel classes before the true labels of the novel class instances arrive. Novel class detection problem becomes more challenging in the presence of concept-drift, when the underlying data distributions evolve in streams. In order to determine whether an instance belongs to a novel class, the classification model sometimes needs to wait for more test instances to discover similarities among those instances. A maximum allowable wait time  $T_c$  is imposed as a time constraint to classify a test instance. Furthermore, most existing stream classification approaches assume that the true label of a data point can be accessed immediately after the data point is classified. In reality, a time delay  $T_l$  is involved in obtaining the true label of a data point since manual labeling is time consuming. We show how to make fast and correct classification decisions under these constraints and apply them to real benchmark data. Comparison with state-of-the-art stream classification techniques prove the superiority of our approach.

Index Terms—Data streams, concept-drift, novel class, ensemble classification, K-means clustering, k-nearest neighbor classification, silhouette coefficient.

## **1** INTRODUCTION

ATA stream classification poses many challenges, some of which have not been addressed yet. Most existing data stream classification algorithms [2], [6], [11], [13], [17], [22], [27], [29] address two major problems related to data streams: their "infinite length" and "concept-drift." Since data streams have infinite length, traditional multipass learning algorithms are not applicable as they would require infinite storage and training time. Concept-drift occurs in the stream when the underlying concept of the data changes over time. Thus, the classification model must be updated continuously so that it reflects the most recent concept. However, another major problem is ignored by most state-of-the-art data stream classification techniques, which is "concept-evolution," that means emergence of a novel class. Most of the existing solutions assume that the total number of classes in the data stream is fixed. But in real-world data stream classification problems, such as intrusion detection, text classification, and fault detection,

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new intrusion). Traditional data stream classification techniques would be unable to detect the novel class until the classification models are trained with labeled instances of the novel class. Thus, all novel class instances will go undetected (i.e., misclassified) until the novel class is manually detected by experts, and training data with the instances of that class is made available to the learning algorithm. We address this concept-evolution problem and provide a solution that handles all three problems, namely, infinite length, concept-drift, and concept-evolution. Novel class detection should be an integral part of any realistic data stream classification technique because of the evolving nature of streams. It can be useful in various domains, such as network intrusion detection [12], fault detection [8], and credit card fraud detection [27]. For example, in case of intrusion detection, a new kind of intrusion might go undetected by traditional classifier, but our approach should not only be able to detect the intrusion, but also deduce that it is a new kind of intrusion. This discovery would lead to an intense analysis of the intrusion by human experts in order to understand its cause, find a remedy, and make the system more secure.

novel classes may appear at any time in the stream (e.g., a

Note that in our technique we consider mining from only one stream. We address the infinite length problem by dividing the stream into equal-sized chunks so that each chunk can be accommodated in memory and processed online. Each chunk is used to train one classification model as soon as all the instances in the chunk are labeled. We handle concept-drift by maintaining an ensemble of M such classification models. An unlabeled instance is classified by taking majority vote among the classifiers in the ensemble.

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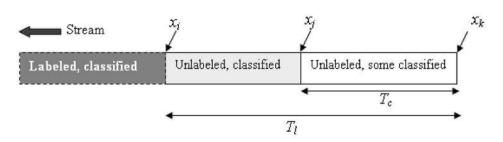


Fig. 1. Illustration of  $T_l$  and  $T_c$ .

The ensemble is continuously updated so that it represents the most recent concept in the stream. The update is performed as follows: As soon as a new model is trained, one of the existing models in the ensemble is replaced by it, if necessary. The victim is chosen by evaluating the error of each of the existing models in the ensemble on the latest labeled chunk, and discarding the one with the highest error. Our approach provides a solution to concept-evolution problem by enriching each classifier in the ensemble with a novel class detector. If all of the classifiers discover a novel class, then arrival of a novel class is declared, potential novel class instances are separated and classified as "novel class." Thus, novel class can be automatically identified without manual intervention.

Our novel class detection technique is different from traditional "one-class" novelty detection techniques [16], [21], [28] that can only distinguish between the normal and anomalous data. That is, traditional novelty detection techniques assume that there is only one "normal" class and any instance that does not belong to the normal class is an anomaly/novel class instance. Therefore, they are unable to distinguish among different types of anomaly. But our approach offers a "multiclass" framework for the novelty detection problem that can distinguish between different classes of data and discover the emergence of a novel class. Furthermore, traditional novelty detection techniques simply identify data points as outliers/anomalies that deviate from the "normal" class. But our approach not only detects whether a single data point deviates from the existing classes, but also discovers whether a group of such outliers possess the potential of forming a new class by showing strong cohesion among themselves. Therefore, our approach is a synergy of a "multiclass" classification model and a novel class detection model.

Traditional stream classification techniques also make impractical assumptions about the availability of labeled data. Most techniques [6], [11], [29] assume that the true label of a data point can be accessed as soon as it has been classified by the classification model. Thus, according to their assumption, the existing model can be updated immediately using the labeled instance. In reality, we would not be so lucky in obtaining the label of a data instance immediately, since manual labeling of data is time consuming and costly. For example, in a credit card fraud detection problem, the actual labels (i.e., authentic/fraud) of credit card transactions usually become available in the next billing cycle after a customer reviews all his transactions in the last statement and reports fraud transactions to the credit card company. Thus, a more realistic assumption would be to have a data point labeled after  $T_l$  time units of its arrival. For simplicity, we assume that the *i*th instance in the stream arrives at *i*th time unit. Thus,  $T_l$  can be considered as a time constraint

imposed on data labeling process. Note that traditional stream classification techniques assume  $T_l = 0$ . Finally, we impose another time constraint,  $T_c$ , on classification decision. That is, an instance must be classified by the classification model within  $T_c$  time units of its arrival. If it is assumed that there is no concept-evolution, it is customary to have  $T_c = 0$ , i.e., an instance should be classified as soon as it arrives. However, when new concepts evolve, classification decision may have to be postponed until enough instances are seen by the model to gain confidence in deciding whether an instance belongs to a novel class or not.  $T_c$  is the maximum allowable time up to which the classification decision can be postponed. Note that  $T_c < T_l$  must be maintained in any practical classification model. Otherwise, we would not need the classifier at all, we could just wait for the labels to arrive.

#### 1.1 Example Illustrating Time Constraints

Fig. 1 illustrates the significance of  $T_l$  and  $T_c$  with an example. Here,  $x_k$  is the last instance that has arrived in the stream. Let  $x_i$  be the instance that arrived  $T_c$  time units earlier, and  $x_i$  be the instance that arrived  $T_l$  time units earlier. Then,  $x_i$  and all instances that arrived before  $x_i$ (shown with dark-shaded area) are labeled since all of them are at least  $T_l$  time units old. Similarly,  $x_i$  and all instances that arrived before  $x_i$  (both the light-shaded and darkshaded areas) are classified by the classifier since they are at least  $T_c$  time units old. However, the instances inside the light-shaded area are unlabeled. Instances that arrived after  $x_i$  (age less than  $T_c$ ) are unlabeled and may or may not be classified (shown with the unshaded area). In summary,  $T_l$ is enforced/utilized by labeling an instance x after  $T_l$  time units of its arrival, and  $T_c$  is enforced by classifying x within  $T_c$  time units of its arrival, for every instance x in the stream.

Integrating classification with novel class detection is a nontrivial task, especially in the presence of concept-drift, and under time constraints. We assume an important property of each class: the data points belonging to the same class should be closer to each other (cohesion) and should be far apart from the data points belonging to other classes (separation). If a test instance is well separated from the training data, it is identified as an Foutlier. Foutliers have potential to be a novel class instance. However, we must *wait* to see whether more such *Foutliers* appear in the stream that observe strong cohesion among themselves. If a sufficient number of such strongly cohesive Foutliers are observed, a novel class is assumed to have appeared, and the Foutliers are classified as a novel class instance. However, we can defer the classification decision of a test instance at most  $T_c$  time units after its arrival, which makes the problem more challenging. Furthermore, we must keep detecting novel class instances in this "unsupervised" fashion for at least  $T_l$  time units from the arrival of the

first novel class instance since labeled training data of the novel class(es) would not be available before that.

We have several contributions. First, to the best of our knowledge, no other data stream classification techniques address the concept-evolution problem. This is a major problem with data streams that must be dealt with. In this light, this paper offers a more realistic solution to data stream classification. Second, we propose a more practical framework for stream classification by introducing time constraints for delayed data labeling and making classification decision. Third, our proposed technique enriches traditional classification model with a novel class detection mechanism. Finally, we apply our technique on both synthetic and real-world data and obtain much better results than state-of-the-art stream classification algorithms.

The rest of the paper is organized as follows: Section 2 discusses related work. Section 3 provides an overview of our approach and Section 4 discusses our approach in detail. Section 5 then describes the data sets and experimental evaluation of our technique. Finally, Section 6 concludes with directions to future works.

## 2 RELATED WORK

Our technique is related to both data stream classification and novelty detection. Data stream classification has been an interesting research topic for years, and many approaches are available. These approaches fall into one of two categories: single model and ensemble classification. Single model classification techniques maintain and incrementally update a single classification model and effectively respond to concept-drift [6], [11], [29]. Several ensemble techniques for stream data mining have been proposed [9], [13], [17], [22], [27]. Ensemble techniques require relatively simpler operations to update the current concept than their single model counterparts, and also handle concept-drift efficiently. Our approach follows the ensemble technique. However, our approach is different from all other stream classification techniques in two different aspects. First, none of the existing techniques can detect novel classes, but our technique can. Second, our approach is based on a more practical assumption about the time delay in data labeling, which is not considered in most of the existing algorithms.

Our technique is also related to novelty/anomaly detection. Markou and Singh study novelty detection in details in [16]. Most novelty detection techniques fall into one of two categories: parametric, and nonparametric. Parametric approaches assume a particular distribution of data and estimate parameters of the distribution from the normal data. According to this assumption, any test instance is assumed to be novel if it does not follow the distribution [19], [21]. Our technique is a nonparametric approach, and therefore, it is not restricted to any specific data distribution. There are several nonparametric approaches available, such as parzen window method [28], k-nearest neighbor (k-NN)-based approach [30], kernelbased method [3], and rule-based approach [15].

Our approach is different from the above novelty/ anomaly detection techniques in three aspects. First, existing novelty detection techniques only consider whether a test point is significantly different from the normal data. However, we not only consider whether a test instance is sufficiently different from the training data, but also consider whether there are strong similarities among such test instances. Therefore, existing techniques discover novelty individually in each test point, whereas our technique discovers novelty collectively among several coherent test points to detect the presence of a novel class. Second, our model can be considered as a "multiclass" novelty detection technique since it can distinguish among different classes of data, and also discover emergence of a novel class. But existing novelty detection techniques can only distinguish between normal and novel, and therefore, can be considered as "one-class" classifiers. Finally, most of the existing novelty detection techniques assume that the "normal" model is static, i.e., there is no concept-drift in the data. But our approach can detect novel classes even if concept-drift occurs in the existing classes.

Novelty detection is also closely related to outlier detection techniques. There are many outlier detection techniques available, such as [1], [4], [5], [14]. Some of them are also applicable to data streams [24], [25]. However, the main difference with these outlier detection techniques from ours is that our primary objective is novel class detection, not outlier detection. Outliers are the by-product of intermediate computation steps in our algorithm. Thus, the precision of our outlier detection technique is not too critical to the overall performance of our algorithm.

Spinosa et al. [23] propose a cluster-based novel concept detection technique that is applicable to data streams. However, this is also a "single-class" novelty detection technique, where authors assume that there is only one "normal" class and all other classes are novel. Thus, it not directly applicable to a multiclass environment, where more than one classes are considered as "normal" or "non-novel." But our approach can handle any number of existing classes, and also detect a novel class that do not belong to any of the existing classes. Therefore, our approach offers a more practical solution to the novel class detection problem, which has been proved empirically.

This paper significantly extends our previous work on novel class detection [18] in several ways. First, in our previous work, we did not consider the time constraints  $T_l$ and  $T_c$ . Therefore, the current version is more practical than the previous one. These time constraints impose several restrictions on the classification algorithm, making classification more challenging. We encounter these challenges and provide efficient solutions. Second, it adds considerable amount of mathematical analysis over the previous version. Third, evaluation is done in a more realistic way (continuous evaluation rather than chunk by chunk evaluation) and a newer version of baseline technique [23] is used (version 2008 instead of 2007). Finally, more figures, discussions, and experiments are added for improved readability, clarity, and analytical enrichment.

## **3** OVERVIEW

At first, we mathematically formulate the data stream classification problem.

• The data stream is a continuous sequence of data points: {*x*<sub>1</sub>,...,*x*<sub>now</sub>}, where each *x<sub>i</sub>* is a *d*-dimensional feature vector. *x*<sub>1</sub> is the very first data point in the stream, and *x<sub>now</sub>* is the latest data point that has just arrived.

TABLE 1 Commonly Used Symbols and Terms

C	Total number of classes in a data stream	$c_i$	A class label			
$D_i$	A data chunk	H	A cluster			
h	A pseudopoint created from cluster $H$	K	Number of clusters created per chunk			
k	Parameter $k$ of a $k$ -NN classifier	M	Ensemble size			
	The ensemble	$L_i$	The <i>i</i> -th model in the ensemble			
q	Parameter $q$ of the $q$ -nearest neighborhood ( $q$ -NH)	S	Chunk size			
$T_l$	Labeling time constraint	$T_c$	Classification time constraint			
<i>Novel class</i> A class that appears due to concept-evolution						

- Each data point  $x_i$  is associated with two attributes:  $y_i$  and  $t_i$  being its class label and time of arrival, respectively.
- For simplicity, we assume that  $t_{i+1} = t_i + 1$  and  $t_1 = 1$ .
- The latest *T<sub>l</sub>* instances in the stream: {*x*<sub>now-*T*<sub>l</sub>+1</sub>,..., *x*<sub>now</sub>} are unlabeled, meaning, their corresponding class labels are unknown. But the class labels of all other data points are known.
- We are to predict the class label of  $x_{now}$  before the time  $t_{now} + T_c$ , i.e., before the data point  $x_{now+T_c}$  arrives, and  $T_c < T_l$ .

Table 1 summarizes the most commonly used symbols and terms used throughout the paper.

## 3.1 Top Level Algorithm

Algorithm 1 outlines the top level overview of our approach. The algorithm starts with building the initial ensemble of models  $L = \{L_1, \ldots, L_M\}$  with the first M labeled data chunks. The algorithm maintains three buffers: buffer buf keeps potential novel class instances, buffer U keeps unlabeled data points until they are labeled, buffer  $\mathcal{L}$  keeps labeled instances until they are used to train a new classifier. After initialization, the while loop begins from line 5, which continues indefinitely. At each iteration of the loop, the latest data point in the stream,  $x_i$  is classified (line 7) using Classify() (Algorithm 2). The novel class detection mechanism is implemented inside Algorithm 2. If the class label of  $x_i$ cannot be predicted immediately, it is stored in buf for future processing. Details of this step will be discussed in Section 4.  $x_i$  is then pushed into the unlabeled data buffer U (line 8). If the buffer size exceeds  $T_l$ , the oldest element  $x_k$  is dequeued and labeled (line 9) since  $T_l$  units of time has elapsed since  $x_k$ arrived in the stream (so it is time to label  $x_k$ ). The pair  $\langle x_k, y_k \rangle$  is pushed into the labeled data buffer  $\mathcal{L}$  (line 9). When we have S instances in  $\mathcal{L}$ , where S is the chunk size, a new classifier L' is trained using the chunk (line 13). Then, the existing ensemble is updated (line 14) by choosing the best M classifiers from the M + 1 classifiers  $L \cup \{L'\}$  based on their accuracies on  $\mathcal{L}$ , and the buffer  $\mathcal{L}$  is emptied to receive the next chunk of training data (line 15). Our algorithm will be mentioned henceforth as "ECSMiner" (pronounced like ExMiner), which stands for Enhanced Classifier for Data Streams with novel class Miner. We believe that any base learner can be enhanced with the proposed novel class detector, and used in ECSMiner. The only operation that needs to be treated specially for a particular base learner is *Train-and-save-decision-boundary*. We illustrate this operation for two base learners in this section.

## Algorithm 1. ECSMiner

- 1:  $L \leftarrow$  Build-initial-ensemble()
- 2:  $buf \leftarrow empty //temporary buffer$
- 3:  $U \leftarrow \text{empty } //\text{unlabeled data buffer}$
- 4:  $\mathcal{L} \leftarrow$  empty //labeled data buffer (training data)

#### 5: while true do

- 6:  $x_j \leftarrow$  the latest data point in the stream
- 7: **Classify**( $L, x_j, buf$ ) //(Algorithm 2, Section 4)
- 8:  $U \Leftarrow x_j$  //enqueue
- 9: **if**  $|U| > T_l$  **then** //time to label the oldest instance
- 10:  $x_k \leftarrow U //\text{dequeue the instance}$
- 11:  $\mathcal{L} \Leftarrow \langle x_k, y_k \rangle$  //label it and save in training buffer
- 12: **if**  $|\mathcal{L}| = S$  **then** //training buffer is full
- 13:  $L' \leftarrow$ **Train-and-save-decision-boundary**( $\mathcal{L}$ ) (Section 3.5)
- 14:  $L \leftarrow \text{Update}(L, L', \mathcal{L})$
- 15:  $\mathcal{L} \leftarrow \text{ empty}$
- 16: **end if**
- 17: end if
- 18: end while

#### 3.2 Nearest Neighborhood Rule

We assume that the instances belonging to a class c are generated by an underlying generative model  $\theta_c$ , and the instances in each class are independently identically distributed. With this assumption, one can reasonably argue that the instances which are close together under some distance metric are supposed to be generated by the same model, i.e., belong to the same class. This is the basic assumption for nearest neighbor classifications [7]. Besides, this assumption is used in numerous semisupervised learning techniques, such as [20], and in many other semisupervised learning works [31]. We generalize this assumption by introducing the concept of "nearest neighborhood."

**Definition 1** ( $\lambda_{c,q}$ -neighborhood).  $\lambda_{c,q}$ -neighborhood, or  $\lambda_{c,q}(x)$  of any instance x is the set of q-nearest neighbors of x within class c.

For example, let there be three classes  $c_+$ , and  $c_-$ , and  $c_0$  denoted by the symbols "+," "-," and black dots,

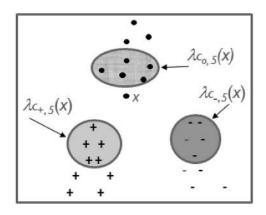


Fig. 2. Illustrating  $\lambda_{e,q}(x)$  for q = 5.

respectively (Fig. 2). Also, let q = 5. Then,  $\lambda_{c_+,q}(x)$  of any arbitrary instance x is the set of five nearest neighbors of x in class  $c_+$ , and so on.

Let  $D_{c,q}(x)$  be the mean distance from x to  $\lambda_{c,q}(x)$ , i.e.,

$$\bar{D}_{c,q}(x) = \frac{1}{q} \sum_{x_i \in \lambda_{c,q}(x)} D(x, x_i), \qquad (1)$$

where  $D(x_i, x_j)$  is the distance between the data points  $x_i$  and  $x_j$  in some appropriate metric.

Let  $c_{min}$  be the class label such that  $\overline{D}_{c_{min},q}(x)$  is the minimum among all  $\overline{D}_{c,q}(x)$ , i.e.,  $\lambda_{c_{min},q}(x)$  is the nearest  $\lambda_{c,q}(x)$  neighborhood (or *q*-nearest neighborhood or *q*-NH) of *x*. For example, in Fig. 2,  $c_{min} = c_0$ , i.e.,  $\lambda_{c_0,q}(x)$  is the *q*-NH of *x*.

**Definition 2 (q-NH Rule).** Let  $c_{min}$  be the class label of the instances in q-NH of x. According to the q-NH rule, the predicted class label of x is  $c_{min}$ .

In the example of Fig. 2,  $c_{min} = c_0$ , therefore, the predicted class label of *x* is  $c_0$ . Our novel class detection technique is based on the assumption that any class of data follow the *q*-NH rule. In Section 4, we discuss the similarity of this rule with k-NN rule, and highlight its significance.

## 3.3 Novel Class and its Properties

**Definition 3 (Existing Class and Novel Class).** Let *L* be the current ensemble of classification models. A class *c* is an existing class if at least one of the models  $L_i \in L$  has been trained with the instances of class *c*. Otherwise, *c* is a novel class.

Therefore, if a novel class c appears in the stream, none of the classification models in the ensemble will be able to correctly classify the instances of c. An important property of the novel class follows from the q-NH rule.

**Property 1.** Let x be an instance belonging to a novel class c, and let c' be an existing class. Then, according to q-NH rule,  $\overline{D}_{c,q}(x)$ , i.e., the average distance from x to  $\lambda_{c,q}(x)$  is smaller than  $\overline{D}_{c',q}(x)$ , the average distance from x to  $\lambda_{c',q}(x)$ , for any existing class c'. In other words, x is closer to the neighborhood of its own class (cohesion), and farther from the neighborhood of any existing classes (separation).

Fig. 3 shows an hypothetical example of a decision tree and the appearance of a novel class. A decision tree and its corresponding feature vector partitioning by its leaf nodes are shown in the figure. The shaded portions of the feature

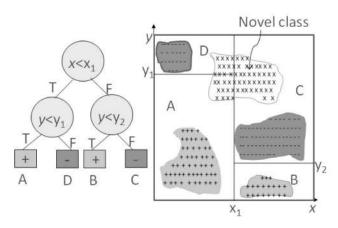


Fig. 3. A decision tree and corresponding feature space partitioning. Shaded areas represent the training data. A novel class arrives in the stream that follows Property 1.

space represent the training data. After the decision tree is built, a novel class appears in the stream (shown with "x" symbol). The decision tree model misclassifies all the instances in the novel class as existing class instance since the model is unaware of the novel class. Our goal is to detect the novel class without having to train the model with that class. Note that instances in the novel class follow Property 1 since the novel-class neighborhood of any novel class instance is much closer to the instance than the neighborhoods of any other classes. If we observe this property in a collection of unlabeled test instances, we can detect the novel class. This is not a trivial task since we must decide when to classify an instance immediately, and when to postpone the classification decision, and wait for more test instances so that Property 1 can be revealed among those instances. Because in order to discover Property 1 (cohesion), we need to deal with a collection of test instances simultaneously. Besides, we cannot defer the decision more than  $T_c$  time units after the arrival of a test instance.

Therefore, the *main challenges* in novel class detection are as follows:

- 1. saving the training data efficiently without using much memory;
- knowing when to classify a test instance immediately, and when to postpone the classification decision;
- 3. classifying the deferred instances within  $T_c$  time unit; and
- 4. predicting the presence of a novel class quickly and correctly.

## 3.4 Base Learners

We apply our technique on two different classifiers: decision tree and k-nearest neighbor. When decision tree is used as a classifier, each training data chunk is used to build a decision tree. When k-NN is used, each chunk is used to build a k-NN classification model. The simplest way to build such a model is to just store all the data points of the training chunk in memory. But this strategy would lead to a inefficient classification model, both in terms of memory and running time. In order to make the model more efficient, we build *K* clusters with the training data [17]. We apply a semisupervised clustering technique using

Expectation Maximization (E-M) that tries to minimize both intracluster dispersion (same objective as unsupervised K-means) and cluster impurity. After building the clusters, we save the cluster summary of each cluster (centroid and frequencies of data points belonging to each class) in a data structure called "microcluster," and discard the raw data points. Since we store and use only K microclusters, both the time and memory requirements become functions of K (a constant number). A test instance  $x_j$  is classified as follows: we find the microcluster whose centroid is nearest from  $x_j$  and assign it a class label that has the highest frequency in that microcluster.

## 3.5 Creating Decision Boundary during Training

The training data are clustered using *K*-means and the summary of each cluster is saved as "pseudopoint." Then, the raw training data are discarded. These pseudopoints form a decision boundary for the training data.

## 3.5.1 Clustering

*K* clusters are built per chunk from the training data. This clustering step is specific to each base learner. For example, For k-NN, existing clusters are used that were created using the approach discussed in Section 3.4. For decision tree, clustering is done at each leaf node of the tree since we need to create decision boundaries in each leaf node separately. This is done as follows: Suppose *S* is the chunk size. During decision tree training, when a leaf node  $l_i$  is reached,  $k_i = (t_i/S) * K$  clusters are built in that leaf, where  $t_i$  denotes the number of training instances belonging to leaf node  $l_i$ . Therefore, the number of training instances built in each leaf node is proportional to the number of training instances that belong to the leaf node. If a leaf node is not empty (has one or more instances), then at least one cluster is built in that node.

## 3.5.2 Storing the Cluster Summary Information

For each cluster, we store the following summary information in a data structure called pseudopoint:

- 1. Weight, w: Total number of points in the cluster;
- 2. Centroid,  $\mu$ ;
- 3. *Radius; R*: Distance between the centroid and the farthest data point in the cluster; and
- 4. *Mean distance,*  $\mu d$ : The mean distance from each point to the cluster centroid.

So, w(h) denotes the "weight" value of a pseudopoint h and so on. After computing the cluster summaries, the raw data are discarded and only the pseudopoints are stored in memory. Any pseudopoint having too few (less than 3) instances is considered as noise and is also discarded. Thus, the memory requirement for storing the training data becomes constant, i.e., O(K).

Each pseudopoint *h* corresponds to a hypersphere in the feature space having center  $\mu(h)$  and radius  $\mathcal{R}(h)$ . Let us denote the portion of feature space covered by a pseudopoint *h* as the "region" of *h* or RE(h). Therefore,  $RE(L_i)$  denotes the union of the regions of all pseudopoints *h* in the classifier  $L_i$ , i.e.,  $RE(L_i) = \bigcup_{h \in L_i} RE(h)$ .  $RE(L_i)$  forms a decision boundary for the training data of classifier  $L_i$ . The decision boundary for the ensemble of classifiers *L* is the union of the decision boundaries of all classifiers in the ensemble, i.e.,

 $RE(L) = \bigcup_{L_i \in L} RE(L_i)$ . The decision boundary plays an important role in novel class detection. It defines the physical boundary of existing class instances. Lemma 1 emphasizes the significance of the decision boundary in distinguishing the existing class instances from novel class instances.

- **Lemma 1.** Let x be a test instance inside the decision boundary RE(L). That is, there is a pseudopoint h such that the distance from x to the center of h is less than or equal to the radius of h, i.e.,  $D(x, \mu(h)) \leq \mathcal{R}(h)$ . Then, x must be an existing class instance.
- **Proof.** Without loss of generality, let D(a, b) be the square of euclidean distance between a and b, i.e.,  $D(a, b) = (a b)^2$ . Note that  $\mathcal{R}(h)$  is the distance between  $\mu(h)$  and the farthest data point in the corresponding cluster H. Let the data point be x'. Therefore,  $D(\mu, x') = \mathcal{R}(h)$ . Also, x' is an existing class data point since it is a training instance that was used to form the cluster. Let  $x_i \in H$  be an arbitrary data point in cluster H, and the total number of data points in H is n, i.e., w(h) = n. In order to simplify notation, we use  $\mu$  instead of  $\mu(h)$  in the proof.

From the lemma statement, we can deduce that

$$\begin{split} D(\mu, x) &\leq D(\mu, x') \Rightarrow (x - \mu)^2 \leq (x' - \mu)^2 \Rightarrow x^2 - 2x\mu \\ &+ \mu^2 \leq x'^2 - 2x'\mu + \mu^2 \\ \Rightarrow x^2 - 2x\mu \leq x'^2 - 2x'\mu \Rightarrow x^2 - 2x\frac{1}{n}\sum_{x_i \in H} x_i \leq x'^2 \\ &- 2x'\frac{1}{n}\sum_{x_i \in H} x_i \text{(by definition of } \mu) \\ \Rightarrow x^2 - 2x\frac{1}{n}\sum_{x_i \in H} x_i + \frac{1}{n}\sum_{x_i \in H} x_i^2 \leq x'^2 - 2x'\frac{1}{n}\sum_{x_i \in H} x_i \\ &+ \frac{1}{n}\sum_{x_i \in H} x_i^2 \left( \text{adding } \frac{1}{n}\sum_{x_i \in H} x_i^2 \text{ on both sides} \right) \\ \Rightarrow \frac{1}{n}\sum_{x_i \in H} (x^2 - 2xx_i + x_i^2) \leq \frac{1}{n}\sum_{x_i \in H} (x'^2 - 2x'x_i + x_i^2) \\ \Rightarrow \frac{1}{n}\sum_{x_i \in H} (x - x_i)^2 \leq \frac{1}{n}\sum_{x_i \in H} (x' - x_i)^2 \\ \Rightarrow \overline{D}(x, H) \leq \overline{D}(x', H). \end{split}$$

where  $\overline{D}(x, H)$  denotes the mean distance from x to the instances in H. Therefore, the mean distance from x to the instances in H is less than the mean distance from x' to the instances in H. Since x' as well as all  $x_i \in H$  are existing class instances, according to Property 1, x must also be an existing class instance.

We deduce from the lemma that a novel class instance must be outside the decision boundary. We call any test instance outside the decision boundary as an *Foutlier*.

**Definition 4 (Foutlier).** A test instance is an Foutlier (i.e., filtered outlier) if it is outside the decision boundary of all classifiers  $L_i \in L$ , i.e., it is outside RE(L).

Again, a novel class instance must be an Foutlier.

## 4 CLASSIFICATION WITH NOVEL CLASS DETECTION

Algorithm 2 (Classify) sketches the classification and novel class detection technique. The algorithm consists of two main parts: classification (lines 1-5) and novel class

detection (lines 6-14). Details of the steps of this algorithm will be explained in the following sections.

## **Algorithm 2.** Classify $(L, x_j, buf)$

**Input:** *L*: Current ensemble of best *M* classifiers  $x_i$ : test instance

*buf*: buffer holding temporarily deferred instances

**Output:** Immediate or deferred class prediction of  $x_j$ 

1: fout  $\leftarrow$  true

- 2: if Foutlier( $L, x_j$ ) = false then
- 3:  $y'_i \leftarrow \text{majority-voting}(L, x_j) //\text{classify immediately}$
- 4: fout  $\leftarrow$  false
- 5: end if
- 6: Filter(buf)
- 7: if fout = true then
- 8:  $buf \leftarrow x_j //enqueue$

```
9: if buf.length > q and last_trial + q \le t_i then
```

10: last\_trial  $\leftarrow t_i$ 

11: novel  $\leftarrow$  DetectNovelClass(L, buf) //

- (Algorithm 3, Section 4.2)
- 12: **if** novel = **true then** remove\_novel (buf)
- 13: **end if**
- 14: end if

## 4.1 Classification

In line 2 of Algorithm 2, we first check whether the test instance  $x_j$  is an *Foutlier*. So, if  $x_j$  is not an *Foutlier*, we classify it immediately using the ensemble voting (line 3). Recall that a novel class instance must be an *Foutlier*. However, an *Foutlier* is not necessarily an existing class instance. Therefore, we perform further analysis on the *Foutliers* to determine whether they really belong to novel class.

## 4.2 Novel Class Detection

The buffer buf temporarily holds potential novel class instances. These instances are analyzed periodically in order to detect novel class, which is explained in the next paragraph. buf needs to be cleared periodically (line 6, Algorithm 2) to remove instances that no longer contribute to novel class detection. Besides, instances in buf that has reached classification deadline  $T_c$  are classified immediately. An instance is removed from *buf* if it fulfills either of the three conditions: 1) Age > S: the front of *buf* contains the oldest element in *buf*. It is removed if its age is greater than *S*, the chunk size. Therefore, at any moment in time, there can be at most S instances in buf; 2) Ensemble update: the ensemble may be updated while an instance  $x_k$  is waiting inside *buf*. As a result,  $x_k$  may no longer be an *Foutlier* for the new ensemble of models, and it must be removed if so. If  $x_k$  is no longer an *Foutlier*, and it is not removed, it could be falsely identified as a novel class instance, and also, it could interfere with other valid novel class instances, misleading the detection process; 3) Existing class: any instance is removed from buf if it has been labeled, and it belongs to one of the existing classes. If it is not removed, it will also mislead novel class detection. When an instance is removed from *buf*, it is classified immediately using the current ensemble (if not classified already).

Lines 7-14 are executed only if  $x_j$  is an *Foutlier*. At first,  $x_j$  is enqueued into *buf* (line 8). Then, we check whether *buf*.length, i.e., the size of *buf* is at least q, and the last check

on *buf* for detecting novel class had been executed (i.e., *last\_trial*) at least q time units earlier (line 9). Since novel class detection is more expensive than simple classification, this operation is performed at most once in every q time units. In line 11, Algorithm 3 (DetectNovelClass) is called, which returns true if a novel class is found. Finally, if a novel class is found, all instances that are identified as novel class are removed from *buf* (line 12).

Next, we examine Algorithm 3 to understand how *buf* is analyzed to detect presence of novel class. First, we define q-neighborhood silhouette coefficient, or q-NSC, as follows:

**Definition 5 (q-NSC).** Let  $\overline{D}_{c_{out},q}(x)$  be the mean distance from an Foutlier x to  $\lambda_{c_{out},q}(x)$  defined by (1), where  $\lambda_{c_{out},q}(x)$  is the set of q-nearest neighbors of x within the Foutlier instances. Also, let  $\overline{D}_{c_{min},q}(x)$  be the minimum among all  $\overline{D}_{c,q}(x)$ , where c is an existing class. Then, q-NSC of x is given by

$$q\text{-}NSC(x) = \frac{\bar{D}_{c_{min},q}(x) - \bar{D}_{c_{out},q}(x)}{max(\bar{D}_{c_{min},q}(x), \bar{D}_{c_{out},q}(x))},$$
(2)

where *q*-NSC, which is a unified measure of cohesion and separation, yields a value between -1 and +1. A positive value indicates that x is closer to the *Foutlier* instances (more cohesion) and farther away from existing class instances (more separation) and vice versa. Note that *q*-NSC(x) of an *Foutlier* x must be computed separately for each classifier  $L_i \in L$ . We declare a *new class* if there are at least q' (>q) *Foutliers* having positive *q*-NSC for all classifiers  $L_i \in L$ . The justification behind this decision is discussed in the next section.

## 4.2.1 Speeding up the Computation of q-NSC

For each classifier  $L_i \in L$ , computing *q*-NSC for all *Foutlier* instance takes quadratic time in the number of Foutliers. Let B = buf.length. In order to compute q-NSC for one element x in *buf*, we need O(B) time to compute the distances from x to all other elements in *buf*, and O(K) time to compute the distances from x to all existing class pseudopoints  $h \in L_i$ . Therefore, the total time to compute *q*-NSC of all elements in buf is  $O(B(B+K)) = O(B^2)$  since  $B \gg K$ . In order to make the computation faster, we create  $K_o(=(B/S) * K)$  pseudopoints from Foutliers using K-means clustering and perform the computations on the pseudopoints (referred to as *Fpseudopoints*), where S is the chunk size. The time required to apply K-means clustering on B instances is  $O(K_oB)$ . The time complexity to compute q-NSC of all of the *Fpseudopoints* is  $O(K_o * (K_o + K))$ , which is constant, since both  $K_o$  and K are independent of the input size. Therefore, the overall complexity for computing q-NSC including the overhead for clustering becomes  $O(K_o * (K_o + K) + K_o B) =$  $O(K_o(B + K_o + K) = O(K_oB)$  since  $B \gg K \ge K_o$ . So, the running time to compute *q*-NSC after speedup is linear in *B* compared to quadratic in B before speedup. q-NSC of a Fpseudopoint computated in this way is actually an approximate average of the *q*-NSC of each *Foutlier* in that Fpseudopoint. By using this approximation, although we gain speed, we also lose some precision. However, this drop in precision is negligible, as shown in the analysis to be presented shortly. This approximate q-NSC of an *Fpseudopoint* h is denoted as q-NSC'(h).

In line 1 of Algorithm 3, we create *Fpseudopoints* using the *Foutliers* as explained earlier. For each classifier  $L_i \in L$ , we compute q-*NSC*"(h) of every *Fpseudopoint* h (line 4). If the total weight of the *Fpseudopoints* having positive q-*NSC*"() is greater than q, then  $L_i$  votes for novel class (line 7). If all classifiers vote for novel class, then we decide that a novel class has really appeared (line 9). Once novel class is declared, we need to find the instances of the novel class. This is done as follows: suppose h is an *Fpseudopoint* having positive q-*NSC*"(h) with respect to all classifiers  $L_i \in L$  (note that q-*NSC*"(h) is computed with respect to each classifier separately). Therefore, all *Foutlier* instances belonging to hare identified as novel class instances.

#### Algorithm 3. DetectNovelClass(L, buf)

**Input:** *L*: Current ensemble of best *M* classifiers *buf*: buffer holding temporarily deferred instances

**Output:** true, if novel class is found; false, otherwise

- Make K<sub>o</sub> = (K\*buf.length/S) clusters with the instances in buf using K-means clustering, and create K<sub>o</sub> Fpseudopoints
- 2: Let  $\mathcal{H}_o$  be the set of *Fpseudopoints*
- 3: for each classifier  $L_i \in L$  do
- 4: **for** each  $h \in \mathcal{H}_o$  **do** Compute q-NSC'(h)
- 5:  $\mathcal{H}_p \leftarrow \{h | h \in \mathcal{H}_o \text{ and } q\text{-}NSC'(h) > 0\}$ //*Fpseudopoints* with positive q-NSC'()6:  $w(\mathcal{H}_n) \leftarrow \sum_{h \in \mathcal{H}_n} w(h), //w(h)$  is the weight of h i.e
- 6:  $w(\mathcal{H}_p) \leftarrow \sum_{h \in \mathcal{H}_p} w(h). / / w(h)$  is the weight of *h* i.e., # of instances in the *Fpseudopoint h*.
- 7: **if**  $w(\mathcal{H}_p) > q$  **then** NewClassVote++

## 8: end for

This algorithm can detect one or more novel classes concurrently as long as each novel class follows Property 1 and contains at least q instances. This is true even if the class distributions are skewed. However, if more than one such novel classes appear concurrently, our algorithm will identify the instances belonging those classes as novel, without imposing any distinction between dissimilar novel class instances (i.e., it will treat them simply as "novel"). But the distinction will be learned by our model as soon as the true labels of those novel class instances arrive, and a classifier is trained with those instances.

It should be noted that the larger the value of q, the greater the confidence with which we can decide whether a novel class has arrived. However, if q is too large, then we may also fail to detect a new class if the total number of instances belonging to the novel class is  $\leq q$ . An optimal value of q is obtained empirically (Section 5).

## 4.2.2 Impact of Evolving Class Labels on Ensemble Classification

As reader might have realized already, arrival of novel classes in the stream causes the classifiers in the ensemble to have different sets of class labels. There are two scenarios to consider. Scenario (a): suppose an older (earlier) classifier  $L_i$  in the ensemble has been trained with classes  $c_0$  and  $c_1$ , and an younger (later) classifier  $L_j$  has been trained with classes  $c_1$  and  $c_2$ , where  $c_2$  is a new class that appeared after  $L_i$  had been trained. This puts a negative effect on voting decision since the  $L_i$  obviously misclassifies instances of  $c_2$ . So, rather than counting the votes from each classifier, we selectively count their votes as follows: If an younger

classifier  $L_i$  classifies a test instance x as class c, but an older classifier  $L_i$  had not been trained with training data of  $c_i$ then the vote for  $L_i$  will be ignored if x is found to be an outlier for  $L_i$ . Scenario (b): the opposite situation may also arise where the oldest classifier is trained with some class c', but none of the newer classifiers are trained with that class. This means class c' has been outdated, and in that case, we remove  $L_i$  from the ensemble. Fig. 4a illustrates scenario (a). The classifiers in the ensemble are sorted according to their age, with  $L_1$  being the oldest and  $L_4$  being the youngest. Each classifier  $L_i$  is marked with the classes with which it has been trained. For example,  $L_1$  has been trained with classes  $c_1, c_2$ , and  $c_3$ , and so on. Note that class  $c_4$  appears only in the two youngest classifiers. *x* appears as an outlier to  $L_1$ . Therefore,  $L_1$ 's vote is not counted since x is classified as  $c_4$  by an younger classifier  $L_3$ , and  $L_1$  does not contain class  $c_4$ . Fig. 4b illustrates scenario (b). Here,  $L_1$  contains class c1, which is not contained by any younger classifiers in the ensemble. Therefore,  $c_1$  has become outdated, and  $L_1$  is removed from the ensemble. In this way, we ensure that older classifiers have less impact in the voting process. If class  $c_1$  later reappears in the stream, it will be automatically detected again as a novel class (see Definition 3).

Although there may be other techniques for updating the ensemble for handling evolving class labels, such as exponentially decaying weighting, we found our approach better than the others because of two reasons. First, uniform voting is preferred to weighted ensemble voting, which is supported by our initial observations, as also by other researchers (e.g., [10]). Second, by removing classifiers that contain outdated class labels, we make sure that if the outdated class reappears, a new classification model will be included in the ensemble. This makes the ensemble more up-to-date with the current trend of that class since the class characteristics might have been modified due to conceptdrift. Note that a new model is trained in each batch anyway (i.e., whether a novel class appears or not); therefore, there is no increase in runtime overhead due to our updating approach.

#### 4.3 Analysis and Discussion

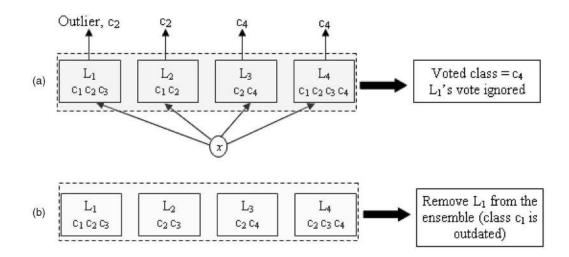
In this section, at first we justify the novel class detection algorithm, then analyze the extent of precision loss in computing *q*-NSC, and finally, analyze the time complexity of ECSMiner.

## 4.3.1 Justification of the Novel Class Detection Algorithm

In Algorithm 3, we declare a novel class if there are at least q'(>q) *Foutliers* that have positive *q*-NSC for all the classifiers in the ensemble. First, we illustrate the significance of this condition, i.e., "more than *q Foutliers* have poisitive *q*-NSC." Equation (2) deals with the mean distance between an *Foutlier* and its *nearest neighborhood*. Now, we go one step further to examine the mean distances between any pair of *Foutliers*.

Let  $\mathcal{F}$  be the set of *Foutliers* having positive *q*-NSC. Therefore, for any  $x \in \mathcal{F}$ 

$$\begin{split} \bar{D}_{c_{min},q}(x) &- \bar{D}_{c_{out},q}(x) > 0 \quad (\text{from } (2)) \\ \Rightarrow \bar{D}_{c_{min},q}(x) &> \bar{D}_{c_{out},q}(x). \end{split}$$



#### Fig. 4. Impact of evolving class label on ensemble.

Summing up for all Foutliers  $x \in \mathcal{F}$ 

$$\begin{split} \sum_{x \in \mathcal{F}} \bar{D}_{c_{min},q}(x) &> \sum_{x \in \mathcal{F}} \bar{D}_{c_{out},q}(x) \\ \Rightarrow \sum_{x \in \mathcal{F}} \frac{1}{q} \sum_{x_i \in \lambda_{c_{min},q}(x)} D(x,x_i) &> \sum_{x \in \mathcal{F}} \frac{1}{q} \\ \sum_{x_j \in \lambda_{c_{out},q}(x)} D(x,x_j) \quad (\text{from (1)}) \\ \Rightarrow \frac{1}{m} \frac{1}{q} \sum_{x \in \mathcal{F}} \sum_{x_i \in \lambda_{c_{min},q}(x)} D(x,x_i) &> \frac{1}{m} \frac{1}{q} \sum_{x \in \mathcal{F}} \\ \sum_{x_j \in \lambda_{c_{out},q}(x)} D(x,x_j) \quad (\text{letting m} = |\mathcal{F}|). \end{split}$$
(3)

Therefore, the mean pairwise distance between any pair  $(x, x_j)$  of Foutliers (such that x is an Foutlier with positive *q*-NSC and  $x_j$  is an Foutlier in  $\lambda_{C_{out},q}(x)$ ) is less than the mean pairwise distance between an Foutlier x and any existing class instance  $x_i$ . In other words, an *Foutlier* with positive q-NSC is more likely to have its k-NN within the Four instances (for k < q). So, each of the Fouriers  $x \in$  $\mathcal F$  should have the same class label as the other *Foutlier* instances and should have a different class label than any of the existing classes. This implies that the Foutliers should belong to a novel class. The higher the value of *q*, the larger the support we have in favor of the arrival of a new class. Furthermore, when all the classifiers unanimously agree on the arrival of a novel class, we have very little choice other than announcing the appearance of a novel class. The *q*-NH rule can be thought of a variation of the k-NN rule and is applicable to any data set irrespective of its data distribution and shape of classes (e.g., convex and nonconvex).

## 4.3.2 Deviation between Approximate and Exact *q*-NSC Computation

As discussed earlier, we compute *q*-NSC for each *Fpseudopoint*, rather than each *Foutler* individually in order to reduce time complexity. The resultant *q*-NSC is an approximation of the exact value. However, following analysis shows that the deviation of the approximate value from exact value is negligible.

Without loss of generality, let  $\phi_i$  be an *Fpseudopoint* having weight  $q_1$ , and  $\phi_j$  be an existing class pseudopoint having weight  $q_2$ , which is the closest existing class pseudopoint from  $\phi_i$  (Fig. 5). We compute q-*NSC*'( $\phi_i$ ), the approximate q-NSC of  $\phi_i$  using the following formula:

$$q\text{-}NSC'(\phi_i) = \frac{D(\mu_i, \mu_j) - D_i}{max(D(\mu_i, \mu_j), \bar{D}_i)},$$
(4)

where  $\mu_i$  is the centroid of  $\phi_i$ ,  $\mu_j$  is the centroid of  $\phi_j$ , and  $\overline{D}_i$  is the mean distance from centroid  $\mu_i$  to the instances in  $\phi_i$ . The exact value of *q*-NSC follows from (2):

$$q\text{-}NSC(\phi_i) = \frac{1}{q_1} \sum_{x \in \phi_i} \frac{\frac{1}{q} \sum_{x_j \in \lambda_{c_{min},q}(x)} D(x, x_j) - \frac{1}{q} \sum_{x_i \in \lambda_{c_{out},q}(x)} D(x, x_i)}{max \left(\frac{1}{q} \sum_{x_j \in \lambda_{c_{min},q}(x)} D(x, x_j), \frac{1}{q} \sum_{x_i \in \lambda_{c_{out},q}(x)} D(x, x_i)\right)}$$
(5)

where  $\lambda_{c_{out},q}(x)$  is the set of *q*-nearest neighbors of *x* within *Fpseudopoint*  $\phi_i$ , and  $\lambda_{c_{min},q}(x)$  is the set of *q*-nearest neighbors of *x* within pseudopoint  $\phi_j$ , for some  $x \in \phi_i$ . Therefore, the deviation from the exact value,  $\mathcal{E}_{qnsc} = q$ - $NSC(\phi_i) - q$ - $NSC'(\phi_i)$ . Applying (4) and (5),

$${\cal E}_{qnsc}$$

$$= \frac{1}{q_1} \sum_{x \in \phi_i} \frac{\frac{1}{q} \sum_{x_j \in \lambda_{c_{min},q}(x)} D(x, x_j) - \frac{1}{q} \sum_{x_i \in \lambda_{c_{out},q}(x)} D(x, x_i)}{max \left(\frac{1}{q} \sum_{x_j \in \lambda_{c_{min},q}(x)} D(x, x_j), \frac{1}{q} \sum_{x_i \in \lambda_{c_{out},q}(x)} D(x, x_i)\right)} - \frac{D(\mu_i, \mu_j) - \bar{D}_i}{max (D(\mu_i, \mu_j), \bar{D}_i)}.$$
(6)

In order to simplify the equations, we assume that  $q_1 = q_2 = q$ , and *q*-NSC is positive for any  $x \in \phi_i$ . Therefore,  $\lambda_{c_{out},q}(x) = \phi_i, \lambda_{c_{min},q}(x) = \phi_j$ . Also, we consider square of euclidean distance as the distance metric, i.e.,  $D(x, y) = (x - y)^2$ . Since *q*-NSC is positive for any  $x \in \phi_i$ , we can deduce following relationships:

 $R_1$ :  $max(D(\mu_i, \mu_j), \overline{D}_i) = D(\mu_i, \mu_j)$ —as the *q*-NSC for each  $x \in \phi_i$  is positive, the overall *q*-NSC of  $\phi_i$  (i.e., *q*-NSC'( $\phi_i$ )) is also positive. Therefore, this relationship follows from (4).

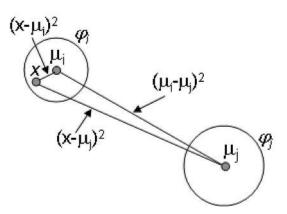


Fig. 5. Illustrating the computation of deviation.  $\phi_i$  is an *Fpseudopoint*, i.e., a cluster of *Foutliers*, and  $\phi_j$  is an existing class pseudopoint, i.e., a cluster of existing class instances. In this particular example, all instances in  $\phi_i$  belong to a novel class.

$$\begin{split} R_2: \max & \left( \frac{1}{q} \sum_{x_j \in \lambda_{c_{\min},q}(x)} D(x,x_j), \frac{1}{q} \sum_{x_i \in \lambda_{c_{out},q}(x)} D(x,x_i) \right) \\ &= \frac{1}{q} \sum_{x_j \in \lambda_{c_{\min},q}(x)} D(x,x_j), \end{split}$$

which follows, since the mean *q*-NSC of the instances in  $\phi_i$  is positive.

Also,  $\overline{D}_i = \frac{1}{q} \sum_{x \in \phi_i} (x - \mu_i)^2 = \sigma_i^2$ , the mean distance of the instances in  $\phi_i$  from the centroid. Therefore, q-NSC'( $\phi_i$ ) can be rewritten as:

$$q\text{-}NSC'(\phi_i) = \frac{(\mu_i - \mu_j)^2 - \sigma_i^2}{(\mu_i - \mu_j)^2} = \frac{1}{q} \sum_{x \in \phi_i} \frac{(\mu_i - \mu_j)^2 - (x - \mu_i)^2}{(\mu_i - \mu_j)^2}$$
$$= \frac{1}{q} \sum_{x \in \phi_i} q\text{-}NSC'(x),$$
(7)

where q-NSC'(x) is an approximate value of q-NSC(x). Now, we can deduce the following inequalities:

 $I_1: (x - \mu_i)^2 \le (\mu_i - \mu_j)^2 \text{--since } q\text{-}NSC'(x) > 0 \text{ for all } x \in \phi_i.$ 

$$I_2: \sigma_i^2 \le (\mu_i - \mu_j)^2$$
—from (7) since  $q$ - $NSC'(\phi_i) > 0$ .

$$\begin{split} I_3: & (x - \mu_j)^2 \leq (x - \mu_i)^2 + (\mu_i - \mu_j)^2 \text{--by triangle in-}\\ \text{equality (see Fig. 5).}\\ & I_4: & \sigma_j^2 \leq (\mu_i - \mu_j)^2 \text{--because } \phi_j \text{ represents an existing} \end{split}$$

 $I_4: \sigma_j^2 \leq (\mu_i - \mu_j)^2$ —because  $\phi_j$  represents an existing class, and similar inequality as  $I_2$  is applicable to the instances of  $\phi_j$ .

Continuing from (6)

$$\begin{aligned} \mathcal{E}_{qnsc} &= \frac{1}{q} \sum_{x \in \phi_i} \frac{\frac{1}{q} \sum_{x_j \in \phi_j} (x - x_j)^2 - \frac{1}{q} \sum_{x_i \in \phi_i} (x - x_i)^2}{\frac{1}{q} \sum_{x_j \in \phi_j} (x - x_j)^2} \\ &- \frac{(\mu_i - \mu_j)^2 - \sigma_i^2}{(\mu_i - \mu_j)^2} \\ &= \frac{1}{q} \sum_{x \in \phi_i} \left( \frac{\frac{1}{q} \sum_{x_j \in \phi_j} (x - x_j)^2 - \frac{1}{q} \sum_{x_i \in \phi_i} (x - x_i)^2}{\frac{1}{q} \sum_{x_j \in \phi_j} (x - x_j)^2} - \frac{(\mu_i - \mu_j)^2 - (x - \mu_i)^2}{(\mu_i - \mu_j)^2} \right). \end{aligned}$$

It is easy to show that  $\frac{1}{q}\sum_{x\in\phi_i}(x-x_i)^2 - (x-u_i)^2 = \sigma_i^2$ and  $\frac{1}{q}\sum_{x\in\phi_j}(x-x_j)^2 - (x-u_j)^2 = \sigma_j^2$ . Substituting these values, we obtain

$$\begin{split} \mathcal{E}_{qnsc} &= \frac{1}{q} \sum_{x \in \phi_i} \left( \frac{\sigma_j^2 + (x - \mu_j)^2 - \sigma_i^2 - (x - \mu_i)^2}{\sigma_j^2 + (x - \mu_j)^2} \right) \\ &\quad - \frac{(\mu_i - \mu_j)^2 - (x - \mu_i)^2}{(\mu_i - \mu_j)^2} \right) \\ &= \frac{1}{q} \sum_{x \in \phi_i} \left( 1 - \frac{\sigma_i^2 + (x - \mu_i)^2}{\sigma_j^2 + (x - \mu_j)^2} - 1 + \frac{(x - \mu_i)^2}{(\mu_i - \mu_j)^2} \right) \\ &= \frac{1}{q} \sum_{x \in \phi_i} \left( \frac{(x - \mu_i)^2}{(\mu_i - \mu_j)^2} - \frac{\sigma_i^2 + (x - \mu_i)^2}{\sigma_j^2 + (x - \mu_j)^2} \right) \\ &= \frac{\sigma_i^2}{(\mu_i - \mu_j)^2} - \frac{1}{q} \sum_{x \in \phi_i} \frac{\sigma_i^2}{\sigma_j^2 + (x - \mu_j)^2} \\ &\quad - \frac{1}{q} \sum_{x \in \phi_i} \frac{(x - \mu_i)^2}{\sigma_j^2 + (x - \mu_j)^2} \\ &\leq \frac{\sigma_i^2}{(\mu_i - \mu_j)^2} - \frac{\sigma_i^2}{\sigma_i^2 + \sigma_j^2 + (\mu_i - \mu_j)^2} \\ &\quad - \frac{1}{q} \sum_{x \in \phi_i} \frac{(x - \mu_i)^2}{\sigma_i^2 + (x - \mu_j)^2}. \end{split}$$

The last line follows since using the relationship between harmonic mean and arithmetic mean it can be shown that

$$\begin{split} &\frac{1}{q} \sum_{x \in \phi_i} \frac{\sigma_i^2}{\sigma_j^2 + (x - \mu_j)^2} \\ &\geq \frac{\sigma_i^2}{\frac{1}{q} \sum_{x \in \phi_i} (\sigma_j^2 + (x - \mu_j)^2)} = \frac{\sigma_i^2}{\sigma_j^2 + \frac{1}{q} \sum_{x \in \phi_i} (x - \mu_j)^2} \\ &= \frac{\sigma_i^2}{\sigma_j^2 + \sigma_i^2 + (\mu_i - \mu_j)^2}. \end{split}$$

Applying inequalities  $I_1$ - $I_4$ , and after several algebraic manipulations, we obtain

$$\mathcal{E}_{qnsc} \le \frac{\sigma_i^2}{(\mu_i - \mu_j)^2} - \frac{\sigma_i^2}{3(\mu_i - \mu_j)^2} - \frac{\sigma_i^2}{3(\mu_i - \mu_j)^2} = \frac{\sigma_i^2}{3(\mu_i - \mu_j)^2}.$$
(8)

Usually, if  $\phi_i$  belongs to a novel class, it is empirically observed that q- $NSC'(\phi_i) \ge 0.9$ . Putting this value in (7), and solving, we obtain  $\sigma_i^2 \le (1 - 0.9)(\mu_i - \mu_j)^2$ . Therefore, from (8), we obtain  $\mathcal{E}_{qnsc} \le 0.1/3 \approx 0.03$ . Since the range of q-NSC is -1 to +1, a deviation of 0.03 (3 percent) from the exact value is really negligible and does not effect the outcome of the algorithm. Similar reasoning can be carried out for the cases where q-NSC of the instances in  $\phi_i$  is negative.

## 4.3.3 Time and Space Complexity

Line 1 of Algorithm 3 (clustering) takes O(KS) time, and the for loop (lines 3-8) takes  $O(K^2M)$  time. The overall time complexity of Algorithm 3 is  $O(K^2M + KS) = O(KS)$  since  $S \gg KM$ . Lines 1-5 of Algorithm 2 take  $O(S(KM + Mf_c))$  per chunk, where  $f_c$  is the time to classify an instance using a classifier, and O(KM) is the time to determine whether an instance is an *Foutlier*. Line 6 takes O(S) time. Line 11 (Algorithm 3) is executed at most once in every q time units. Therefore, the worst-case complexity of lines 7-14 is O((KS) \* (S/q)), where O(KS) is the time required to execute line 11 (Algorithm 3). So, the overall complexity of Algorithm 2 is  $O(S(KM + Mf_c + KSq^{-1}))$  per chunk. For most classifiers,  $f_c = O(1)$ . Also, let S/q = m. So, the overall complexity of Algorithm 2 becomes O(KMS + MS + mS) = O(mS) since  $m \gg KM$ . Finally, the overall complexity of Algorithm 1 (ECSMiner) is  $O(mS + f_t(S))$  per chunk, where  $f_t(S)$  is the time to train a classifier with S training instances, and  $m \ll S$ .

ECSMiner keeps three buffers: buf, the training buffer  $\mathcal{L}$ , and the unlabeled data buffer U. Both buf and  $\mathcal{L}$  hold at most S instances, whereas U holds at most  $T_l$  instances. Therefore, the space required to store all three buffers is  $O(max(S, T_l))$ . The space required to store a classifier (along with the pseudopoints) is much less than S. So, the overall space complexity remains  $O(max(S, T_l))$ .

## 5 EXPERIMENTS

In this section, we describe the data sets, experimental environment, and report the results.

#### 5.1 Data Sets

## 5.1.1 Synthetic Data with only Concept-Drift (SynC)

SynC simulates only concept-drift, with no novel classes. This is done to show that concept-drift does not erroneously trigger a new-class detection in our approach. SynC data are generated with a moving hyperplane. Equation of a hyperplane is as follows:  $\sum_{i=1}^{d} a_i x_i = a_0$ . If  $\sum_{i=1}^{d} a_i x_i \leq a_0$ , then an example is negative, otherwise it is positive. Each example is a randomly generated *d*-dimensional vector  $\{x_1, \ldots, x_d\}$ , where  $x_i \in [0, 1]$ . Weights  $\{a_1, \ldots, a_d\}$  are also randomly initialized with a real number in the range [0, 1]. The value of  $a_0$  is adjusted so that roughly the same number of positive and negative examples are generated. This can be done by choosing  $a_0 = \frac{1}{2} \sum_{i=1}^{d} a_i$ . We also introduce noise randomly by switching the labels of p percent of the examples, where p = 5 is set in our experiments.

There are several parameters that simulate concept-drift. Parameter *m* specifies the percent of total dimensions whose weights are involved in changing, and it is set to 20 percent. Parameter *t* specifies the magnitude of the change in every *N* examples. In our experiments, *t* is set to 0.1, and *N* is set to 1,000.  $s_i, i \in \{1, \ldots, d\}$  specifies the direction of change for each weight. Weights change continuously, i.e.,  $a_i$  is adjusted by  $s_i.t/N$  after each example is generated. There is a possibility of 10 percent that the change would reverse direction after every N examples are generated. We generate a total of 250,000 records.

## 5.1.2 Synthetic Data with Concept-Drift and Novel-Class (SynCN)

This synthetic data simulates both concept-drift and novel class. Data points belonging to each class are generated using Gaussian distribution having different means (-5.0 to + 5.0)

and variances (0.5-6) for different classes. Besides, in order to simulate the evolving nature of data streams, the probability distributions of different classes are varied with time. This caused some classes to appear and some other classes to disappear at different times. In order to introduce conceptdrift, the mean values of a certain percentage of attributes have been shifted at a constant rate. As done in the SynC data set, this rate of change is also controlled by the parameters m, t, s, and N in a similar way. The data set is normalized so that all attribute values fall within the range [0, 1]. We generate the SynCN data set with 20 classes, 40 real valued attributes, having a total of 400K data points.

## 5.1.3 Real Data—KDDCup 99 Network Intrusion Detection

We have used the 10 percent version of the data set, which is more concentrated, hence more challenging than the full version. It contains around 490,000 instances. Here, different classes appear and disappear frequently, making the new class detection challenging. This data set contains TCP connection records extracted from LAN network traffic at MIT Lincoln Labs over a period of two weeks. Each record refers to either to a normal connection or an attack. There are 22 types of attacks, such as buffer-overflow, portsweep, guess-passwd, neptune, rootkit, smurf, spy, etc. So, there are 23 different classes of data. Most of the data points belong to the normal class. Each record consists of 42 attributes, such as connection duration, the number bytes transmitted, number of root accesses, etc. We use only the 34 continuous attributes and remove the categorical attributes. This data set is also normalized to keep the attribute values within [0, 1].

#### 5.1.4 Real Data—Forest Cover (UCI Repository)

The data set contains geospatial descriptions of different types of forests. It contains seven classes, 54 attributes, and around 581,000 instances. We normalize the data set, and arrange the data so that in any chunk at most three and at least two classes co-occur, and new classes appear randomly.

#### 5.2 Experimental Setup

We implement our algorithm in Java. The code for decision tree has been adapted from the Weka machine learning open source repository (http://www.cs.waikato.ac.nz/ml/ weka/). The experiments were run on an Intel P-IV machine with 2 GB memory and 3 GHz dual processor CPU. Our parameter settings are as follows, unless mentioned otherwise:

- 1. *K* (number of pseudopoints per classifier) = 50;
- 2. q (minimum number of instances required to declare novel class) = 50;
- 3. M (ensemble size) = 6; and
- 4. S (chunk size) = 2,000. These values of parameters are tuned to achieve an overall satisfactory performance.

#### 5.3 Baseline Method

To the best of our knowledge, there is no approach that can classify data streams and detect novel class. So, we compare ECSMiner with a combination of two baseline techniques: *OLINDDA* [23] and Weighted Classifier Ensemble (*WCE*) [27], where the former works as novel class detector and the latter performs classification. This is done as follows: For

each test instance, we delay its classification for  $T_c$  time units. That is, *OLINDDA* is given  $T_c$  time units to determine whether the instance is novel. If by that time the test instance is identified as a novel class instance, then it is considered novel and not classified using *WCE*. Otherwise, the instance is assumed to be an existing class instance, and its class is predicted using *WCE*. We use *OLINDDA* as the novelty detector since it is a recently proposed algorithm that is shown to have outperformed other novelty detection techniques in data streams [23].

However, OLINDDA assumes that there is only one "normal" class, and all other classes are "novel." So, it is not directly applicable to the multiclass novelty detection problem, where any combination of classes can be considered as the "existing" classes. Therefore, we propose two alternative solutions. First, we build parallel OLINDDA models, one for each class, which evolve simultaneously. Whenever the instances of a novel class appear, we create a new *OLINDDA* model for that class. A test instance is declared as novel, if all the existing class *models* identify this instance as novel. We will refer to this baseline method as WCE-OLINDDA\_PARALLEL. Second, we initially build an OLINDDA model using all the available classes with the first *init\_number* instances. Whenever a novel class is found, the class is absorbed into the existing OLINDDA model. Thus, only one "normal" model is maintained throughout the stream. This will be referred to as WCE-OLINDDA\_SINGLE. In all experiments, the ensemble size and chunk size are kept the same for all three baseline techniques. Besides, the same base learner is used for WCE and ECSMiner. The parameter settings for OLINDDA are following:

- 1. number of clusters built in the initial model, K = 30;
- 2. least number of normal instances needed to update the existing model = 100;
- 3. least number of instances needed to build the initial model = 100; and
- 4. maximum size of the "unknown memory" = 200.

These parameters are chosen either according to the default values used in [23] or by trial and error to get an overall satisfactory performance. *We will henceforth use the acronyms XM for ECSMiner, W-OP for WCE-OLINDDA\_PARAL-LEL, and W-OS for WCE-OLINDDA\_SINGLE.* 

## 5.4 Performance Study

#### 5.4.1 Evaluation Approach

Let  $F_n$  = total novel class instances misclassified as existing class,  $F_p$  = total existing class instances misclassified as novel class,  $F_e$  = total existing class instances misclassified (other than  $F_p$ ),  $N_c$  = total novel class instances in the stream, N = total instances the stream. We use the following performance metrics to evaluate our technique:  $\mathbf{M}_{new} = \%$ of novel class instances Misclassified as existing class =  $\frac{F_n * 100}{N_c}$ ,  $\mathbf{F}_{new} = \%$  of existing class instances Falsely identified as novel class =  $\frac{F_p * 100}{N - N_c}$ ,  $\mathbf{ERR}$  = Total misclassification error (percent) (including  $M_{new}$  and  $F_{new}$ ) =  $\frac{(F_p + F_n + F_c) * 100}{N}$ . From the definition of the error metrics, it is clear that ERR is not necessarily equal to the sum of  $M_{new}$  and  $F_{new}$ .

Evaluation is done as follows: we build the initial models in each method with the first *init\_number* instances. In our experiments, we set *init\_number* = 3S (first three chunks). From the fourth chunk onward, we evaluate the performances of each method on each data point using the time constraints. We update the models with a new chunk whenever all data points in that chunk are labeled.

## 5.4.2 Results

Figs. 6a, 6b, and 6c show the total number of novel class instances missed (i.e., misclassified as existing class) and Figs. 6d, 6e, and 6f show the overall error rates (ERR) of each of the techniques for decision tree classifier up to a certain point in the stream in different data sets. We omit SynC from the figures since it does not have any novel class. k-NN classifier also has similar results. For example, in Fig. 6a at X-axis = 100, the Y values show the total number of novel class instances missed by each approach in the first 100K data points in the stream (Forest Cover). At this point, XM misses only 15 novel class instances, whereas W-OP and W-OS misses 1,937 and 7,053 instances, respectively. Total number of novel class instances appeared in the stream by this point of time is shown by the corresponding Y value of the curve "Total," which is 12,226. Likewise, in Fig. 6d, the ERR rates are shown throughout the stream history. In this figure, at the same position (X = 100), Y values show the ERR of each of the three techniques upto the first 100K data points in the stream. The ERR rates of XM, W-OP, and W-OS at this point are following: 9.2, 14.0, and 15.5 percent, respectively.

Table 2 summarizes the error metrics for each of the techniques in each data set for decision tree and KNN. The columns headed by ERR,  $M_{new}$ , and  $F_{new}$  report the value of the corresponding metric on an entire data set. For example, while using decision tree in KDD data set, XM, W-OP, and W-OS have 1.0, 5.8, and 6.7 percent ERR, respectively. Also, their corresponding  $M_{new}$  are 1.0, 13.2, and 96.9 percent, respectively. Note that there is no novel class in SynC, and so, there is no  $M_{new}$  for any approach. Both W-OP and W-OS have some  $F_{new}$  in SynC data set, which appears since W-OP and W-OS are less sensitive to concept-drift than XM. Therefore, some existing class instances are misclassified as novel class because of concept-drift. In general, XM outperforms the baseline techniques in overall classification accuracy and novel class detection.

Figs. 7a and 7b show how XM and W-OP respond to the constraints  $T_l$  and  $T_c$  in Forest Cover data set. Similar characteristics are observed for other data sets and W-OS. From Fig. 7a, it is evident that increasing  $T_l$  increases error rates. This is because of the higher delay involved in labeling, which makes the newly trained models more outdated. Naturally,  $M_{new}$  rate decreases with increasing  $T_c$  as shown in Fig. 7b because higher values of  $T_c$  means more time to detect novel classes. As a result, ERR rates also decreases.

Figs. 8a, 8b, 8c, and 8d illustrate how the error rates of XM change for different parameter settings on Forest cover data set and decision tree classifier. These parameters have similar effects on other data sets and k-NN classifier. Fig. 8a shows the effect of chunk size on ERR,  $F_{new}$ , and  $M_{new}$  rates for default values of other parameters. We note that ERR and  $F_{new}$  rates decrease upto a certain point (2,000) then increases. The initial decrement occurs because larger chunk size means more training data for the classifiers, which leads

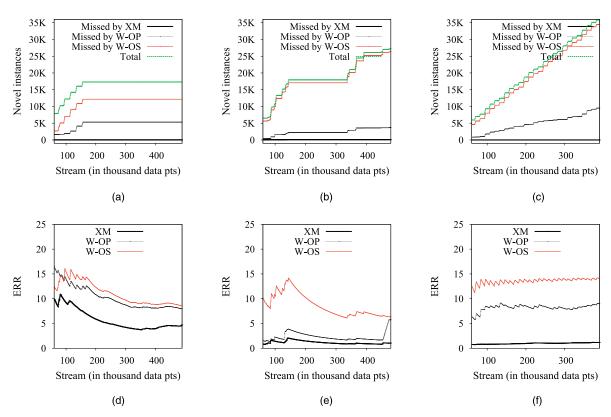


Fig. 6. (a)-(c): Novel class instances missed by each method, (d)-(f): overall error of each method ( $T_l = 1,000, T_c = 400$ ). (a) Forest cover, (b) KDDcup, (c) SynCN, (d) Forest cover, (e) KDDcup, and (f) SynCN.

to lower error rates. However, if chunk size is increased too much, then we have to wait much longer to build the next classifier. As a result, the ensemble is updated less frequently than desired, meaning, the ensemble remains outdated for longer period of time. This causes increased error rates.

Fig. 8b shows the effect of ensemble size (M) on error rates. We observe that the ERR and  $F_{new}$  rates keep decreasing with increasing M. This is because when M is increased, classification error naturally decreases because of the reduction of error variance [26]. But the rate of decrement is diminished gradually. However,  $M_{new}$  rate starts increasing after some point (M = 6) because a larger ensemble means more restriction on declaration of the arrival of novel classes. Therefore, we choose a value where the overall error (ERR) is considerably low and also  $M_{new}$  is low. Fig. 8c shows the effect of number of clusters (K) on error. The x-axis in this chart is drawn on a logarithmic scale. Although the overall error is not much sensitive on  $K, M_{new}$  rate is. Increasing K reduces  $M_{new}$  rate because outliers are more correctly detected. Fig. 8d shows the effect of q (Minimum neighborhood size to declare a novel class) on error rates. The x-axis in this chart is also drawn on a logarithmic scale. Naturally, increasing q up to a certain point (e.g., 200) helps reducing  $F_{new}$  and ERR since a higher value of q gives us a greater confidence (i.e., reduces possibility of false detection) in declaring a new class (see Section 4). But a too large value of q increases  $M_{new}$  and ERR rates (which is observed in the chart) since a novel class is missed by the algorithm if there are less than q instances of

TABLE 2 Performance Comparison

Classifier	Deternt	ERR			M new			Fnew		
Classifier	Dataset	XM	W-OP	W-OS	XM	W-OP	W-OS	XM	W-OP	W-OS
	SynC	6.9	14.1	12.8	8 <del>7</del> 8	-	-	0.0	2.4	1.1
Decision tree	SynCN	1.2	8.9	13.9	0.0	26.5	96.2	0.02	1.6	0.1
	KDD	1.0	5.8	6.7	1.0	13.2	96.9	0.9	4.3	0.03
	Forest Cover	4.7	7.9	8.5	0.2	30.7	70.1	3.0	1.1	0.2
	SynC	0.0	2.4	1.1	-	-		0.0	2.4	1.1
k-NN	SynCN	0.01	8.9	13.9	0.0	26.5	96.2	0.0	1.6	0.1
	KDD	1.2	4.9	5.2	5.9	12.9	96.5	0.9	4.4	0.03
	Forest Cover	3.6	4.1	4.6	8.4	32.0	70.1	1.3	1.1	0.2

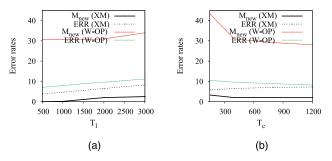


Fig. 7.  $M_{new}$  and overall error (ERR) rates on forest cover data set for (a)  $T_c = 400$  and different values of  $T_l$  and (b)  $T_l = 2,000$  and different values of  $T_c$ .

the novel class in a window of S instances. We have found that any value between 20-100 is the best choice for q.

Finally, we compare the running times of all three competing methods on each data set for decision tree in Table 3. k-NN also shows similar performances. The columns headed by "Time (sec)/1K" show the average running times (train and test) in seconds per 1,000 points, the columns headed by "Points/sec" show how many points have been processed (train and test) per second on average, and the columns headed by "speed gain" shows the ratio of the speed of XM to that of W-OP and W-OS, respectively. For example, XM is 26.9 times faster than W-OP on KDD data set. Also, XM is 1.2, 8.5, and 8.9 times faster than W-OP in SynC, SynCN, and Forest cover data sets, respectively. In general, W-OP is roughly C times slower than XM in a data set having *C* classes. This is because W-OP needs to maintain *C* parallel models, one for each class. Besides, OLINDDA model creates cluster using the "unknown memory" every time a new instance is identified as unknown, and tries to validate the clusters. As a result, the processing speed becomes diminished when novel classes occur frequently, as observed in KDD data set. However, W-OS seems to run a bit faster than XM in three data sets although W-OS shows much poorer performance in detecting novel classes and in overall error rates (see Table 2). For example, W-OS fails to detect 70 percent or more novel class instances in all data sets, but XM correctly detects 91 percent or more novel class instances in any data set. Therefore, W-OS is virtually incomparable to XM for the novel class detection task. Thus, XM outperforms W-OP both in speed and accuracy and dominates W-OS in accuracy. We also test the scalability of XM on higher dimensional data having larger number of classes. Fig. 9 shows the results. The tests are done on synthetically generated data, having different dimensions (20-60) and number of classes (10-40). Each data set has

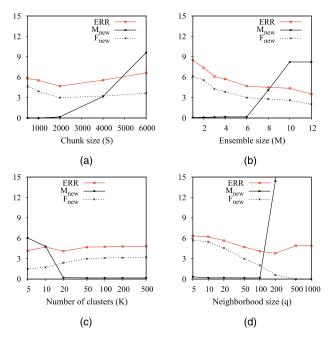


Fig. 8. Parameter sensitivity.

250,000 instances. It is evident from the results that the time complexity of XM increases linearly with total number of dimensions in the data, as well as total number of classes in the data. Therefore, XM is scalable to high-dimensional data.

## 6 DISCUSSION AND CONCLUSION

## 6.1 Discussion

We observe in the evaluation that XM outperforms both W-OP and W-OS in detecting novel classes. The main reason behind the poorer performance of W-OP and W-OS in detecting novel classes is the way OLINDDA detects novel class. OLINDDA makes two strong assumptions about a novel class and normal classes. First, it assumes a spherical boundary (or, convex shape) of the normal model. It updates the radius and centroid of the sphere periodically and declares anything outside the sphere as a novel class if there is evidence of sufficient cohesion among the instances outside the boundary. The assumption that a data class must have a convex/spherical shape is too strict to be maintained for a real-world problem. Second, it assumes that the data density of a novel class must be at least that of the normal class. If a novel class is more sparse than the normal class, the instances of that class would never be recognized as a novel class. But in a real-world problem, two different classes may have different data densities.

TABLE 3 Running Time Comparison in All Data Sets

Dataset	Time(sec)/1K				Points/se	ec	Speed gain		
Dataset	XM	W-OP	W-OS	XM	W-OP	W-OS	XM over W-OP	XM over W-OS	
SynC	0.33	0.41	0.2	2,960	2,427	5,062	1.2	0.6	
SynCN	1.7	14.2	2.3	605	71	426	8.5	1.4	
KDD	1.1	30.6	0.5	888	33	1,964	26.9	0.45	
Forest Cover	0.93	8.3	0.36	1,068	120	2,792	8.9	0.4	

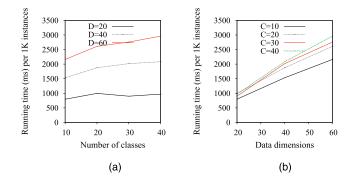


Fig. 9. Scalability test.

OLINDDA would fail in those cases where any of the assumptions are violated. On the other hand, XM does not require that an existing class must have convex shape or that the data density of a novel class should match that of the existing classes. Therefore, XM can detect novel classes much more efficiently. Besides, OLINDDA is less sensitive to concept-drift, which results in falsely declaring novel classes when drift occurs in the existing class data. On the other hand, XM correctly distinguishes between conceptdrift and concept-evolution, avoiding false detection of novel classes in the event of concept-drift. W-OS performs worse than W-OP since W-OS "assimilates" the novel classes into the normal model, making the normal model too generalized. Therefore, it considers most of the future novel classes as normal (non-novel) data, yielding very high false negative rate.

In general, existing novel class detection techniques have limited applicability since those are similar to one-class classifiers. That is, they assume that there is only one "normal" class, and all other classes are novel. However, our technique is applicable to the more realistic scenario where there are more than one existing classes in the stream. Besides, our novel class detection technique is nonparametric, and it does require any specific data distribution or does not require the classes to have convex shape. We have also shown how to effectively classify stream data under different time constraints. Our approach outperforms the state-of-the-art data-stream-based classification techniques in both classification accuracy and processing speed. We believe that our work will inspire more research toward solving real-world stream classification problems.

It might appear to readers that in order to detect novel classes we are in fact examining whether new clusters are being formed, and therefore, the detection process could go on without supervision. But supervision is necessary for classification. Without external supervision, two separate clusters could be regarded as two different classes although they are not. Conversely, if more than one novel classes appear simultaneously, all of them could be regarded as a single novel class if the labels of those instances are never revealed.

### 6.2 Conclusion

We have addressed several real-world problems related to data stream classification. We have proposed a solution to the concept-evolution problem, which has been ignored by most of the existing data stream classification techniques. Existing data stream classification techniques assume that total number of classes in the stream is fixed. Therefore, instances belonging to a novel class are misclassified by the existing techniques. We show how to detect novel classes automatically even when the classification model is not trained with the novel class instances. Novel class detection becomes more challenging in the presence of concept-drift.

In future we would like to apply our technique to network traffic. Besides, we would like to address the data stream classification problem under dynamic feature sets.

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