

Online Streaming Feature Selection via Multi-Conditional Independence and Mutual Information Entropy[†]

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ABSTRACT

The goals of feature selection are to remove redundant and irrelevant features from high-dimensional data, extract the “optimal feature subset” of the original feature space to improve the classification accuracy, and reduce the time complexity. Traditional feature selection algorithms are based on static feature spaces that are difficult to apply in dynamic streaming data environments. Existing works, such as *Alpha-investing* and Online Streaming Feature Selection (OSFS), and Scalable and Accurate OnLine Approach (SAOLA), have been proposed to serve the feature selection with streaming feature, but they have drawbacks, including low prediction accuracy and a large number of selected features if the streaming features exhibit characteristics such as low redundancy and high relevance. To address the limitations of the abovementioned works, we propose the algorithm of Online Streaming Feature Selection via Conditional dependence and Mutual information (OSFSCM) for streaming feature, which is found to be superior to *Alpha-investing* and OSFS for datasets with low redundancy and high relevance. The efficiency of the proposed OSFSCM algorithm is validated through a performance test on widely used datasets, e.g., *NIPS 2003* and *Causality Workbench*. Through extensive experimental results, we demonstrate that OSFSCM significantly improves the prediction accuracy and requires fewer selected features compared with *Alpha-investing* and OSFS.

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1. INTRODUCTION

Dimensionality reduction can combat the curse of dimensionality [1–4]. Feature selection chooses an “optimal feature subset” by following a certain criterion while removing redundant and irrelevant features of the class attribute during classification from the original high-dimensional feature set [2,5–7]. This process maintains relevant features for learning and provides faster and more cost-effective predictors with respect to efficiency and effectiveness [8].

In real-world applications, many datasets have very high-dimensional and increasing feature spaces [9,10], such as the datasets from the real-time monitoring and analysis of environments [11], capturing CCTV streaming images [12], collecting features for intrusion detection [13,14], real-time updates of popular topics in *Microblog* and *Twitter* [15,16], and real-time spam filtering of email [17]. In such streaming feature spaces, the new, individual features flow into the model dynamically and must be considered sequentially. As an emerging research direction during the past few years, online feature selection with streaming features

offers a complementary algorithmic methodology to enrich online feature selection [9,18,19] that addresses high-dimensionality in big data analyses. This methodology assumes that sequential features flow into the model dynamically, and the feature selection is performed when features arrive. This approach is different from classical online learning in which the feature space remains consistent while samples flow in sequentially [19,20]. The representative classical methods [19] include *Grafting*, *Alpha-investing*, online streaming feature selection (OSFS), Online Feature Selection with Group Structure (OGFS), and scalable and accurate online approach (SAOLA).

Feature sets with a continuous flow of new features, such as in medical diagnosis, relevant features might be generated with low redundancy. This scenario results in many selected features being generated in the candidate feature set. Experiments from the previously mentioned studies demonstrate that feature selection is restricted in these datasets. For example, the *Grafting* algorithm is weak at handling streaming features because the entire feature space must be known. The algorithm of *Alpha-investing* has low prediction accuracy and is unstable. Although the OSFS algorithm offers high prediction accuracy for these types of datasets, the running time increases exponentially with increasing of selected features, so OSFS becomes very slow. The SAOLA algorithm offers good efficiency in its running time while having a few features, but its prediction accuracy is not excellent.

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Motivated by the above analysis of existing studies of online feature selection with streaming features, we investigate the problem with a novel algorithm called Online Streaming Feature Selection via Conditional dependence and Mutual (OSFSCM) information to solve the following challenges: (1) how to select the feature subset in a streaming feature space with low redundancy and obtain high relevancy, (2) how to provide effective continuous mechanisms to discover the pattern of the running time among all generated features with increasing feature volumes, and (3) how to validate the performance.

The contributions of this paper that extend beyond the existing methods include our proposal of a novel algorithm for streaming feature selection, an analysis of its theoretical properties, and a validation of its empirical performance through extensive experiments.

The remainder of the paper is organized as follows: Section 2 surveys the related work and Section 3 introduces notations, definitions, and a framework for streaming feature filtering as well as the proposal of our OSFSCM algorithm and analysis. Section 4 reports our experimental results and describes its application in a real scenario, followed by our conclusion in Section 5.

2. RELATED WORK

Feature selection is simple, interpretable, and essential when working with high-dimensional data [18,19]. Based on the available label information, feature selection can include supervised, unsupervised, and semi-supervised methods. From the perspective of a selection strategy, these algorithms can include a wrapper, filter, or embedded method [19].

Online feature selection for streaming features is an important branch of feature selection [4,8,21] that focuses on scenarios where the number of features changes over time, which must be processed in real-time and not after all the features arrive. Representative algorithms of this field include *Grafting*, *Alpha-investing*, *OSFS*, *OGFS*, and *SAOLA*.

Grafting [22] is an embedded feature selection method applied to streaming features with irrelevant and redundant features. However, *Grafting* is weak for this approach because it must know about the entire feature space to select a good value for its regularization parameter. *Alpha-investing* [23] can handle infinitely large feature sets but also must transform the initial features based on prior knowledge of candidate features. Only judging the relevance of features can give many redundant features in the candidate set resulting in a reduction of classification accuracy. *OSFS* [24] uses the G^2 test to denote conditional independence or dependence between features from which it identifies irrelevant and redundant features. While *OSFS* can remove these irrelevant and redundant features from streaming features, when the number of weak relevant features increases, the running time increases exponentially. *OGFS* [25] leverages the group structure information as a priori knowledge for feature selection and generates a better feature subset through two stages of intra-group and inter-group feature selection. However, *OGFS* must select a few parameters in advance, which is relatively difficult without prior information. *SAOLA* [15] can handle an extremely high-dimensional feature space by using a strategy of online pairwise comparisons. However, *SAOLA* only judges if the features are redundant under a single condition and cannot remove

all redundant features. Besides, the algorithm alone cannot obtain the judgment threshold of the optimal relevance between features.

Based on the above observations, we propose an efficient OSFSCM algorithm for feature selection in a streaming feature space that offers low redundancy and high relevance.

3. THE OSFSCM ALGORITHM FOR ONLINE FEATURE SELECTION

3.1. Notations and Definitions

The selected features of the class attribute represent an optimal feature subset [24,26] containing all strongly relevant and nonredundant features [24], as shown in Figure 1. The optimal feature subset is named the Markov blanket of the class attribute, and this criterion only removes the attributes that are unnecessary for inclusion into the feature set because they are irrelevant or redundant [7,24,27]. We consider this problem in the context of online feature selection with streaming features where S is the feature space set containing all available features.

Table 1 lists all symbols and notations used in this paper. Assume that x_i denotes the i th new incoming feature at time t_i , S_{i-1} is the selected feature set at time t_{i-1} , $S_i \subseteq S$, and T is the class attribute.

Definition 1 (Independence [28]). In a set S , two variables $x, y \in S$ are independent with respect to a probability distribution P , denoted as $x \perp y$, iff $P(x, y) = P(x)P(y)$. Similarly, x and y are not independent, denoted as $x \not\perp y$, iff $P(x, y) \neq P(x)P(y)$.

Definition 2 (Conditional Independence [8]). In a set S , two random variables $x, y \in S$ are conditionally independent given a set of variables $S_i \subseteq S - \{x, y\}$ with respect to a probability distribution P iff there exists an assignment of x and y , such that $P(x | y, S_i) = P(x | S_i)$, denoted as $x \perp y | S_i$.

Conditional independence is a generalization of the traditional notion of statistical independence in Bayesian networks due to the

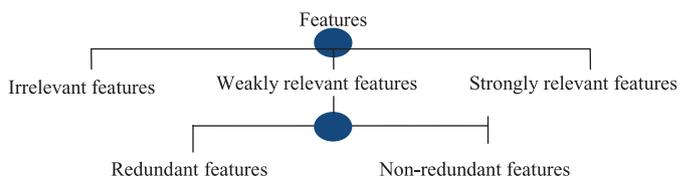


Figure 1 | Feature relevance and redundancy [24].

Table 1 | Notations with mathematical meanings.

Notation	Mathematical Meanings
S	Feature space set
f	Feature $f, f \in S$
CFS	Candidate feature set at current time
t	A time point
T	Class attribute
$P(x)$	Event probability of feature x
$P(. .)$	Conditional probability
ρ	A threshold
$MB(T)$	Markov blanket of T
$a \perp b$	a is independent of b

factorizations of the allowed joint probability distribution. If x and y are dependent in the condition of S_i , then we write $P(x, y | S_i) = P(x | S_i)P(y | S_i)$.

Definition 3 (Markov blanket [15,29,30], MB). A Markov blanket of the class attribute T , denoted as $MB(T)$, is a minimal set of features that makes $\forall x \in S - MB(T) \cup \{T\}$, such that $x \perp T | MB(T)$.

$MB(T)$ is the set of parents, children, and children's parents of T [31]. These Markov blankets can eliminate a conditionally independent feature without increasing the distance from the desired distribution. The Markov blanket criterion only removes the attributes that are unnecessary based on those that are irrelevant to the class attribute and redundant given other attributes [26].

Definition 4 (Strong relevance [24]). A feature x is strongly relevant to the class attribute, T , iff $\forall S_i \subseteq S - \{x\}$, such that $P(T | S_i, x) \neq P(T | S_i)$.

Definition 5 (Weak relevance [24]). A feature x is weakly relevant to the class attribute, T , iff $\exists S_i \subseteq S - \{x\}$, such that $P(T | x, S_i) \neq P(T | S_i)$.

Definition 6 (Irrelevance [24]). A feature x is irrelevant to a class attribute, T , iff $\forall S_i \subseteq S - \{x\}$, s.t. $P(T | S_i, x) = P(T | S_i)$.

Definition 7 (Redundant features [24]). A feature x is redundant to the class attribute T iff it is weakly relevant to T and has a $MB(x)$ that is a subset of $MB(T)$.

3.2. Using Mutual Information to Filter Partial Redundant Features

Mutual information theory and Fisher's z-test are used to judge the correlation between features [15]. The evaluation standard of entropy information quantity, also known as Shannon entropy, uses a numerical value to express the uncertainty degree of a random variable. The entropy of feature Y is defined as

$$H(Y) = - \sum_{y_i \in Y} P(y_i) \log_2 P(y_i) \quad (1)$$

Conditional entropy refers to the uncertainty degree of another variable after it is known, i.e., the dependence degree of the variable on the known variable. Assuming the feature Z is known, the conditional entropy of Y under the Z condition is defined as

$$H(Y|Z) = \sum_{z_i \in Z} P(z_i) \sum_{y_i \in Y} P(y_i | z_i) \log_2 P(y_i | z_i) \quad (2)$$

Conditional mutual information is the degree of correlation between two features under the condition that a certain feature is known. The expression of conditional mutual information is established according to Eqs. (3) and (4) [15].

$$\begin{aligned} I(X; Y|Z) &= H(X|Z) - H(X|YZ) \\ &= H(X|Z) + H(Y|Z) - H(Z, Y, Z) - H(Z) \end{aligned} \quad (3)$$

When a new feature f arrives, to determine if it is relevant for the class attribute T , the algorithm calculates a correlation threshold δ , such that if $I(f; T) > \delta$ ($0 \leq \delta \leq 1$), and $f_i \in CFS - \{f\}$, then f is considered a relevant feature for the class attribute T . Meanwhile, if

Eq. (4) is hold [15], the f term is considered a redundant feature, so it is then filtered and not considered in the next process.

$$I(f; T) \geq I(f_i; T) \text{ and } I(f_i; f) \geq I(f_i; T) \quad (4)$$

3.3. Using Fisher's z-Test to Filter Irrelevant and Partially Redundant Features with Continuous Data

Fisher's z-test calculates the correlation degree between features [15], as in shown in Eq. (5). In the Gaussian distribution $N(\mu, \Sigma)$, after the feature subset S is provided, the expression of the partial correlation coefficient $r_{(f_i, T|S)}$ between the feature f_i and the class attribute T is expressed as [15].

$$r_{(f_i, T|S)} = \frac{r_{(f_i, T)} - r_{(f_i, S)}r_{(TS)}}{\sqrt{1 - r_{(f_i, S)}^2} \sqrt{1 - r_{(TS)}^2}} \quad (5)$$

In Fisher's z-test, under the null hypothesis of the conditional independence between the feature f_i and the class attribute T of the given feature subset S , $r_{(f_i, T|S)} = 0$. Suppose α is a given significance level, and ρ is a p -value returned by Fisher's z-test. Under the null hypothesis of the conditional independence of f_i and T , if $\rho > \alpha$, f_i and T are not related to each other when the subset S is given. If $\rho \leq \alpha$, then f_i and T are relevant to each other.

3.4. Using the G² Test to Filter Irrelevant and Partially Redundant Features with Discrete Data

As an alternative to the X^2 test, the G^2 test is a statistic defined as Eq. (6) [24]

$$G^2 = 2 \sum_{i,j,k} S_{abc}^{ijk} \ln \frac{S_{abc}^{ijk} S_c^{ik}}{S_{ac}^{ik} S_{bc}^{jk}} \quad (6)$$

where S_{abc}^{ijk} represents the number of features satisfying $f_i = a, f_j = b$, and $f_k = c$ in a dataset, and $S_{ac}^{ik}, S_{bc}^{jk}, S_c^{ik}$ are defined similarly. If three features f_i, f_j , and f_k , and f_i, f_j , are conditionally independent given f_k . Under the condition where α has a given significance level of 0.05, then ρ is the p -value returned and $f_i \perp f_j | f_k$ defines the null hypothesis (H_0) of f_i , and f_j is conditionally independent given f_k iff $\rho > \alpha$. Otherwise, for (H_1), f_i and f_j are nonconditionally independent given f_k iff $\rho \leq \alpha$, i.e. $f_i \not\perp f_j | f_k$.

3.5. The OSFSCM Algorithm and Analysis

In this section, we propose a new approach for feature filtering via multi-conditional independence and mutual information entropy to process data with streaming features. In this approach, the data stream is fixed, while the features continue arriving as each feature is evaluated. The process of feature selection can be performed in three phase, as shown in Table 2. First, the irrelevant features are filtered by the nonconditional independence and leaving the

relevant features. Second, part of the redundant features is further discarded from the weakly relevant features by filtering through mutual information. Finally, the remaining redundant features are discarded by filtering through conditional independence.

According to the above approach, The *OSFSCM* algorithm for streaming feature filtering is proposed, as is shown in Algorithm 1.

Algorithm 1: The OSFSCM algorithm

1. Initialization
 - (1) Selected features set $CFS = \{\}$, class attribute T
 2. Analysis of irrelevant features: filtering via non-conditional independence
 - (2) Get a new feature f ;
 - (3) If f is an irrelevant feature, discard f , enter step (2);
 - (4) If not, $CFS = CFS \cup \{f\}$, enter step 3;
 3. Analysis of redundant features 1: the filtering via mutual information
 - (5) If $\exists x \in CFS$, x is the redundant feature in the filtering via mutual information, discard x , and enter step (2); Once all features in CFS is non-redundant according to Eq. (4); then, step (6);
 4. Analysis of redundant features 2: the filtering via conditional independence
 - (6) If $\exists y \in CFS$, $subSet \subseteq CFS - \{y\}$, f is the redundant feature according to Eq. (5) or Eq. (6), discard y ;
 5. Repeat steps 2–5 until no new features or stopping criteria is met;
 6. Output the selected features CFS .
-

In the *OSFSCM* algorithm, CFS is a candidate feature set at the current time, and f is the new feature. In step 2, nonconditional independence filtering is executed. If it returns an irrelevant feature of class attribute T , then the feature f is discarded. Otherwise, the feature f goes into further analysis of the redundant features.

The filtering of the redundant features is divided into two sequential steps. Step 3 includes the filtering via mutual information and step 4 filters via conditional independence.

For filtering via mutual information, according to Eq. (4), a new feature f is relevant with T , $\exists x \in CFS - \{f\}$, if $I(x; T) \geq I(f; T)$ and $I(f; x) \geq I(f; T)$, then x can be removed from CFS until any feature in $CFS - \{f\}$ is nonredundant in the condition of f .

For filtering via conditional independence, the candidate feature set CFS includes the new feature f , and the expression of $y \perp T \mid subSet$ is determined on the condition of $subSet \subseteq CFS - \{y\}$. If $y \perp T \mid subSet$, then the feature y is redundant, and y is discarded from CFS . Through continuous looping of steps 3–4, all redundant features in CFS are discarded due to the inclusion of new features.

The *OSFSCM* uses the notation $x \perp T \mid S$, $S \subseteq CFS - \{x\}$ to denote the conditional independence. To measure $x \perp T \mid S$, *OSFSCM* uses the p -value returned by the G^2 test for discrete data and Fisher's

Table 2 | Phase of filtering for streaming features.

Phase of Filtering	Feature Analysis
Phase1: filtering via nonconditional independence	Analysis of irrelevant features
Phase2: filtering via mutual information	Analysis of redundant features
Phase3: filtering via conditional independence	

z -test for continuous data of its measurements with significance levels of 0.05 or 0.01, where the former threshold is used in this paper.

3.6. The Time Complexity of OSFSCM

The complexity of the *OSFSCM* algorithm depends on the tests of nonconditional independence filtering, mutual information filtering, and conditional independence filtering. Assume that $|N|$ is the number of features arrived, and $|N|$ also is the number of remaining features before filtering via non-conditional independence, $|S_1|$ is the number of remaining features before filtering via mutual information, and $|CFS|$ is the number of remaining features before filtering via conditional independence. For *OSFSCM*, we use the k -greedy search strategy with $k = 3$. The complexity of each filtering phase is shown in Table 3. Therefore, the complexity is represented as $O(|N| |S_1| |CFS|^2 C_{|CFS|}^3)$. Therefore, the search times exponentially decline. Meanwhile, numerous irrelevant and redundant features exist in the large data condition, so the size of $|CFS|$ is very small. The time complexity determined by the number of features within CFS and the worst-case complexity is $O(|N|^4 C_{|N|}^3)$ when the size of $|S_1|$ is $|N|$ during step (5) and $|CFS|$ is $|N|$ during step (6). With new features continuously arriving, the complexity of *OSFSCM* becomes very high. This means that almost all features is strongly related with class attributes are, and there are almost no irrelevant features and redundant features in the dataset. Few features are filtered out by steps 2 to 5. Obviously, this situation nearly does not exist in real flow datasets.

4. EXPERIMENTAL RESULTS

4.1. Experiment Setup

We empirically evaluate the performance of the *OSFSCM* algorithm on 14 benchmark datasets listed in Table 4. All experiments are performed on a computer with an Intel(R) Xeon(R) CPU E3-1505M 3.0 GHz with 32 G RAM.

The arcene, colon, ionosphere, and leukemia datasets is sourced from the *NIPS 2003* feature selection challenge [8] and a frequently studied public microarray datasets (wdbc). We also downloaded the datasets from the *Causality Workbench* that includes sylvia, lung, cina0, reged1, lucas0, marti1, and lucap0. The cina0 is a marketing dataset derived from census data. The reged1 is a genomics dataset for studying the causes of lung cancer. The marti1 is obtained from the data generative process of simulated genomic data. The lucas0 is a lung cancer simple set, and lucap0 is a lung cancer set with probes, which are used for modeling a medical application for the

Table 3 | The time complexity of OSFSCM algorithm.

Phase of filtering	Cost
Phase1: filtering via nonconditional independence	$O(N)$
Phase2: filtering via mutual information	$O(S_1 CFS)$
Phase3: filtering via conditional independence	$O\left(C_{ CFS }^3 \cdot CFS \right)$

OSFSCM, online streaming feature selection via conditional dependence and mutual information.

diagnosis, prevention, and cure of lung cancer. The number of features available in these datasets ranges from 11 to 10000, and the number of samples varies from 72 to 145252. In particular, the number of features of seven datasets is larger than the number of samples, including mart1, reged1, lung, prostate_GE, leukemia, arcene, and Smk_can_187. These 14 datasets cover a wide range of real-world application domains, including gene expressions, ecology, and casual discovery, making the construction of feature selection very challenging. We preprocess the data by turning to a standard score and deleting the same columns, such as in leukemia and other datasets.

Our analysis of the *OSFSCM* algorithm compares against the state-of-the-art online feature selection algorithms, *Alpha-investing*, and *OSFS*, using 10-fold cross-validation on each training dataset. First, we compared the prediction accuracy of *OSFSCM* with that of the state-of-the-art using 12 classifiers, including Decision Tree, KNN, SVM, and Ensemble, as implemented in MATLAB. Second, we analyzed the number of selected features and the running time for each

algorithm. Third, the *OSFSCM* algorithm is applied to a real-world data scenario and compared with the other algorithms.

4.2. Comparison of *OSFSCM* with Two Online Algorithms

The algorithms are implemented in Library of Online Streaming Feature Selection (LOFS) [18], an open-source library available in MATLAB 2017. To evaluate the selected features in the experiments, we use twelve available classifiers in the app of “classification learner” in MATLAB 2017, including Decision Trees (Complex Tree, Medium Tree, and Simple Tree), SVM (Linear, Quadratic, and Cubic), KNN (Fine, Medium, and Cubic), and Ensemble (Bagged Trees, Subspace discriminant, and RUSBoosted Trees) classifiers. In the app of classification learner, we automatically train these classification models with default parameters, as is shown in the Table 5. We choose the results of *OSFSCM*, *OSFS*, and *Alpha-investing* as the datasets to train and validate classification

Table 4 | The benchmark datasets used for algorithm evaluation.

Dataset	#Features	#Instances	Dataset	#Features	#Instances
wdbc	30	569	mart1	1024	500
colon	2000	62	reged1	999	500
lucas0	11	2000	lung	3312	203
sylva	216	13086	prostate_GE	5966	102
ionosphere	34	351	leukemia	7066	72
cina0	132	16033	arcene	10000	100
lucap0	143	2000	Smk_can_187	19993	187

#Features: the number of features; #Instances: the number of instances.

Table 5 | Parameter settings in 12 classifiers.

Classifier Parameters	Parameter Values		
	Complex Tree	Medium Tree	Simple Tree
Maximum number of splits	100	20	10
Split criterion		Gini's diversity index	
Surrogate decision splits		Off	
Maximum surrogates per node		10	
	Linear SVM	Quadratic SVM	Cubic SVM
Kernel function	Linear	Quadratic	Cubic
Box constraint level		1	
Kernel scale mode		Auto	
Manual kernel scale		1	
	Fine KNN	Medium KNN	Cubic KNN
Number of neighbors	1	10	10
Distance metric	Euclidean	Euclidean	Minkowski
Distance weight		Equal	
Box constraint level		1	
Kernel scale mode		Auto	
Manual kernel scale		1	
	Bagged Trees	Subspace Discriminant	RUSBoosted Trees
Ensemble method	Bag	Subspace	RUSBoost
Learner type	Decision tree	Discriminant	RUSBoost
Maximum number of splits	266	20	20
Number of learners		30	
Learning rate		0.1	
Subspace dimension		1	

models. After training multiple models, compare these algorithms' performance side-by-side in these classifiers.

As described above, we evaluate *OSFSCM* against the others based on prediction accuracy, the number of selected features, and the running time. In the following, we perform statistical comparisons to analyze the prediction accuracies further.

4.2.1. Prediction Accuracy

Figure 2 summarizes the prediction accuracies with 12 classifiers applied to 14 datasets during online learning. We conduct these tests using G^2 for discrete data and Fisher's z-test for continuous data at an $\alpha = 0.05$ significance level. The prediction accuracies of *OSFSCM* and *OSFS* are higher compared with *Alpha-investing* for most of the 5-14 datasets for these classifiers. *OSFSCM* achieves a higher accuracy for all the classifiers except for *RUSBoosted Trees*. With some datasets, as seen in Figure 2, the accuracies of the classifiers are reduced too much, such as in leukemia, marti1, and reged1.

In addition, from the three curves in Figure 2, the prediction accuracies from *Alpha-investing*, *OSFS*, and *OSFSCM* are nearly the same for some datasets, including six datasets using Complex Tree, three each using Medium Tree, Liner SVM, Quadratic SVM, Cubic SVM, Fine KNN, Medium KNN, and Subspace discriminant, two datasets with Cubic KNN, Bagged Trees, and *RUSBoosted Trees*,

and one dataset with Simple Tree. For the *wdbc*, *lucas0*, and *sylva* datasets, the prediction accuracies of these classifiers, as well as *RUSBoosted Trees*, remain nearly the same.

4.2.2. Number of Selected Features and Running Time

To further examine these three algorithms, Table 6 lists the performance of each with respect to the 14 datasets comprised of different numbers of features.

- *Summary of the numbers of selected features*
While the prediction accuracy of *OSFSCM* is higher than *Alpha-investing* and *OSFS* for most datasets, as previously described in Figure 2, another observation is apparent in Table 6. The number of selected features from *OSFSCM* is more than *Alpha-investing* and *OSFS* for many datasets. This result is due to the following reasons:

For the *Alpha-investing* algorithm: From its low prediction accuracy, the ability of the algorithm to mine features is low, so part of the Markov blanket elements cannot be obtained.

For the *OSFS* algorithm: During the redundant feature analysis phase, the non-redundant features could be discarded under the condition of redundant features due to its low prediction accuracy, so fewer features are selected.

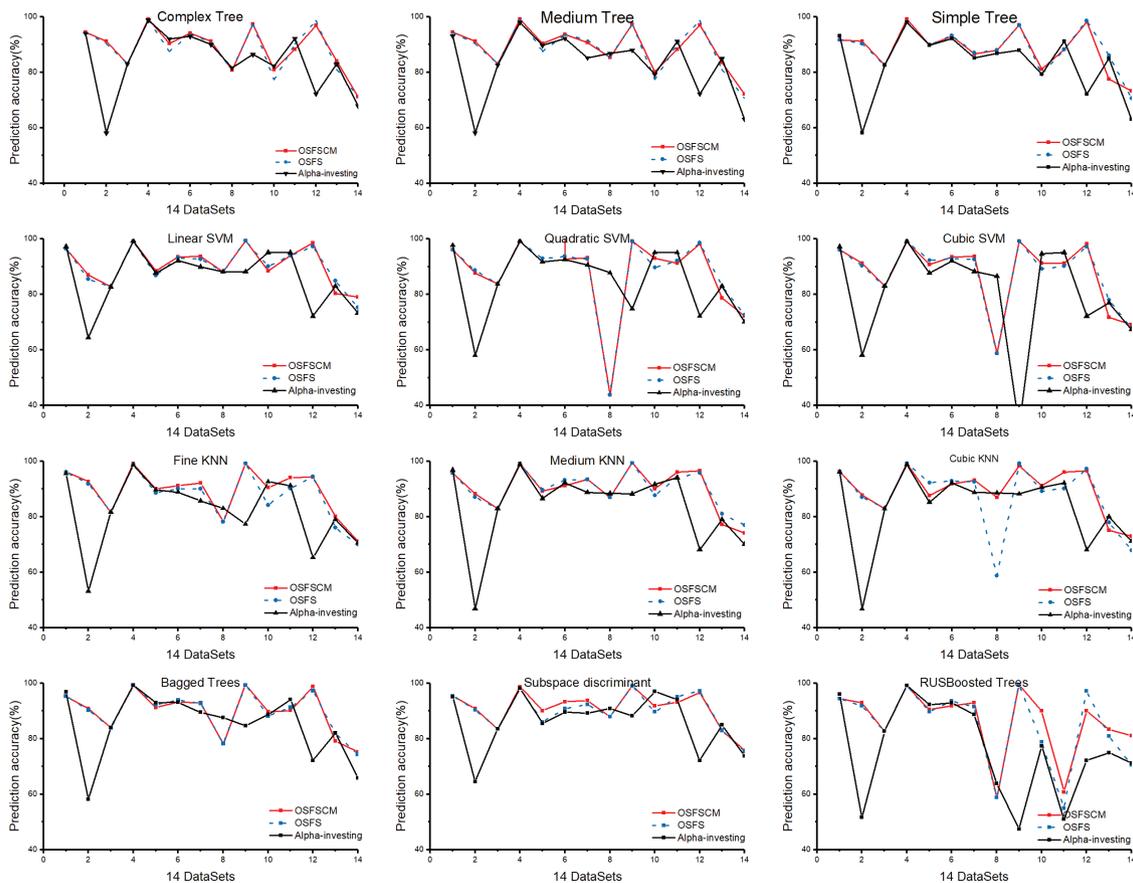


Figure 2 | Prediction accuracies of the three algorithms on 14 datasets using classifiers on the selected features. The x-axis labels represent the 14 datasets as (1) *wdbc*, (2) *colon*, (3) *lucas0*, (4) *sylva*, (5) *ionosphere*, (6) *cina0*, (7) *lucap0*, (8) *marti1*, (9) *reged1*, (10) *lung*, (11) *prostate_GE*, (12) *leukemia*, (13) *arcene*, and (14) *Smk_can_187*.

For the *OSFSCM* algorithm: A large number of selected features is potential due to its significant outperformance over *OSFS* and *Alpha-investing* in mining the elements from the Markov blanket, enabling it to find more elements.

- *Summary of the running times*

The running time performances for the three algorithms are also reported in Table 6. The *Alpha-investing* is faster than *OSFS* and *OSFSCM* for all datasets because it only considers the newest feature added, and never considered the discarded features again. This approach also results in low prediction accuracy (as seen in Figure 2) with the classifiers, as with the datasets colon, lucap0, reged1, leukemia, and Smk_can_187.

The performances of the running times in *OSFSCM* and *OSFS* are very different, as seen in the results in Table 6, with *OSFS* being much faster than *OSFSCM* on the datasets of wdcb, colon, lucas0, ionosphere, lung, prostate_GE, arcene, leukemia, and Smk_can_187.

On the other hand, *OSFSCM* is much faster compared with *OSFS* on the datasets of sylvia, cina0, lucap0, reged1, and lung, as highlighted in bold. These datasets include features with higher feature selection, as shown in Table 4, and have low redundancy with high strong and weak relevance because the number of candidate features significantly influences the runtime for the two algorithms.

4.3. Application in a Real Scenario

The PEMS-SF dataset from the UCI website is selected for our real scenario test for algorithm evaluation. This dataset contains 440 instances and 138672 features based on 963 sensors recordings between January 1, 2008, and September 30, 2009, of daily lane occupancy rates on highways. Each feature represents a lane occupancy rate from a sensor for a day (between 0 and 1) with a classification label of 1 through 7, representing Monday to Sunday. The dataset deleted data on public holidays and anomalies observed on March 8 and March 9, 2008. This evaluation only considers the *OSFSCM*, *OSFS*, and *SAOLA* algorithms because the *Alpha-investing* algorithm cannot run in the PEMS-SF dataset. The significance level α is set to 0.05, and the experiment performs multiple 10-fold cross-validations.

Table 7 reports the running time of the *OSFSCM* algorithm on the PEMS-SF dataset. As the features flow in, *OSFSCM* removes many irrelevant and redundant features. First, irrelevant features can be removed by the filtering of nonconditional independence, followed by the removal of redundant features by the filtering of mutual information and conditional independence. Finally, the approximate Markov blanket of the classification label is obtained by the three filters.

Compared with the low prediction accuracy of *Alpha-investing* and a high running time cost (greater than three days) of *OSFS* on the

Table 6 | Number of selected features and running time.

Dataset	Alpha-Investing		OSFS		OSFSCM	
	#Features	Time(s)	# Features	Time(s)	# Features	Time(s)
wdbc	20	0.0138	3	0.1577	3	0.0567
colon	1	0.0663	3	0.6778	4	0.0582
lucas0	4	0.0008	4	0.0142	4	0.0061
sylvia	70	1.6717	18	247.9366	22	32.5611
ionosphere	10	0.0147	4	0.1315	7	0.0669
cina0	8	0.1046	22	721.3638	28	62.1947
lucap0	10	0.0197	36	1.43E+07	41	72.5523
marti1	28	0.116	1	0.1081	1	0.1045
reged1	1	0.0417	13	121.2839	13	27.3695
lung	45	0.7523	11	420.5678	20	85.3698
prostate_GE	12	0.4308	3	7.7915	4	3.1189
leukemia	1	0.4346	3	12.7647	5	8.3657
arcene	8	1.4139	5	20.8445	6	5.8624
Smk_can_187	6	2.7929	4	42.8323	7	12.1648

OSFS, online streaming feature selection; OSFSCM, online streaming feature selection via conditional dependence and mutual information.

Table 7 | Running time and the number of selected features from *OSFSCM* algorithm.

Algorithm	Time(s)	#Features			
		Ratio (%)	% Filtering via Nonconditional Independence	Filtering via Mutual Information	Filtering via Conditional Independence
<i>OSFSCM</i>	476.3568	25	14295	98	21
		50	31461	110	33
		75	46636	150	35
		100	61618	210	41

OSFSCM, online streaming feature selection via conditional dependence and mutual information.

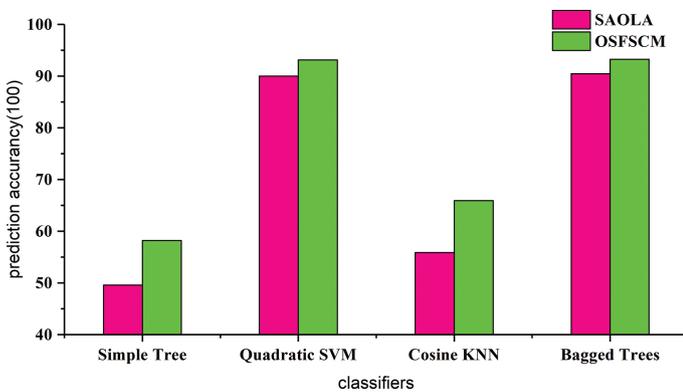


Figure 3 | Classification accuracies on the PEMS-SF dataset using four classifiers.

PEMS-SF dataset, *OSFSCM* and *SAOLA* complete the dataset filtering with many instances and features in a limited time. As is shown in Figure 3, the classification accuracies of *OSFSCM* on Decision Tree (simple tree), SVM (qualitative SVM), KNN (cosine KNN), and embedded (bagged trees) are higher compared with *SAOLA*, with differences of 8.63%, 3.11%, 10.03%, and 2.81%, respectively.

5. CONCLUSIONS

We explored OSFS through a proposed algorithm called *OSFSCM* designed with conditional dependence and mutual information. Using benchmark datasets, we compared the *OSFSCM* algorithm with two state-of-the-art online feature selection methods, and our empirical results demonstrated the following: (1) The prediction accuracy of *OSFSCM* is higher compared with *Alpha-investing* and *OSFS* for many datasets. (2) The efficacy of the *OSFSCM* algorithm is high for these datasets, especially those with low redundancy and high relevance. (3) The number of selected features from *OSFSCM* is greater compared with *Alpha-investing* and *OSFS* for many datasets. (4) The validity of the *OSFSCM* algorithm is verified by tests using a real-world dataset scenario.

CONFLICT OF INTEREST

The authors declare they have no conflicts of interest.

AUTHORS' CONTRIBUTIONS

Hongyi Wang: conceptualization, validation, investigation, resources; Dianlong You: methodology, formal analysis, writing—original draft, data curating, writing—review and editing, supervision.

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