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Adaptive Feature Selection based on the Most Informative Graph-based Features^{*}

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Abstract. In this paper, we propose a novel method to adaptively select the most informative and least redundant feature subset, which has strong discriminating power with respect to the target label. Unlike most traditional methods using vectorial features, our proposed approach is based on graph-based features and thus incorporates the relationships between feature samples into the feature selection process. To efficiently encapsulate the main characteristics of the graph-based features, we probe each graph structure using the steady state random walk and compute a probability distribution of the walk visiting the vertices. Furthermore, we propose a new information theoretic criterion to measure the joint relevance of different pairwise feature combinations with respect to the target feature, through the Jensen-Shannon divergence measure between the probability distributions from the random walk on different graphs. By solving a quadratic programming problem, we use the new measure to automatically locate the subset of the most informative features, that have both low redundancy and strong discriminating power. Unlike most existing state-of-the-art feature selection methods, the proposed information theoretic feature selection method can accommodate both continuous and discrete target features. Experiments on the problem of P2P lending platforms in China demonstrate the effectiveness of the proposed method.

1 Introduction

Many real-world applications, including image processing, bioinformatics analysis, face recognition, and P2P lending analysis [16], are represented by high dimensional data. However, only a small number of features are really significant to describe the target label [12]. One way to overcome this problem is to use feature selection.

Mutual information (MI) [8, 18, 15, 9] is a well-known means of measuring the mutual dependency of two variables, and has received much attention for developing new feature selection methods. Typical examples include 1) the Information-based Feature Selection method (MIFS) [8], 2) the Maximum-Relevance Minimum-Redundancy Feature Selection method (MRMR) [18], 3) the Joint-Information Feature Selection method (JMI) [20], and 4) the MIFS method under the assumption of a uniform distribution for input features (MIFS-U) [14]. Unfortunately, these methods suffer from two widely known drawbacks. First, these methods require the number of selected features in advance. Second, these methods mine subsets of the most informative features in a greedy

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manner [10]. To overcome the shortcomings, Liu et al. [15] have developed the Adaptive MI based Feature Selection method (AMIF) that can automatically determine the size of most informative feature subset, by maximizing the average pairwise informativeness. Zhang and Hancock [21] have developed a Hypergraph based Information-Theoretic Feature Selection method (HITF) that can automatically determine the most informative feature subset through dominant hypergraph clustering [17].

Unfortunately, the aforementioned information theoretic feature selection methods cannot incorporate the relationship between pairwise samples of each feature dimension. More specifically, for a dataset with N features denoted as $\mathcal{X} = \{\mathbf{f}_1, \dots, \mathbf{f}_i, \dots, \mathbf{f}_N\}$, each feature \mathbf{f}_i has M samples as $\mathbf{f}_i = \{f_{i1}, \dots, f_{ia}, \dots, f_{ib}, \dots, f_{iM}\}^T$. Traditionally, existing information theoretic feature selection methods accommodate each feature \mathbf{f}_i as a vector, and thus ignore the relationship between pairwise samples f_{ia} and f_{ib} in \mathbf{f}_i . This drawback limits the precise information theoretic measure between pairwise features. To address this shortcoming, Cui et al. [11] have proposed a new feature selection method in terms of graph-based features. They transform each vectorial feature into a graph structure that encapsulates the relationship between pairwise samples from the feature. The most relevant vectorial features are identified by selecting the graph-based features that are most similar to the graph-based target feature, in terms of the Jensen-Shannon divergence measure between graphs. Unfortunately, this method cannot adaptively determine the most relevant feature subset. It is fair to say that developing effective information theoretic feature selection method still remains a challenge.

This paper aims to overcome the shortcomings of existing information theoretic feature selection methods by developing a new algorithm that can incorporate the relationship between feature samples into the feature selection process. In summary, the main contributions are threefold. First, like Cui et al. [11], for the above dataset \mathcal{X} having N features, we transform each vectorial feature \mathbf{f}_i into a graph-based feature \mathbf{G}_i . Here, \mathbf{G}_i is a complete weighted graph, where each vertex v_a represents a corresponding sample f_{ia} in \mathbf{f}_i and each weighted edge $\{v_a, v_b\}$ represents the relationship between pairwise samples f_{ia} and f_{ib} . We use the Euclidean distance to measure the relationship between f_{ia} and f_{ib} . Similarly, for the target feature \mathbf{Y} (e.g., the class labels), we also compute a target feature graph \mathbf{G}_Y . We argue that the graph-based features can reflect richer characteristics than the original vectorial features. Furthermore, for the feature graphs \mathbf{G}_i and \mathbf{G}_Y , we probe each graph structure in terms of the steady state random walk (SSRW) [3] and compute a probability distribution of the walk visiting the vertices. Second, with the probability distributions of the feature graphs \mathbf{G}_i and \mathbf{G}_Y to hand, we propose a new information theoretic criterion to measure the joint relevance of different pairwise feature combinations with respect to the target feature, through the Jensen-Shannon divergence (JSD). Third, we use the new information theoretic measure to automatically locate the subset of the most informative and less redundant features by solving a quadratic program problem [17]. We show that, unlike most existing feature selection methods, the proposed feature selection method can accommodate both continuous and discrete target variables. Experimental results on the analysis of P2P lending platforms in China demonstrate the effectiveness of the proposed method.

2 Preliminary Concepts

2.1 The Steady State Random Walk (SSRW)

As mentioned in the previous section, we propose to use the SSRW to capture the main characteristics of the graph-based features. The main advantages of using SSRWs are twofold. First, SSRWs can accommodate weighted information residing on edges. Second, the computational complexity of probing a graph structure using SSRWs is quadratic in the number of vertices, i.e., SSRWs can be efficiently performed on graphs. As a result, SSRWs represent an elegant way of efficiently characterizing the graph-based features. Below, we review the main concepts underpinning SSRWs.

Let $G(V, E)$ be a weighted graph, V be the vertex set, and E be edge set. Assume $\omega : V \times V \rightarrow \mathbb{R}^+$ is a edge weight function. If $\omega(u, v) > 0$ ($\omega(u, v) = \omega(v, u)$), we say that (u, v) is an edge of G , i.e., the vertices $u \in V$ and $v \in V$ are adjacent. The vertex degree matrix of G is a diagonal matrix D whose elements are given by $D(v, v) = d(v) = \sum_{u \in V} \omega(v, u)$. Based on [3], the probability of the steady state random walk visiting each vertex v is $p(v) = d(v) / \sum_{u \in V} d(u)$. Furthermore, from the probability distribution $P = \{p(1), \dots, p(v), \dots, p(|V|)\}$, we can straightforwardly compute the Shannon entropy of G as

$$H_S(G) = - \sum_{v \in V} p(v) \log p(v). \quad (1)$$

2.2 The Jensen-Shannon Divergence

In information theory, the JSD is a dissimilarity measure between probability distributions. Let two (discrete) probability distributions be $\mathcal{P} = (p_1, \dots, p_a, \dots, p_A)$ and $\mathcal{Q} = (q_1, \dots, q_b, \dots, q_B)$, then the JSD between \mathcal{P} and \mathcal{Q} is defined as

$$I_D(\mathcal{P}, \mathcal{Q}) = H_S\left(\frac{\mathcal{P} + \mathcal{Q}}{2}\right) - \frac{1}{2}H_S(\mathcal{P}) - \frac{1}{2}H_S(\mathcal{Q}), \quad (2)$$

where $H_S(\mathcal{P}) = \sum_{a=1}^A p_a \log p_a$ is the Shannon entropy of the probability distribution \mathcal{P} . In [3], the JSD has been used as a means of measuring the information theoretic dissimilarity between graphs associated with their probability distributions. In this work, we are also concerned with the similarity measure between graph-based features. Therefore, we transform the JSD into its negative form and obtain the corresponding exponential function value to denote the information theoretic similarity measure I_S between probability distributions, i.e.,

$$I_S(\mathcal{P}, \mathcal{Q}) = \exp\{-I_D(\mathcal{P}, \mathcal{Q})\}. \quad (3)$$

3 Methodology of The Proposed Feature Selection Method

3.1 Graph-based Features from Vectorial Features

In this subsection, we introduce how to transform each vectorial feature into a complete weighted graph. The advantages of using the graph-based representation are twofold.

First, graph structures have stronger ability to encapsulate global topological information than vectors. Second, the graph-based features can incorporate the relationships between samples of each original vectorial feature into the feature selection process, thus reducing information loss.

Given a dataset of N features denoted as $\mathcal{X} = \{\mathbf{f}_1, \dots, \mathbf{f}_i, \dots, \mathbf{f}_N\} \in \mathbb{R}^{M \times N}$, \mathbf{f}_i represents the i -th vectorial feature and has M samples as $\mathbf{f}_i = \{f_{i1}, \dots, f_{ia}, \dots, f_{ib}, \dots, f_{iM}\}^T$. We transform each feature \mathbf{f}_i into a graph-based feature $\mathbf{G}_i(V_i, E_i)$, where each vertex $v_{ia} \in V_i$ indicates the a -th sample f_{ia} of \mathbf{f}_i , each pair of vertices v_{ia} and v_{ib} is connected by a weighted edge $(v_{ia}, v_{ib}) \in E_i$, and the weight $\omega(v_{ia}, v_{ib})$ of (v_{ia}, v_{ib}) is the Euclidean distance between f_{ia} and f_{ib} , i.e.,

$$\omega(v_{ia}, v_{ib}) = \|f_{ia} - f_{ib}\|_2. \quad (4)$$

Similarly, if the sample values of the target feature $\mathbf{Y} = \{y_1, \dots, y_a, \dots, y_b, \dots, y_M\}^T$ are continuous, its graph-based feature $\hat{\mathbf{G}}(\hat{V}, \hat{E})$ can be computed using Eq.(4) and each vertex \hat{v}_a represents the a -th sample y_a . However, for classification problems, the sample of the target feature Y is the class label c and thus takes the discrete value $c = 1, 2, \dots, C$, i.e., the samples of each feature \mathbf{f}_i belong to the C different classes. In this case, we propose to compute the graph-based target feature $\hat{\mathbf{G}}_i(\hat{V}_i, \hat{E}_i)$ for each feature \mathbf{f}_i , where the weight $\omega(\hat{v}_{ia}, \hat{v}_{ib})$ of each edge $(\hat{v}_{ia}, \hat{v}_{ib}) \in \hat{E}_i$ is

$$\omega(\hat{v}_{ia}, \hat{v}_{ib}) = \|\mu_{ia} - \mu_{ib}\|_2, \quad (5)$$

where μ_{ia} is the mean value of all samples in \mathbf{f}_i from the same class c .

Note that, constructing the graph-based feature from the original vectorial feature is an open problem. In fact, in addition to the distance measure employed in this paper, one could employ a number of alternative measures, e.g., covariance, cosine similarity, etc. Moreover, instead of a complete graph, one may want to define a sparser graph.

3.2 The Information Theoretic Criterion for Feature Selection

We propose to use the following information theoretic criterion to measure the joint relevance of different pairwise feature combinations with respect to either the continuous or discrete target feature. For a set of N features $\mathbf{f}_1, \dots, \mathbf{f}_i, \dots, \mathbf{f}_j, \dots, \mathbf{f}_N$ and the associated continuous target feature \mathbf{Y} , the relevance degree of a feature pair $\{\mathbf{f}_i, \mathbf{f}_j\}$ is

$$W_{i,j} = I_S(\mathbf{G}_i, \hat{\mathbf{G}}) \times I_S(\mathbf{G}_j, \hat{\mathbf{G}}) \times I_D(\mathbf{G}_i, \mathbf{G}_j), \quad (6)$$

where \mathbf{G}_i and $\hat{\mathbf{G}}$ are the graph-based features of \mathbf{f}_i and \mathbf{Y} , I_S is the JSD based information theoretic similarity measure defined in Eq(3), and I_D is the JSD based information theoretic dissimilarity measure defined in Eq(2). The above relevance measure consists of three terms. The first and second terms $I_S(\mathbf{G}_i, \hat{\mathbf{G}})$ and $I_S(\mathbf{G}_j, \hat{\mathbf{G}})$ are the relevance degrees of individual features \mathbf{f}_i and \mathbf{f}_j with respect to the target feature \mathbf{Y} , respectively. The third term $I_S(\mathbf{G}_i, \mathbf{G}_j)$ measures the non-redundancy between the feature pair $\{\mathbf{f}_i, \mathbf{f}_j\}$. Therefore, $W_{\mathbf{f}_i, \mathbf{f}_j}$ is large if and only if both $I_S(\mathbf{G}_i, \hat{\mathbf{G}})$ and $I_S(\mathbf{G}_j, \hat{\mathbf{G}})$ are large (i.e., both \mathbf{f}_i and \mathbf{f}_j are informative themselves with respect to the target feature \mathbf{Y}) and $I_D(\mathbf{G}_i, \mathbf{G}_j)$ is also large (i.e., \mathbf{f}_i and \mathbf{f}_j are not redundant).

For classification problems, the samples of the target feature \mathbf{Y} take the discrete value c and $c = 1, 2, \dots, C$. In this case, we compute the individual graph-based target feature $\hat{\mathbf{G}}_i$ for each feature \mathbf{f}_i , and the relevance measure defined in Eq.(6) can re-written as

$$W_{i,j} = \{S(\mathbf{f}_i)I_S(\mathbf{G}_i, \hat{\mathbf{G}}_i)\} \times \{S(\mathbf{f}_j)I_S(\mathbf{G}_j, \hat{\mathbf{G}}_j)\} \times \{I_D(\mathbf{G}_i, \mathbf{G}_j)\}, \quad (7)$$

where $S(\mathbf{f}_i)$ is the Fisher score of feature \mathbf{f}_i [13] and is defined as

$$S(\mathbf{f}_i) = \sum_{c=1}^L n_l(\mu_l - \mu)^2 / \sum_{c=1}^C n_c \sigma_c^2, \quad (8)$$

where μ_c and σ_c^2 are the mean and variance of the samples belonging to the c -th class in feature \mathbf{f}_i , μ is the mean of feature \mathbf{f}_i , and n_c is the sample number of the c -th class in feature \mathbf{f}_i . For Eq.(8), the Fisher score $S(\mathbf{f}_i)$ indicates the quality of the graph-based target feature $\hat{\mathbf{G}}_i$ for \mathbf{f}_i , i.e., a higher Fisher score $S(\mathbf{f}_i)$ means a better target feature graph $\hat{\mathbf{G}}_i$. This follows the definition of Eq.(5). More specifically, the graph-based target feature $\hat{\mathbf{G}}_i$ of original vectorial feature \mathbf{f}_i is preferred, if the distances between samples in different classes are as large as possible and the distances between data points in the same class are as small as possible. Similar to Eq.(6), the three terms of Eq.(7) have the same corresponding theoretical significance.

3.3 Determination of the Most Informative Feature Subset

We adaptively determine the most informative subset of features by solving a quadratic program problem [17]. More specifically, for a set of N features $\mathbf{f}_1, \dots, \mathbf{f}_i, \dots, \mathbf{f}_j, \dots, \mathbf{f}_N$ and the target feature \mathbf{Y} , we commence by transforming each feature into a graph-based feature. Moreover, based on the graph-based features, we construct a feature informativeness matrix \mathbf{W} , where each element $W_{i,j} \in \mathbf{W}$ represents the information theoretic measure between a feature pair $\{\mathbf{f}_i, \mathbf{f}_j\}$ based on Eq.(6) (for \mathbf{Y} is continuous) or Eq.(7) (for \mathbf{Y} is discrete). As we have stated in Section 3.2, $W_{\mathbf{f}_i, \mathbf{f}_j}$ is large if and only if both \mathbf{f}_i and \mathbf{f}_j are informative themselves with respect to the target feature \mathbf{Y} , and \mathbf{f}_i and \mathbf{f}_j are not redundant. Therefore, we locate the most informative feature subset by finding the solution of the following quadratic program problem [17]

$$\max \quad f(\mathbf{a}) = \frac{1}{2} \mathbf{a}^T \mathbf{W} \mathbf{a} \quad (9)$$

subject to $\mathbf{a} \in \mathbb{R}^N$, $\mathbf{a} \geq 0$ and $\sum_{i=1}^N a_i = 1$. The solution vector \mathbf{a} to the above quadratic program is an N -dimensional vector. When $a_i > 0$, the i -th feature \mathbf{f}_i belongs to the most informative feature subset. Therefore, the number of the selected features n can be determined by counting the positive components of vector \mathbf{a} . Pavan and Pelillo [17] have shown that the local maximum of $f(\mathbf{a})$ can be solved using the following equation

$$a_i(t+1) = a_i(t) \frac{(\mathbf{W}\mathbf{a}(t))_i}{\mathbf{a}(t)^T \mathbf{W} \mathbf{a}(t)}. \quad (10)$$

where $a_i(t)$ corresponds to the i -th feature \mathbf{f}_i at iteration t of the update process. According to the value of the element in \mathbf{a} , all features $\mathbf{f}_1, \dots, \mathbf{f}_N$ fall into two disjoint

subsets, i.e., $\mathbf{S}_1(a) = \{\mathbf{f}_i \mid a_i > 0\}$ and $\mathbf{S}_2(a) = \{\mathbf{f}_j \mid a_j = 0\}$. Clearly, the set \mathbf{S}_1 that has nonzero variables is the selection of the most informative feature subset. The features in \mathbf{S}_1 have both low redundancy and strong discriminative power.

3.4 Complete Feature Ranking

The proposed feature selection method aims to adaptively select a compact most informative feature subset that falls into the subset $\mathbf{S}_1(a) = \{\mathbf{f}_i \mid a_i > 0\}$. We can rank the feature $\mathbf{f}_i \in \mathbf{S}_1$ by evaluating the values of their indicators a_i . A higher indicator a_i means a more informative feature. Moreover, we can also rank the features contained in the unselected feature subset $\mathbf{S}_2(a) = \{\mathbf{f}_j \mid a_j = 0\}$ based on the selection method in [15]. More specifically, we compute the reward of each feature $\mathbf{f}_j \in \mathbf{S}_2$ as

$$r_j = \sum_{\mathbf{f}_i \in \mathbf{S}_1, a_i > 0} W_{i,j} a_i, \quad (11)$$

which summarizes the pairwise informativeness between the feature $\mathbf{f}_j \in \mathbf{S}_2$ and each feature $\mathbf{f}_i \in \mathbf{S}_1$. A higher r_j means a more informative feature in \mathbf{S}_2 , thus providing a measure to rank the features in \mathbf{S}_2 . Based on the feature ranking of \mathbf{S}_1 and \mathbf{S}_2 , we can obtain a Complete Feature Ranking List L , from 1 to a user-specified number.

4 Experimental Evaluations

To validate the effectiveness of the proposed feature selection approach, we perform the following experimental evaluation on a P2P dataset collected from the Peer-to-Peer (P2P) lending sector in China. The reasons for using this dataset are twofold. **First**, P2P lending data are usually high-dimensional, highly correlated, and unstable, thus representing a challenge for traditional statistical and machine learning techniques. To better analyze the P2P data, the sample relationship of the P2P data encapsulating significant information should be incorporated, when designing feature selection methods. Unfortunately, most existing feature selection methods ignore the sample relationships and may cause significant information loss. By contrast, our proposed adaptive feature selection method is able to encapsulate the sample relationship of P2P data and overcome these shortcomings. **Second**, the P2P lending industry in China has developed rapidly since 2007, with more than 3,000 P2P lending platforms and an accumulative loan amount of 12 trillion by 2015. It is of great significance to develop an effective decision aid for the credit risk analysis of the P2P platforms.

The **P2P dataset** is collected from a reputable P2P lending portal in China¹, which consists of the most popular 200 platforms (i.e., 200 samples) until Aug 2014. For each platform, we collect 19 features including 1) transaction volume, 2) total turnover, 3) average annualized interest rate, 4) total number of borrowers, 5) total number of investors, 6) online time, which refers to the foundation year of the platform, 7) operation time, i.e., number of months since the foundation of the platform, 8) registered capital, 9) weighted turnover, 10) average term of loan, 11) average full mark time, i.e., tender

¹ See the website <http://www.wdzj.com/> for more details

period of a loan raised to the required full capital, 12) average amount borrowed, i.e., average loan amount of each successful borrower, 13) average amount invested, which is the average investment amount of each successful investor, 14) loan dispersion, i.e., the ratio of the repayment amount to the total capital, 15) investment dispersion, the ratio of the invested amount to the total capital, 16) average times of borrowing, 17) average times of investment, 18) loan balance, and 19) popularity.

4.1 Identification of The Most Influential Factors for Credit Risk

We evaluate the performance of the proposed feature selection approach with respect to **continuous target features**. Specifically, we use the proposed method to perform credit risk evaluation of the P2P lending platforms. As it is difficult to obtain sufficient data of the platforms which encountered problem, we use the annualized average interest rate as an indicator of the credit risk of the P2P lending platforms. In finance, interest rate is the amount charged, expressed as a percentage of principal, by a lender to a borrower for the use of assets. When the borrower is a low-risk party, they will usually be charged a low interest rate. On the other hand, if the borrower is considered high risk, the interest rate charged will be higher. Likewise, a higher annualized average interest rate of the P2P lending platforms often indicates greater likelihood of default, i.e., higher credit risk of the platforms. Identifying the most relevant features to the interest rate can help investors effectively manage the credit risks involved in P2P lending. Therefore, in our experiment, we set the average annualized interest rate as the target feature which takes continuous values. Our purpose is to identify the most informative subset of features for the credit risk of the P2P platforms by using the proposed feature selection method. To further strengthen our findings, we also compare the proposed adaptive feature selection method associated with the SSRW (AFS-RW) with three alternative methods. These include correlation analysis (CA) and multiple linear regression (MLR), which are simple but widely applied. Furthermore, we also compare the proposed method to the most relevant graph-based feature selection method associated with the SSRW (FS-RW) [11], since it can also accommodate the continuous target feature.

Table 1 presents a comparison of the results obtained using these methods. For each method, we display the top 10 features in terms of correlation to the average annualized interest rate. Because the number of the most informative features adaptively located by AFS-RW is six, we display these results in bold. It is worth noting that the most influential factors located by the proposed AFS-RW method is in general different from the remaining three methods used for comparison. This is due to the unique characteristics of the proposed feature selection method which encourages the most informative and least redundant features to be selected. For instance, AFS-RW identifies average full mark time, transaction volume, and average amount borrowed as the most informative features. This is reasonable because a longer full mark time of the loan often reflects a higher level of credit risk and a higher amount of total transaction volume and a higher level of the average amount borrowed indicate a higher preference of both the borrowers and investors for the P2P lending platform due to a higher degree of security. Also, AFS-RW and CA consider loan balance as a relevant feature. This is also reasonable because a higher amount of loan balance often indicates a higher level of credit risk and can result in a higher interest rate. In addition, the total number of borrowers reflects

the borrowers preference for the P2P lending platforms and is a significant influential factor. A platform with a relatively low average annualized interest rate is often more attractive to the borrowers because this indicates both a lower transaction cost and a lower credit risk of the platform. However, only the proposed AFS-RW method is able to select this factor, whereas the remaining three methods rank this factor much lower. These results demonstrate the effectiveness of the proposed method for identifying the most influential factors for credit risk of P2P lending platforms.

Table 1. Comparison of four methods

Ranking	AFS-RW	FS-RW	Correlation Analysis	Multiple Linear Regression
1#	Average full mark time	Registered capital	Popularity	Loan dispersion
2#	Transaction volume	Operation time	Loan balance	Investment dispersion
3#	Average amount borrowed	Average amount invested	Average times of investment	Online tim
4#	Loan balance	Loan dispersion	Average times of borrowing	Popularity
5#	Investment dispersion	Average times of investment	Investment dispersion	Operation time
6#	Total number of borrowers	Online time	Loan dispersion	Average times of borrowing
7#	Average times of borrowing	Average term of loan	Average amount invested	Total number of borrowers
8#	Total turnover	Total number of investors	Average amount borrowed	Loan balance
9#	Average amount invested	Investment dispersion	Average full mark time	Transaction volume
10#	Weighted turnover	Popularity	Average term of loan	Weighted turnover

4.2 Classification for The Credit Rating of The P2P Lending Platforms

We evaluate the performance of the proposed feature selection approach with respect to **discrete target features**. Specifically, we aim to locate the most informative subset of features for the credit rating of the P2P platforms in China, which takes discrete values and is collected from the Report on the Development of the P2P lending industry in China, 2014-2015, issued by the *Financial Research Institute of the Chinese Academy of Social Sciences*. In this Report, only 104 platforms are included due to the strict evaluation criteria involved, among which only 42 platforms belong to the 200 platforms used in the above P2P dataset. Thus, we use the 42 platforms (i.e., samples) for the evaluation. We set the credit ranking for these platforms as the discrete target feature, and aim to locate the most informative feature subset using the proposed approach.

To evaluate the effectiveness of the features selected by the proposed approach, we set the discrete credit ranking targets as classification labels. Since there are only 42 samples and these need to be classified into four classes, it is a very challenging classification problem. In the experiment, we randomly select 50% samples as training data and the remainder as testing data. By repeating this selection process 10 times, we obtain 10 random partitions of the original data. For each partition, we identify the most relevant features via the proposed method based on the train data, and perform a 10-fold cross-validation using a C-Support Vector Machine (C-SVM) to evaluate the classification accuracy associated with the selected features based on the testing data, i.e., we use 9 folds for training and 1 fold for testing. For the C-SVM on each partition, we repeat the process 10 times and compute the average classification accuracy. Finally, we compute the average classification accuracy over the 10 partitions. To further evaluate our study, we compare the proposed method (AFS-QW) with several alternative feature selection methods. These alternative methods include: 1) the Fisher Score method (FS) [13], 2) the Mutual Information based method (MI) [19], and 3) most relevant graph-based feature selection method (FS-RW) [11]. The classification accuracy of each method is shown in Fig. 1 as a function of the number of features selected.

Fig. 1 indicates that the proposed method AFS-RW achieves the best classification accuracy (34.50%) while requiring the lowest number of features, i.e., 3 adaptively

selected features. In contrast, FS and MI both require 4 features to generate best classification accuracies. Like the proposed method, the FS-RW also achieves best accuracy with 3 features. However, only the proposed method can adaptively determine the most informative feature subset. Finally, recall that there are only 42 samples divided into 4 class for the evaluation, making this classification task very challenging. Thus, these results demonstrates the effectiveness of the proposed method.

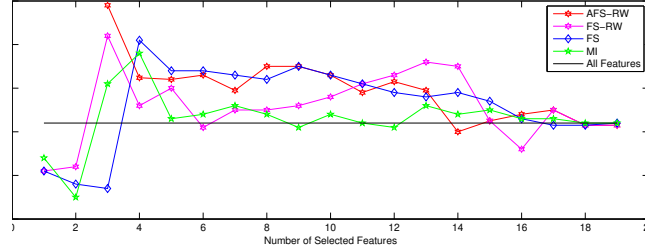


Fig. 1. Accuracy vs. Number of Selected Features for Different Feature Selection Methods.

5 Conclusion

In this paper, we have proposed an adaptive feature selection method, based on a new information theoretic criterion between graph-based features. Unlike most existing information theoretic feature selection methods, our approach has two advantages. First, it is based on graph-based features and thus incorporates the relationships between feature samples into the feature selection process. Second, it can accommodate both continuous and discrete target features. Experiments on the analysis of P2P lending platforms in China demonstrate the effectiveness of the proposed feature selection method.

We will extend our method in a number of ways. First, in our previous works [5, 4], we have developed a number of quantum Jensen-Shannon kernels using both the continuous-time and discrete-time quantum walks. It is interesting to extend the proposed feature selection method using the classical Jensen-Shannon divergence to that using its quantum counterpart. Second, we will also use our previous graph kernel measures as the graph similarity measures for our feature selection frameworks [7, 2, 6]. We will explore the performance of our feature selection method associated with different graph kernels. Third, the proposed feature selection method only considers the relationship between pairwise features, i.e., it only evaluates the two-order relationship between features. Our future work will extend the proposed method into a high-order feature selection method by establishing higher order relationship between features. Finally, it is interesting to establish hypergraph-based features [1] and thus develop a new hypergraph-based feature selection method.

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