A DYNAMIC WRAPPER METHOD FOR FEATURE DISCRETIZATION AND SELECTION

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Abstract: In many learning problems, an adequate (sometimes discrete) representation of the data is necessary. For instance, for large number of features and small number of instances, learning algorithms may be confronted with the *curse of dimensionality*, and need to address it in order to be effective. Feature selection and feature discretization techniques have been used to achieve adequate representations of the data, by selecting an adequate subset of features with a convenient representation. In this paper, we propose static and dynamic methods for feature discretization. The static method is unsupervised and the dynamic method uses a wrapper approach with a quantizer and a classifier, and it can be coupled with any static (unsupervised or supervised) discretization procedure. The proposed methods attain efficient representations that are suitable for learning problems. Moreover, using well-known feature selection methods with the features discretized by our methods leads to better accuracy than with the features discretized by other methods or even with the original features.

1 INTRODUCTION

Datasets with large numbers of features and (relatively) smaller numbers of instances are challenging for machine learning methods. In fact, it is often the case that many features are irrelevant or redundant for the task at hand (*e.g.*, learning a classifier) (Yu et al., 2004; Peng et al., 2005); this may be specially harmful with relatively small training sets, where these irrelevancies/redundancies are harder to detect.

To deal with such datasets, *feature selection* (FS) and *feature discretization* (FD) methods have been proposed to obtain data representations that are more adequate for learning. FD aims at finding a representation of each feature that contains enough information for the learning task at hand, while ignoring minor (possibly irrelevant) fluctuations. FS aims at reducing the number of features, thus directly targeting the curse of dimensionality problem, often allowing the learning algorithms to obtain classifiers with better performance. A byproduct of FD and FS is a reduction of the memory required to represent the data.

Both FD and FS are topics with a long research history and a vast literature; regarding FD, see (Dougherty et al., 1995), (Liu et al., 2002), (Witten and Frank, 2005) for extensive reviews of many methods; regarding FS, see (Guyon et al., 2006), (Hastie et al., 2009), and (Escolano et al., 2009) for comprehensive coverage and pointers to the literature.

1.1 Our Contribution

In this paper, we propose an unsupervised method for static FD and a new method for dynamic FD. The dynamic discretization method uses a wrapper approach with a quantizer and a classifier, and can be coupled with any static (unsupervised or supervised) discretization procedure. The dynamic method assesses the performance of each feature as discretization is carried out; if it is found that the original representation is preferable to the discretized one, the original feature is kept.

The remaining text is organized as follows. Section 2 reviews supervised and unsupervised FD and FS techniques. Section 3 presents the proposed static and dynamic methods for FD. Section 4 reports the experimental evaluation of our methods in comparison with other techniques. Finally, Section 5 ends the paper with some concluding remarks and directions for future work.

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2 BACKGROUND

This section briefly reviews some of the most common unsupervised and supervised FD and FS techniques that have proven effective for many learning problems. This description is hugely far from being exhaustive, as FD and FS are two fields with a long research history. The interested reader is referred to the works of (Dougherty et al., 1995), (Kotsiantis and Kanellopoulos, 2006), (Liu et al., 2002), and (Witten and Frank, 2005), for reviews of FD methods. Reviews of FS methods were done by (Guyon and Elisseeff, 2003), (Guyon et al., 2006), (Hastie et al., 2009), and (Escolano et al., 2009); see also the following special issue: jmlr.csail.mit.edu/papers/special/feature03.html.

2.1 Feature Discretization

FD can be performed in supervised or unsupervised modes, *i.e.*, using or not the class labels, and aims at reducing the amount of memory as well as improving classification accuracy (Witten and Frank, 2005). The supervised mode may lead, in principle, to better classifiers. In the context of unsupervised scalar FD (Witten and Frank, 2005), two efficient techniques are commonly used:

- . *equal-interval binning* (EIB), *i.e.*, uniform quantization with a given number of bits per feature;
- . *equal-frequency binning* (EFB), *i.e.*, non-uniform quantization yielding intervals such that, for each feature, the number of occurrences in each interval is the same, leading to a uniform (*i.e.*, maximum entropy) distribution; this technique is also known as *maximum entropy quantization*.

In EIB, the range of values is divided into bins of equal width. It is simple and easy to implement, but it is very sensitive to outliers, thus may lead to inadequate discrete representations. The EFB method is less sensitive to outliers. The quantization intervals have smaller width in regions where there are more occurrences of the values of each feature.

Recently, we have proposed (Ferreira and Figueiredo, 2011) an unsupervised scalar discretization method, based on the well-known Linde-Buzo-Gray (LBG) algorithm (Linde et al., 1980). The LBG algorithm is applied individually to each feature and stopped when the MSE distortion falls below a threshold Δ or when the maximum number of bits q per feature is reached (setting Δ to 5% of the range of each feature and $q \in \{4, ..., 10\}$ were found to be adequate choices). That algorithm, named *unsupervised* LBG (U-LBG 1), which produces a variable number of bits

per feature, has been shown to lead to better classification results than EFB on different kinds of (sparse and dense) data (Ferreira and Figueiredo, 2011). The key idea of using the LBG algorithm in this context is that if we can represent a feature with low MSE, we have a discrete version that approximates well the continuous version of that feature, thus this representation should be adequate for learning. Algorithm 1 presents the U-LBG1 procedure.

Algorithm 1: U-LBG1.

Input: X, n × p matrix training set (p features, n patterns). Δ: maximum expected distortion. q: the maximum number of bits per feature.
Dutput: \widetilde{X} : $n \times p$ matrix, discrete feature training set. $Q^1,, Q^p$: set of p quantizers (one per feature).
1: for $i = 1$ to p do
2: for $b = 1$ to q do
3: Apply the LBG algorithm to the <i>i</i> -th feature to
obtain a <i>b</i> -bit quantizer $Q_b(\cdot)$;
4: Compute $MSE_i = \frac{1}{n} \sum_{j=1}^n (X_{ij} - Q_b(X_{ij}))^2$;
5: if (MSE _{<i>i</i>} $\leq \Delta$ or $b = q$) then
6: $Q^{i}(\cdot) = Q_{b}(\cdot); \{/\text{* Store the quantizer. */}\}$
7: $\widetilde{X}_i = Q^i(X_i);$ {/* Quantize feature. */}
8: break; {/* Proceed to the next feature. */}
9: end if
10: end for
11: end for

It has been found that unsupervised FD methods tend to perform well in conjunction with several classifiers; in particular, the EFB method in conjunction with *naive Bayes* (NB) classification produces very good results (Witten and Frank, 2005). It has also been found that applying FD with both EIB and EFB to microarray data, in conjunction with *support vector machine* (SVM) classifiers, yields good results (Meyer et al., 2008).

There are also many supervised approaches to feature discretization. (Fayyad and Irani, 1993) have applied an entropy minimization heuristic to choose the cut points, and thus the discretization intervals. The experimental results show that the proposed method leads to the construction of better decision trees than the previous methods. An efficient FD algorithm for use in the construction of Bayesian belief networks (BBN), was proposed by (Clarke and Barton, 2000). The partitioning minimizes the information loss, relative to the number of intervals used to represent the variable. Partitioning can be done prior to BBN construction or extended for repartitioning during construction. A supervised static, global, incremental, and top-down discretization algorithm based on class-attribute contingency coefficient was proposed by (Tsai et al., 2008).

Very recently, a supervised discretization algo-

rithm based on *correlation maximization* (CM) was proposed by (Zhu et al., 2011); it uses *multiple correspondence analysis* (MCA) to capture correlations between multiple variables. For each numeric feature, the correlation information obtained from MCA is used to build the discretization algorithm that maximizes the correlations between feature intervals/items and classes.

2.2 Feature Selection

In this subsection, we briefly describe two FS methods that have prove successful in many problems. For this reason, we have included them on the experimental evaluation of our methods.

The well-known (supervised) *Fisher ratio* (FiR) (Furey et al., 2000) of each feature is defined as

$$\operatorname{FiR}_{i} = \frac{\left| \bar{X}_{i}^{(-1)} - \bar{X}_{i}^{(+1)} \right|}{\sqrt{\operatorname{var}(X_{i})^{(-1)} + \operatorname{var}(X_{i})^{(+1)}}}, \qquad (1)$$

where $\bar{X}_i^{(\pm 1)}$ and $var(X_i)^{(\pm 1)}$, are the mean and variance of feature *i*, for the patterns of each class. The FiR measures how well each feature alone separates the two classes.

The (supervised) *minimum redundancy maximum relevancy* (mRMR) method of (Peng et al., 2005) adopts a filter approach to the problem of FS, thus being fast and applicable with any classifier. The key idea is to compute both the redundancy between features and the relevance of each feature. The redundancy is computed by the *mutual information* (MI) (Cover and Thomas, 1991) between pairs of features, whereas relevance is measured by the MI between features and class label.

3 PROPOSED METHODS

This section presents our proposals for FD. Subsection 3.1 presents a static unsupervised FD method, whereas Subsections 3.2 and 3.3 detail our dynamic wrapper methods for FD and joint FS/FD, respectively.

3.1 Static Discretization

We address static discretization with a new unsupervised proposal. The first new proposal for FD is named U-LBG2, and it is a minor modification of previous method U-LBG1, described as Algorithm 1 in Subsection 2.1. The proposed modification is to use a fixed, rather than variable, number of bits per feature, q, according to the MSE distortion criterion. This method exploits the same key idea as the previous one, that is, a discretization with a low MSE will provide an accurate representation of each feature, being suited for learning purposes. Algorithm 2 describes the proposed U-LBG2 method.

Algorithm 2: U-LBG2.	
Input: <i>X</i> , $n \times p$ matrix train <i>q</i> : the maximum num	ing set (p features, n patterns). mber of bits per feature.
Output: \widetilde{X} : $n \times p$ matrix, di $Q^1,, Q^p$: set of	screte feature training set. p quantizers (all with q bits).
1: for $i = 1$ to p do	10
2: Apply the LBG algor	rithm to the <i>i</i> -th feature to ob-
tain a q-bit quantizer	$Q(\cdot);$
3: $Q^i(\cdot) = Q(\cdot);$	$\{/*$ Store the quantizer. $*/\}$
4: $\widetilde{X}_i = Q^i(X_i);$	{/* Quantize feature. */}
5: end for	:0

As compared to U-LBG1 algorithm, the key differences are: now we are using more bits, since each discretized feature will be given the same (maximum) number of bits q; only one quantizer is learned for each feature. Both unsupervised LBG-based procedures aim at obtaining quantizers that represent the features with a small distortion. The proposed procedures are more complex than either EIB or EFB, thus may be expected to perform better.

3.2 Dynamic Discretization Wrapper

The key motivations to propose a wrapper dynamic discretization method are as follows. This method, by adopting a wrapper working mode in conjunction with a classifier, has higher complexity than a static discretization method. However, it is expected that the increased complexity pays off in the sense that we should be able to choose a more adequate number of bits per feature, as compared to static discretization methods. As a consequence, we may hope to attain better classification accuracy with the dynamically discretized features.

The proposed approach for dynamic discretization relies on the use of a static unsupervised or supervised FD algorithm (such as EIB, EFB, U-LBG1, or U-LBG2) which is applied sequentially to the set of features. The key idea is to discretize each feature with an increasing number of bits and to evaluate how the classification accuracy evolves with the discretization of each feature. The classification accuracy is compared against the accuracy obtained with the feature in its original representation. The number of bits that leads to the maximum accuracy is chosen to discretize the feature.

Before giving the details of our algorithm, we show some experimental results that motivate the de-



Figure 1: Test set error rate of naïve Bayes using only a single feature, discretized with q = 4 bit by the U-LBG2 algorithm, on the WBCD dataset. The horizontal dashed line is the test set error rate obtained with the full set of p=30 features.



Figure 2: Test set error rate of naïve Bayes using only a single feature, discretized with q = 8 bit by the U-LBG2 algorithm, on the WBCD dataset. The horizontal dashed line is the test set error rate obtained with the full set of p=30 features.

tails of the method. Fig. 1 and Fig. 2 show the test set error rate of each feature obtained for the WBCD dataset using the naïve Bayes classifier, on discrete features with the static U-LBG2 procedure with q = 4 and q = 8, respectively. We observe that the test set error rates achieved individually by several features are quite close to the error attained with the full set of features (for example, features 8, 23, or 28 in Fig. 1). Another interesting remark about Figs. 1 and 2 is that increasing the number of bits per feature does not necessarily lead to a decrease in the classification error; again, if we look into the individual test set error rates of features 8, 23, and 28, now with q = 8 bits (Fig. 2), we observe an increase in the test set error rates with respect to q = 4 bits (Fig. 1).

In order to gain insight into how the test set error rate of each feature evolves during discretization, we compare the test set error rate of each feature with its U-LBG2 discretized version with $q \in \{1, ..., 10\}$. Fig. 3 and Fig. 4 show the test set error rate of features 17 and 25, again with the naïve Bayes classifier, respectively, for the WBCD dataset.



Figure 3: Test set error rate of naïve Bayes using solely feature 17 of the WBCD dataset (original feature and U-LBG2 discretized with $q \in \{1, ..., 10\}$).



Figure 4: Test set error rate of naïve Bayes using solely feature 25 of the WBCD dataset (original feature and U-LBG2 discretized with $q \in \{1, ..., 10\}$).

In the case of feature 17, discretization never leads to a lower test set error rate, as compared to the original representation. On the other hand, for feature 25, the use of a larger number of bits leads to an improvement in the accuracy. These experimental results show typical situations that we observe with different types of data and depend on the statistics of each feature, leading us to the following observations:

- some features are worth to be discretized up to number of bits *q*;
- for other features, it is preferable *not* to discretize them.

Algorithm 3 details our *dynamic wrapper discretization* (DWD) method. We use the following notation: @class(X,y) denotes a function that learns

some classifier (*e.g.*, support vector machine, *naive Bayes*, *K*-nearest-neighbors) from the training set in matrix *X* with labels *y* and returns the obtained error rate; ($@quant(\cdot,b)$) denotes any of the unsupervised (or supervised) scalar static discretization algorithms, such as those mentioned above or any other, with *b* bits. This algorithm performs its actions solely on the training set portion of the data; it does not require the existence of a separate hold-out test set.

Algorithm 3: DwD - Dynamic wrapper Discretiza-
tion.
Input: X, $n \times p$ matrix training set (p features, n patterns)
y. <i>n</i> -tengui vector with class labers.
q: the maximum number of ons per reature.
@quant: a static discretizer.
\widetilde{w} class: a supervised classifier.
Output: X: $n \times p$ matrix, discretized version on X.
$Q^1,, Q^p$: set of p quantizers (one per feature).
1: for $i = 1$ to p do
2: $\operatorname{err}_{i0} \leftarrow @class(X_i, y)$: training error rate using only
the i -th feature;
3: for $b_i = 1$ to q do
4: $\widetilde{X}_i = @quant(X_i, b_i);$
5: $\operatorname{err}_{ib} \leftarrow @class(\widetilde{X}_i, y);$
6: end for
7: end for
8: Find $b_i = \arg\min_{j \in \{0, 1,, q\}} \operatorname{err}_{ij}$, for $i = 1,, p$.
9: for $i = 1$ to <i>p</i> do
10: if $b_i = 0$ then
11: $\widetilde{X}_i = X_i$; {/* Don't discretize the <i>i</i> -th feature */
12: else
13: $Q^{i}(\cdot) = @quant(\cdot, b_{i}); $ {/* Store quantizer*/
14: $\widetilde{X}_i = Q_{h_i}^i(X_i);$
15: end if
16: end for

In line 2, @*class* provides the baseline error, that is, the training error rate with the original representation of each feature. A similar idea is applied in line 5, where the classifier is applied to each discretized feature. Notice that if a discretized feature never reaches a training error below the baseline, the original representation is kept. We thus have a *dynamic wrapper discretization* procedure that produces a hybrid dataset in the sense that it may contain both discretized and non-discretized features. As a final note on DWD (Algorithm 3), notice that the for loop in line 3 does not need to start at $b_i = 1$; we can set the minimum number of bits per feature to some small value, such as 3 or 4, thus computing fewer quantizers and performing fewer evaluations.

3.3 Optimized Dynamic Discretization

For medium to high-dimensional datasets, the pro-

posed DWD method, as detailed in Algorithm 3, becomes computationally demanding. The need discretize each feature several times, and evaluate the corresponding classification accuracies, can make this method prohibitive for higher dimensions (as it happens with many wrapper methods). The efficiency of both the quantizer and the classifier is a key point to avoid this computational burden. In order to decrease the running time of our DWD algorithm, we have considered that:

- the evolution of the test set error rates shown in Fig. 1 and Fig. 2 suggest that for some features there is no improvement on the classification performance if we use a larger number of bits; moreover, some (irrelevant) features will always lead to low accuracy, regardless of the number of bits we use for discretization;
- the results in Fig. 3 and Fig. 4, show that for some features, there is no gain in discretizing them.

Combining these two ideas we propose the following optimization in order to delete the irrelevant features as a pre-processing stage: after computing the error of each feature, $\operatorname{err}_{i0} \leftarrow @class(X_i, y)$ in line 2 of Algorithm 3, we keep only a fraction η of the top rank features. We thus avoid the discretization of many irrelevant features, saving execution time while simultaneously improving the classification accuracy and reducing the amount of memory needed to represent the dataset. This optimization can been seen as wrapped FS process acting as a pre-processing stage for the DWD algorithm; for this reason, we name this optimized version of DWD as DWD-FS, being a wrapper for both discretization and selection.

4 EXPERIMENTAL EVALUATION

This section reports experimental results obtained by our FD techniques on several public domain datasets. We use linear *support vector machines* (SVM), *naïve Bayes* (NB), and *K*-nearest-neighbors (KNN) (with K = 3) classifiers, implemented in the PRTools¹ toolbox (Duin et al., 2007). We start by assessing the behavior of static discretization methods in Subsection 4.1 and proceed to the analysis of the dynamic discretization methods in Subsection 4.2. In Subsection 4.3, we perform a running time analysis of these methods. Finally, in Subsection 4.4, we apply FS methods on the original and on the discretized features to check if the discrete features lead to an increase in the classification performance. The experi-

¹http://www.prtools.org/prtools.html

Dataset Name	р	С	n
Phoneme	5	2	5404
Pima	8	2	768
Abalone	8	2	4177
Contraceptve	9	2	1473
Wine	13	3	178
Hepatitis	19	2	155
WBCD	30	2	569
Ionosphere	34	2	351
SpamBase	54	2	4601
Lung	55	3	32
Arrhythmia	279	16	452
Colon	2000	2	62
SRBCT	2309	4	83
Lymphoma	4026	9	96
Leukemia 1	5327	3	72
9-Tumors	5726	9	60

Table 1: 11 UCI datasets and 5 microarray datasets used in the experiments; p, c, and n are the number of features, classes, and patterns, respectively.

ments were carried out on a common laptop computer with 2.16 GHz CPU and 4 Gb of RAM.

Table 1 briefly describes the public domain benchmark datasets from the UCI Repository (Frank and Asuncion, 2010) that were used in our experiments. We chose several well-known datasets with different kinds of data. We have also included public domain microarray gene expression datasets².

4.1 Analysis of Static Discretization

Fig. 5 shows the typical MSE decay obtained by the U-LBG1 algorithm using up to q = 10 bits, when discretizing features 1 and 18 of the Hepatitis dataset. This plot shows that even for features that start with a high distortion (with a single bit), the MSE drops fast (roughly exponentially fast).

Table 2 shows a comparison of three static FD methods, namely EFB, U-LBG1, and U-LBG2 for some standard datasets, using up to q = 7 bits. The EIB method is not considered here, since it usually attains poorer performance than the EFB method. For each FD method, we show the average test set error rate of the naïve Bayes classifier for ten runs with different training/test replications and the total number of bits allocated for the set of quantizers. Table 3 shows a similar set of results, using linear SVM classifiers. Comparing the results in Table 2 and Table 3, we see that on these datasets, the linear SVM classifiers attain better results than the naïve Bayes classifiers. In some cases, the linear SVM classifiers attains better performance on the original features than on the discretized ones. The EFB method has good results,



Figure 5: MSE evolution as a function of the number of bits $q \in \{1, ..., 10\}$, for the features 1 and 18 of the Hepatitis dataset, using the U-LBG1 algorithm.

despite its simplicity; however, on microarray data which we have a large number of features and a small number of patterns (large p, small n) as well as multiclass problems, the U-LBG methods tend to perform better. Moreover, on the micoarray datasets, the naïve Bayes classifier performs poorly, so we don't even report those results.

4.2 Dynamic Discretization

We now compare static discretization versus DWD and its optimized version DWD-FS, as described in Subsection 3.3. Table 4 shows a comparison of the static FD methods, EFB and U-LBG1 with their dynamic versions incorporated into our DWD method (Algorithm 2), with linear SVM classifiers for wrapping and evaluation.

The results in Table 4 suggest that the DWD method tends to produce better results for datasets with higher number of features. For low-dimensional datasets, (roughly with p < 20), the additional complexity of the dynamic wrapper method does not lead to better results as compared to the static versions. The DWD method tends to produce discrete representations with a smaller number of bits, as compared to the static counterparts. For the higher-dimensional datasets in Table 4, the use of the DWD algorithm generally improves the performance of the wrapper static discretizer for both EFB an U-LBG1. The Phoneme and Abalone datasets exhibit a behavior such that the use of the original features is preferable; none of the static or dynamic versions attains better results.

Table 5 shows the test results of static, DWD, and DWD-FS U-LBG2 discretization with q = 7 bits. For the DWD-FS algorithm we keep the percentage η of the top rank features; the choice of η leads to select *m* features. In these tests, we use the KNN classifier.

²http://www.gems-system.org/

Table 2: Total number of bits per pattern (T. Bits) and Test Set Error Rate (Err, average of ten runs, with different random training/test partitions) for the static discretization methods, with up to q = 7 bits, using the naïve Bayes classifier. The best test set error rate is in bold face (in case of a tie, the best is considered the one with fewer bits).

		EFB		U-LBG1		U-LI	BG2
Dataset	Original Err	T. Bits	Err	T. Bits	Err	T. Bits	Err
Phoneme	21.30	30	22.30	9	22.80	30	20.60
Pima	25.30	48	25.20	30	25.20	48	25.80
Abalone	28.00	48	27.60	15	27.20	48	27.70
Contraceptive	34.80	54	31.40	15	38.00	54	34.80
Wine	3.73	78	4.80	27	3.20	78	3.20
Hepatitis	20.50	95	21.50	32	21.00	39	18.00
WBCD	5.87	180	5.13	60	5.87	180	5.67
Ionosphere	10.60	198	9.80	49	17.40	198	11.00
SpamBase	15.27	324	13.40	54	15.73	324	15.67
Lung	35.00	318	35.00	74	35.83	318	35.00
Arrhythmia	32.00	1392	51.56	553	30.22	1392	41.56

Table 3: Total number of bits per pattern (T. Bits) and Test Set Error Rate (Err, average of ten runs, with different random training/test partitions) for the static discretization methods, with up to q = 7 bits, using the linear SVM classifier. The best test set error rate is in bold face (in case of a tie, the best is considered the one with less bits).

		EFB		U-LBG1		U-LI	BG2
Dataset	Original Err	T. Bits	Err	T. Bits	Err	T. Bits	Err
Phoneme	22.60	30	24.00	8	21.80	30	22.70
Pima	27.50	48	28.50	27	29.30	48	27.60
Abalone	24.30	48	23.20	16	27.80	48	23.70
Contraceptive	36.50	54	34.60	15	39.80	54	37.20
Wine	4.80	78	1.87	25	5.33	78	1.33
Hepatitis	14.50	114	16.00	33	16.00	114	13.00
WBCD	4.67	180	3.07	59	2.80	180	2.80
Ionosphere	16.60	198	19.60	43	13.80	198	16.00
SpamBase	12.93	324	16.73	54	20.40	324	18.53
Lung	18.33	330	17.50	85	19.17	330	20.83
Arrhythmia	33.33	1392	31.78	550	32.89	1392	33.33
Colon	14.44	12000	11.11	9723	12.78	12000	10.56
SRBCT	0.00	13848	0.53	2793	0.20	13848	0.37
Lymphoma	0.57	24156	0.57	5910	0.00	24156	0.57
Leukemia1	11.43	26635	10.48	24306	5.71	26635	7.62
9-Tumors	22.22	28630	14.67	24906	12.00	28630	14.22

Table 4: Total number of bits per pattern (T. Bits) and Test Set Error Rate (Err, average of ten runs, with different random training/test partitions) for the static and dynamic discretization methods for EFB and U-LBG1, with up to q = 7 bits, using linear SVMs. The best test set error rate is in **bold** face (in case of a tie, the best is considered the one with fewer bits).

	11	Static					DWD A	lgorithm	
	0	EFB		U-LBG1		EFB		U-LBG1	
Dataset	Original Err	T. Bits	Err	T. Bits	Err	T. Bits	Err	T. Bits	Err
Phoneme	21.17	35	23.33	8	23.33	29	24.33	7	21.50
Pima	27.67	56	27.33	28	28.67	28	26.83	25	28.50
Abalone	23.00	56	24.67	16	27.67	26	25.33	13	26.00
Contraceptive	34.00	63	31.67	15	36.67	22	34.50	14	34.00
Wine	2.40	65	0.53	25	4.27	55	1.60	20	1.87
Hepatitis	20.50	95	21.50	32	21.00	39	18.00	29	19.00
WBCD	4.22	210	3.22	59	2.11	125	2.78	52	4.56
Ionosphere	13.40	165	20.00	43	13.00	126	18.40	40	12.40
SpamBase	14.20	260	15.87	52	17.73	82	11.87	52	14.20
Lung	22.22	371	23.61	76	25.00	81	22.22	65	22.22

These results show the adequacy of the DWD-FS algorithm as compared to the other two. In the major-

ity of these tests, it attains the lowest test set error rate using fewer features and fewer bits per feature; it thus Table 5: Total number of bits per pattern (T. Bits) and Test Set Error Rate (Err, average of ten runs, with different random training/test partitions) for the static, DWD, and DWD-FS discretization methods for U-LBG2, with up to q = 7 bits, using the KNN (K = 3) classifier. η is the percentage of the top rank features and *m* is the average number of features for the ten runs. The best test set error rate is in bold face (in case of a tie, the best is considered the one with less bits).

		Static U	-LBG2	DWD U	-LBG2		DWD-	FS U-LB(52
Dataset	Original Err	T. Bits	Err	T. Bits	Err	η	m	T. Bits	Err
Phoneme	22.15	35	22.35	35	22.15	0.8	4.0	28	21.90
Pima	26.10	56	26.10	50	26.00	0.8	6.0	40	27.00
Abalone	26.20	56	24.60	49	26.50	0.8	6.0	42	24.20
Contraceptive	44.60	63	45.30	13	45.50	0.8	7.0	11	43.60
Wine	27.20	91	6.13	84	14.67	0.8	10.0	65	12.27
Hepatitis	29.50	133	29.00	42	30.00	0.8	15.0	38	29.50
WBCD	7.07	210	3.93	209	6.93	0.8	24.0	167	3.80
Ionosphere	18.40	231	18.80	215	16.00	0.8	26.0	174	20.80
SpamBase	16.07	378	17.20	186	23.00	0.8	42.4	169	26.47
Lung	23.33	378	28.33	63	24.17	0.8	42.8	52	22.50

Table 6: Total time (in seconds) taken to discretize features by the three static FD methods, namely EFB, U-LBG1, and U-LBG2, using up to q = 7 bits. The fastest discretization method is in bold face.

Dataset	EFB	U-LBG1	U-LBG2
Phoneme	0.20	0.27	0.79
Wine	0.09	0.05	0.14
Hepatitis	0.10	0.05	0.10
WBCD	0.24	0.29	0.86
SpamBase	0.32	0.07	0.40
Arrhythmia	1.23	0.96	1.55
Colon	11.66	21.81	18.12
SRBCT	32.60	23.57	52.57
Lymphoma	32.17	18.72	49.25
Leukemia 1	31.56	47.39	38.74
9-Tumors	43.49	64.85	55.77

leads to an improvement on the results of the KNN classifier on both the original and U-LBG2 features. The only test in which there are no benefits from the discretization is on the sparse data SpamBase dataset; in this case, the original features are preferable for the KNN classifier.

4.3 **Running Time Analysis**

Table 6 shows a comparison of the time taken to discretize features by the three static FD methods, EFB, U-LBG1, and U-LBG2, using up to q = 7 bits. The U-LBG1 (Algorithm 1) also uses $\Delta = 0.05$ range(X_i). These results show that the EFB and U-LBG2 tend to take roughly the same time, allocating a maximum of q bits per feature. The U-LBG1 algorithm usually is faster since it stops before reaching the maximum q bits; in fact, many features are discretized with a number of bits much smaller than q.

Table 7 shows a similar comparison of the time taken to discretize features by the DWD and DWD-FS methods (with $\eta = 0.8$) using EFB discretization and naïve Bayes classifier, using up to q = 7 bits. The

Table 7: Time (in seconds) taken to discretize features by EFB, DWD and DWD-FS methods (wrapped with EFB discretization and naïve Bayes classifier), using up to q = 7 bits. We show the average time for ten runs.

Dataset	EFB	DWD EFB	DWD-FS EFB
Phoneme	0.04	1.94	3.04
Wine	0.10	5.30	6.97
Hepatitis	0.09	7.01	9.80
WBCD	0.20	11.40	21.94
SpamBase	0.34	19.92	28.13
Arrhythmia	1.26	141.27	175.56
Colon	12.11	725.51	962.57
SRBCT	33.95	1270.97	1621.65

dynamic versions take much more time than the static version. The choice of the classifier also influences the time taken for the discretization. For medium to high-dimensional datasets (p > 200), the implementations (without optimizations) of DWD and DWD-FS are too time-consuming for practical applications.

4.4 Leveraging Feature Selection

In order to asses the quality (informativeness) of the discretized features, we run a few tests using FS on the original and on the discretized features. The key idea of these tests is to show how the use of FS on the discretized features can have benefits, as compared to the original ones.

In Fig. 6, we report naïve Bayes classifier test set error rates with the features selected by the mrMR method; we compare the original features with the discretized versions obtained with the static and dynamic U-LBG1 methods. We observe that the use discretized features seems to help the FS criterion, since lower test error rates are achieved when compared with the original ones.

In Fig. 7 and Fig. 8, we report naïve Bayes test set error rates with the features selected by the FiR method, for the Hepatitis and the Ionosphere datasets,



Figure 6: Test set error rate (average of ten runs, with different training/test partitions), as functions of the number of features, for the naïve Bayes classifier, using FS by the mrMR method, for the WBCD dataset on original, static, and dynamically discretized features.



Figure 7: Test set error rate (average of ten runs, with different training/test partitions), as functions of the number of features, for the naïve Bayes classifier, using FS by the FiR method, for the Hepatitis dataset on original, static, and dynamically discretized features.

respectively; we compare the original features with the discretized versions obtained with the static and dynamic EFB methods. For both datasets, the discrete features are more adequate for FS as compared to the original features.

5 CONCLUSIONS

In this paper, we have proposed static and dynamic methods for feature discretization (FD). The static method is unsupervised and the dynamic method uses a wrapper approach with a quantizer and a classifier, and it can be coupled with any static unsupervised (or supervised) discretization procedure. The key idea of the dynamic method is to assess the performance of



Figure 8: Test set error rate (average of ten runs, with different training/test partitions), as functions of the number of features, for the naïve Bayes classifier, using FS by the FiR method, for the Ionosphere dataset on original, static, and dynamically discretized features.

each feature as discretization is carried out; if the original representation is preferable to the discrete feature, then it is kept. An optimized version of this dynamic method uses a pre-processing stage which consists in a wrapper feature selection process.

The proposed methods, equally applicable to binary and multi-class problems, attain efficient representations, suitable for learning problems. Our experimental results, on public-domain datasets with different types of data, show the competitiveness of our techniques when compared with previous approaches. The use of the features discretized by our methods lead to better accuracy than using the original or discretized features by other methods.

As future work, we plan to optimize the implementation of our dynamic discretization method, and to devise its filter version, suitable to tackle dynamic discretization on (very) high-dimensional datasets.

REFERENCES

- Clarke, E. and Barton, B. (2000). Entropy and MDL discretization of continuous variables for Bayesian belief networks. *International Journal of Intelligent Systems*, 15(1):61–92.
- Cover, T. and Thomas, J. (1991). *Elements of Information Theory*. John Wiley & Sons.
- Dougherty, J., Kohavi, R., and Sahami, M. (1995). Supervised and unsupervised discretization of continuous features. In *International Conference Machine Learning — ICML'95*, pages 194–202. Morgan Kaufmann.
- Duin, R., Juszczak, P., Paclik, P., Pekalska, E., Ridder, D., Tax, D., and Verzakov, S. (2007). PRTools4.1: A Matlab Toolbox for Pattern Recognition. Technical report, Delft Univ. Technology.

Republications

- Escolano, F., Suau, P., and Bonev, B. (2009). *Information Theory in Computer Vision and Pattern Recognition*. Springer.
- Fayyad, U. and Irani, K. (1993). Multi-interval discretization of continuous-valued attributes for classification learning. In *Proceedings of the International Joint Conference on Uncertainty in AI*, pages 1022–1027.
- Ferreira, A. and Figueiredo, M. (2011). Unsupervised joint feature discretization and selection. In 5th Iberian Conference on Pattern Recognition and Image Analysis - IbPRIA2011, pages 200–207, Las Palmas, Spain.
- Frank, A. and Asuncion, A. (2010). UCI machine learning repository, available at http://archive.ics.uci.edu/ml.
- Furey, T., Cristianini, N., Duffy, N., Bednarski, D., Schummer, M., and Haussler, D. (2000). Support vector machine classification and validation of cancer tissue samples using microarray expression data. *Bioinformatics*, 16(10):906–914.
- Guyon, I. and Elisseeff, A. (2003). An introduction to variable and feature selection. *Journal of Machine Learning Research*, 3:1157–1182.
- Guyon, I., Gunn, S., Nikravesh, M., and Zadeh (Editors), L. (2006). Feature Extraction, Foundations and Applications. Springer.
- Hastie, T., Tibshirani, R., and Friedman, J. (2009). The Elements of Statistical Learning. Springer, 2nd edition.
- Kotsiantis, S. and Kanellopoulos, D. (2006). Discretization techniques: A recent survey. GESTS International Transactions on Computer Science and Engineering, 32(1).
- Linde, Y., Buzo, A., and Gray, R. (1980). An algorithm for vector quantizer design. *IEEE Trans. on Communications*, 28:84–94.
- Liu, H., Hussain, F., Tan, C., and Dash, M. (2002). Discretization: An Enabling Technique. *Data Mining and Knowledge Discovery*, 6(4):393–423.
- Meyer, P., Schretter, C., and Bontempi, G. (2008). Information-theoretic feature selection in microarray data using variable complementarity. *IEEE Journal* of Selected Topics in Signal Processing (Special Issue on Genomic and Proteomic Signal Processing), 2(3):261–274.
- Peng, H., Long, F., and Ding, C. (2005). Feature selection based on mutual information: Criteria of maxdependency, max-relevance, and min-redundancy. *IEEE Transactions on Pattern Analysis and Machine Intelligence*, 27(8):1226–1238.
- Tsai, C.-J., Lee, C.-I., and Yang, W.-P. (2008). A discretization algorithm based on class-attribute contingency coefficient. *Inf. Sci.*, 178:714–731.
- Witten, I. and Frank, E. (2005). Data Mining: Practical Machine Learning Tools and Techniques. Elsevier, Morgan Kauffmann, 2nd edition.
- Yu, L., Liu, H., and Guyon, I. (2004). Efficient feature selection via analysis of relevance and redundancy. *Journal of Machine Learning Research*, 5:1205–1224.
- Zhu, Q., Lin, L., Shyu, M., and Chen, S. (2011). Effective supervised discretization for classification based on correlation maximization. In *IEEE International Conference on Information Reuse and Integration (IRI)*, pages 390–395, Las Vegas, Nevada, USA.