Feature Selection Based on the Rough Set Theory and EM Clustering Algorithm

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Abstract. We study the Rough Set theory as a method of feature selection based on tolerant classes that extends the existing equivalent classes. The determination of initial tolerant classes is a challenging and important task for accurate feature selection and classification. In this paper the EM clustering algorithm is applied to determine similar objects. This method generates fewer features with either a higher or the same accuracy compared with two existing methods, i.e., Fuzzy Rough Feature Selection and Tolerance-based Feature Selection, on a number of benchmarks from the UCI repository.

1 Introduction

The problem of reducing dimensionality has been investigated for a long time in a wide range of fields, e.g., statistics, pattern recognition, machine learning, and knowledge discovery. In order to reduce the input dimensionality, there exist two main approaches, i.e., feature extraction and feature selection (FS). Feature extraction maps the primitive feature space into a new space with a lower dimensionality. Two of the most popular feature extraction approaches include Principal Components Analysis [12], and Partial Least Squares [2]. There are numerous applications of feature extraction in the literature, such as image processing [9], visualization[28], and signal processing [20]. In contrast, the FS approach chooses the most informative features from the original features according to a selection method, e.g., t -statistic [16], f -statistic [14], correlation [33], separability correlation measure [7], or information gain [31]. The irrelevant and redundant features in the dataset lead to slow learning and low accuracy. Finding the subset of features that are enough informative is NP complete. Some heuristic algorithms are proposed to search through the feature space. The selected subset can be evaluated from some issues, such as the complexity of the learning algorithm and the accuracy.

The Rough Set (RS) theory can be used as a tool to reduce the input dimensionality and to deal with vagueness and uncertainty in datasets.

The reduction of attributes is based on data dependencies. The RS theory partitions a dataset into some equivalent (indiscernibility) classes, and approximates uncertain and vague concepts based on the partitions. The measure of dependency is calculated by a function of the approximations. The dependency measure is employed as a heuristic to guide the FS process. In order to obtain a significant measure, proper approximations of the concepts are required. Hence, the initial partitions play an important rule. Given a discrete dataset, it is possible to find the indiscernibility classes; however, in case of datasets with real-valued attributes, it is impossible to say whether two objects are the same, or to what extent they are the same, using the indiscernibility relation. A number of research groups [6, 19, 25, 26, 27, 29] extended the RS theory using the tolerant or similarity relation (termed tolerance-based Rough Set). The similarity measure between two objects is delineated by a distance function of all attributes. Two objects are considered to be similar when their similarity measure exceeds a similarity threshold value. Finding the best threshold boundary is both important and challenging. [13] used genetic algorithms to find the best similarity threshold. [8, 10, 21, 22, 24] used fuzzy similarity to cope with real-valued attributes. In this paper we use Expectation-Maximization (EM) [3, 5, 15, 23, 32, 34] clustering algorithm to determine the tolerance classes. The EM algorithm is a general statistical method for finding the maximum likelihood estimations of parameters in probabilistic models. In particular it can be applied in clustering problems. The EM algorithm allows for overlapping clusters and it is robust to noise and to highly skewed data.

The paper is organized as follows. Section 2 summarizes basics of the RS theory. A brief overview of the EM algorithm is represented in section 3. In Section 4, the proposed method of feature selection using the RS theory and EM clustering algorithm is outlined. Section 5 shows the potential of the proposed method on some real datasets. We discuss our results and draw some conclusions in the final section.

2 Basics of the Rough Set Theory

An information decision table is a table where each row presents a situation, event or sample and each column presents a feature or attribute of samples. The RS theory [18] is the approximation of uncertain concepts via the two lower and upper approximation sets. The lower approximation presents those objects that can exactly be classified but the upper approximation is a description of objects that possibly classified. Some basic definitions in the RS theory are considered.

Let T(U, A, C, D) be a decision table, where U is a universe of objects, A is a set of primitive features, C is a set of conditional attribute, D is a decision attribute or class label, and $C, D \subseteq A$.

Indiscernibility relation:

For an arbitrary set $P \subseteq A$, an indiscernibility relation is defined as follows,

$$IND(P) = \{(x, y) \in U \times U : \forall a \in P, a(x) = a(y)\}$$
(1)

An indiscernibility relation partitions the universe U into disjoint subsets. Let U/IND(P) denote the family of all equivalent classes generated by IND(P). The equivalence classes U/IND(C) and U/IND(D) will be called condition and decision equivalent classes, respectively.

Given a subset $X \subseteq U$, it may be infeasible to describe X with a combination of the equivalent classes. It means that a rigorous representation of X with the available information is impossible. The Rough Set Theory describes X using the lower and upper approximation sets as follow, **Approximations:**

If $P \subseteq C$ and $X \subseteq U$ then the lower and upper approximations of X, with respect to P, are respectively defined as follow,

$$\underline{P}X = \{x \in U : [x]_{IND(P)} \subseteq X\}$$
(2)

$$\overline{P}X = \{x \in U : [x]_{IND(P)} \cap X \neq \phi\}$$
(3)

where

$$[x]_{IND(P)} = \{ y \in U : a(y) = a(x), \forall a \in P \}$$
(4)

is the equivalence class of x in U/IND(P).

Positive region:

A *P*-positive region of *D* is a set of all objects from the universe *U* which can be classified with certainty to one class of U/IND(D) employing attributes from *P*,

$$POS_P(D) = \bigcup_{x \in U/IND(D)} \underline{P}X$$
(5)

Dependency of attributes:

A dependency of D on P is defined as,

$$\gamma_p(D) = \frac{|POS_P(D)|}{|U|}.$$
(6)

where |A| is the cardinality of a set A.

A feature $a \in C$ is dispensable in P, if $\gamma_p(D) = \gamma_{p-a}(D)$; otherwise a is an indispensable attribute in P with respect to D. An arbitrary set $B \subseteq C$ is called independent if all its attributes are indispensable. **Reduct:**

From these definitions a reduct set of features can be defined as follows, a set of features $R \subseteq C$ is called the reduct of C, if R is independent and $POS_R(D) = POS_C(D)$. In other words, the reduct is a set of attributes that conserves the partitions generated by C. It means that a reduct is the smallest subset of features that generates the same classification of objects in the universe as the whole set of features. In other words, features that do not belong to the reduct set are redundant with regard to classification of objects in the universe.

In [4] the QUICKREDUCT algorithm for determining the reduct set is proposed. It is a heuristic algorithm that avoids exhaustively generating all possible subsets. The greedy algorithm starts with an empty set and in each iteration adds the attribute that results in the greatest increase in the rough set dependency metric to the reduct set.

3 EM Algorithm

The mixture model is an effective representation of the probability density function and consists of k component density functions. The objective of a mixture model is to fit the density functions to a given dataset to approximate the data distribution. Let D be a dataset with m objects and d attributes and $\mathbf{x} \in D$ be an object in the dataset. The mixture model probability density function, evaluated at \mathbf{x} , is defined as follows,

$$pr(\mathbf{x}) = \sum_{l=1}^{k} W_l.pr(\mathbf{x}|\phi_l)$$
(7)

where

- W_l is the fraction of data points belonging to the cluster l, and $\sum_{l=1}^{k} W_l = 1$.
- $\overline{pr}(\mathbf{x}|\phi_l)$ is the cluster or component distribution models the records of the *l*-th cluster.
- ϕ_l is the model parameters of density function of cluster *l*. In case of Gaussian distribution, ϕ_l is the mean (μ_l) and covariance matrix (Σ_l) .

In this work a Gaussian distribution is used. The EM algorithm is used to determine the value of mean (μ_l) , covariance matrix (Σ_l) , and sampling probability (W_l) for each cluster. The algorithm is as follows,

1. **E Step:** For each object $\mathbf{x} \in D$, compute the membership probability of \mathbf{x} in each cluster $l = 1 \cdots k$ at iteration j:

$$pr(l|\mathbf{x}) = \frac{W_l^j \cdot pr^j(\mathbf{x}|\mu_l^j, \Sigma_l^j)}{pr^j(\mathbf{x})}$$
(8)

2. M Step: Update mixture model parameters for each cluster $l = 1, 2, \dots, k$:

$$W_l^{j+1} = \frac{1}{N} \sum_{\mathbf{x} \in D} pr(l|\mathbf{x})$$
(9)

$$\mu^{j+1,l} = \frac{\sum_{\mathbf{x}\in D} \mathbf{x}.pr(l|\mathbf{x})}{\sum_{\mathbf{x}\in D} pr(l|\mathbf{x})}$$
(10)

$$\Sigma^{j+1,l} = \frac{\sum_{\mathbf{x}\in D} pr(l|\mathbf{x})(\mathbf{x}-\mu_{j+1,l})(\mathbf{x}-\mu_{j+1,l})^T}{\sum_{\mathbf{x}\in D} pr(l|\mathbf{x})}$$
(11)

3. If $|L^j - L^{j+1}| \le \epsilon$, stop. Else set j = j + 1 and go to 1. L^j is the log likelihood of the mixture model at iteration j

$$L^{j} = \sum_{\mathbf{x}\in D} log(pr^{j}(\mathbf{x})) = \sum_{\mathbf{x}\in D} log(\sum_{l=1}^{k} W_{l}^{j}.pr^{j}(\mathbf{x}|\mu_{l}^{j},\Sigma_{l}^{j}))$$
(12)

4 Proposed Method

In the proposed method, each cluster represents a tolerance class. The tolerance classes that are generated by the EM clustering algorithm for an object x are defined as:

$$Clus_P(x) = \{Y \in U \mid x, and Y belongs to the same cluster\}$$
 (13)

In a similar way to the original RS theory, the lower and upper approximations are then delineated as follow,

$$\underline{P}X = \{x \in U : Clus_P(x) \subseteq X\}$$
(14)

$$\overline{P}X = \{x \in U : Clus_P(x) \cap X \neq \phi\}$$
(15)

Based on this, the positive region and dependency functions can respectively be defined as follow,

$$POS_P(D) = \bigcup_{x \in U/IND(D)} \underline{P}X,$$
(16)

$$\dot{\gamma}_P(D) = \frac{|POS_P(D)|}{|U|} \tag{17}$$

Following the above definitions, a feature selection algorithm can be constructed that uses the tolerance-based degree of dependency, $\gamma_P(D)$, to evaluate the significance of feature subsets. The proposed FS algorithm are presented in Figure 1.

EM-CLUSTERING-REDUCT(C, D). Inputs : C, the set of all conditional attributes; D, the set of decision attributes; Output : R, the Reduct Set

(1) $R = \phi$ (2) $\dot{\gamma}_{best} = 0$ (3) **do** $\acute{\gamma}_{tmp}=\acute{\gamma}_{best}$ (4)(5)T = Rfor x in (C-R)(6)if $\dot{\gamma}_{R\cup\{x\}}(D) > \dot{\gamma}_T(D)$ $T = R \cup \{x\}$ (7)(8)(9) $\dot{\gamma}_{best} = \dot{\gamma}_T(D)$ R = T(10)(11) **until** $\dot{\gamma}_{best} == \dot{\gamma}_{tmp}$ (12) return R

Fig. 1. EM Clustering QuickReduct

Table 1. Example Table

Object	a	b	с	q
1	-0.4	-0.3	-0.5	0
2	-0.4	0.2	-0.1	1
3	-0.3	-0.4	-0.3	0
4	0.3	-0.3	0	1
5	0.3	-0.3	0	1
6	0.2	0	0	0

4.2 An illustrative example

In this section, a simple example is used to demonstrate the procedure of the proposed method (see Table 1). There are three continuous conditional attributes and a crisp-valued class attribute in the dataset. In this example, the number of clusters is set to 3.

The greedy algorithm starts with an empty reduct set. It checks each attribute separately and chooses the attribute that has the highest dependency degree. In this example the attribute c is chosen with the dependency degree of 0.33. Then the attribute c is added to the reduct set.

$$\begin{split} U/clust_{\{q\}} &= \{\{1,3,6\},\{2,4,5\}\}\\ U/clus_{\{a\}} &= \{\{3\},\{4,5,6\},\{1,2\}\}\\ \dot{\gamma}_a &= \frac{|\{3\}|}{|\{1,2,3,4,5,6\}|} = \frac{1}{6} = 0.17\\ U/clust_{\{b\}} &= \{\{1,3,4,5\},\{6\},\{2\}\}\\ \dot{\gamma}_b &= \frac{|\{2,6\}|}{|\{1,2,3,4,5,6\}|} = \frac{2}{6} = 0.33\\ U/clust_{\{c\}} &= \{\{3\},\{2,4,5,6\},\{1\}\}\\ \dot{\gamma}_c &= \frac{|\{1,3\}|}{|\{1,2,3,4,5,6\}|} = \frac{2}{6} = 0.33\\ R \leftarrow \{c\} \end{split}$$

The hill climbing forward selection algorithm chooses other attributes in the reduct set as follow,

$$\begin{split} U/clust_{\{a,c\}} &= \{\{1,3\},\{4,5,6\},\{2\}\}\\ \dot{\gamma}_{a,c} &= \frac{|\{1,2,3\}|}{|\{1,2,3,4,5,6\}|} = \frac{3}{6} = 0.5\\ U/clust_{\{b,c\}} &= \{\{1,3\},\{4,5\},\{2,6\}\}\\ \dot{\gamma}_{b,c} &= \frac{|\{1,2,4,5\}|}{|\{1,2,3,4,5,6\}|} = \frac{4}{6} = 0.67 \end{split}$$

Dataset Objects Features Reduct Size FRFS^a Tol.^b EMRS^c Glass 214109 7 5Heart 2701411103 Ionosphere 230351110 $\mathbf{5}$ Iris 1505 54 4 Water2 390 39 11 3 8 Wine 178 14 10 8 8

 Table 2. Reduct Size For FRFS, Tolerance, and EM Clustering Methods

^a FRFS : Fuzzy Rough Set Feature Selection [11]. ^b Tol. : Tolerance-based Feature Selection [11].

^c EMRS : The proposed method, i.e., Feature Selection using the RS theory and EM algorithm

$$R \leftarrow \{b, c\}$$

 $U/clust_{\{a,b,c\}}: \{\{1,3\},\{4,5,6\},\{2\}\}$

$$\dot{\gamma}_{a,b,c} = \frac{|\{1,2,3\}|}{|\{1,2,3,4,5,6\}|} = \frac{3}{6} = 0.5$$

Finally, it returns $\{b, c\}$ as the reduct set which has the same size as the reduct set provided by the Fuzzy Rough Feature Selection (FRFS) and tolerance based FS methods in [11].

5 Simulation Result

In order to evaluate the proposed method, we applied it to a number of real datasets from the UCI repository [1] in Table 2. The EM clustering algorithm from the Weka software [30] was chosen where the number of clusters was selected empirically. The obtained reducts are evaluated via the accuracy of classification. J48, JRIP, and PART classifier in the Weka [30] are chosen as the classifier algorithms.

The obtained accuracies are compared with the accuracy of the FRFS and Tolerance-based FS in [11]. Table 3 shows the average classification accuracy of 10-fold cross validation as a percentage. The classification algorithms are performed on the original dataset and reduced datasets were obtained by the feature selection algorithms, i.e., the FRFS [11], the Tolerance-based FS [11], and the proposed method.

It is evident from Table 2 that the proposed method generated fewer features compared with the two other FS methods. For the J48 classifier, the clustering based FS improved the average accuracy of the unreduced datasets except for the water2 dataset. The proposed method either unchanged or improved upon the performance of the reduced datasets with the other two FS algorithms in all but in the Ionosphere dataset. For the

Table 3. Classification Accuracies(%) For Unreduced, FRFS, Tolerance, and Clustering Methods.

CA^{a}	J48				JRIP			PART				
FSb	$\operatorname{Original}^{\mathbf{C}}$	FRFS^d	$\mathrm{Tol.}^{\mathbf{e}}$	$\mathrm{EMRS}^{\mathrm{f}}$	$\operatorname{Original}^{\mathbf{C}}$	FRFS^d	$\mathrm{Tol.}^{\mathbf{e}}$	$\mathrm{EMRS}^{\mathrm{f}}$	$\operatorname{Original}^{c}$	FRFS^d	$\mathrm{Tol.}^{\mathbf{e}}$	$\mathrm{EMRS}^{\mathrm{f}}$
Glass	67.29	69.63	69.16	69.16	69.16	67.76	67.76	69.16	67.76	68.22	69.62	69.16
Heart	76.67	78.89	80.37	79.59	79.63	81.85	82.59	79.59	73.33	78.52	80.37	79.59
Ionosphere	87.83	91.30	87.39	88.32	86.96	86.52	86.96	86.61	88.26	91.30	86.52	90.03
Iris	96.00	96.00	96.00	96.00	95.33	95.33	94.67	95.33	94.00	94.00	95.33	95.33
Water2	83.33	80.26	81.79	81.77	81.03	80.51	82.31	81.57	85.64	82.56	81.28	82.34
Wine	94.38	92.14	94.94	94.94	91.57	90.45	94.38	92.7	93.82	93.82	94.38	94.38
^a CA : Classification Algorithm.												

⁶ CA : Classification Algorithm.
 ⁶ FS : Feature Selection Algorithm used for each Classification Algorithm.
 ⁶ Original : Original dataset.

^d FRFS : Fuzzy Rough Set Feature Selection [11]. ^e Tol. : Tolerance-based Feature Selection [11].

f EMRS : The proposed method, i.e., Feature Selection using the RS theory and EM algorithm

JRip classifier, the proposed method maintained the average accuracy of the unreduced datasets in all. It either improved or maintained the performance of the reduced dataset with the other two FS algorithms in all but two cases. For PART, the proposed method improved the average accuracy of unreduced datasets in all except the water2 dataset. It has the same behavior as the other two FS methods.

Overall, the proposed algorithm produced a smaller number of attributes compared to the other two FS algorithms and the average accuracy of classifiers is improved or in a few instances remains unchanged. For example, in the water2 dataset the proposed method chose 3 features among 39 features whereas the FRFS chose 10 and the Tolerance-based FS method chose 8 features. In addition, the proposed method has a similar average accuracy compared with the other two approaches.

6 Conclusion

In this work the EM clustering algorithm was applied to deal with the problem of determining initial tolerant classes to obtain a significant classification accuracy. Through some experiments, it was concluded that the proposed method generated a smaller size of feature sets in all datasets compared with the FRFS [11] and tolerance-based FS methods [11]. Beside that, the proposed method either improved or unchanged the average accuracy in all except a few datasets. For future work, an improvement of searching algorithm for finding the reduct set with the new definition of approximations is required. In Addition, an evaluation of the proposed method through experimental comparisons with the other methods in the literature is recommended.

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