AN ALGORITHM FOR DECISION RULES AGGREGATION

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Abstract: Decision trees and decision rules are usually applied for the classification problems in which legibility and possibility of interpretation of the obtained data model is important as well as good classification abilities. Beside trees, rules are the most frequently used knowledge representation applied by knowledge discovery algorithms. Rules generated by traditional algorithms use conjunction of simple conditions, each dividing input space by a hyperplane parallel to one of the hyperplanes of the coordinate system. There are problems for which such an approach results in a huge set of rules that poorly models real dependencies in data, is susceptible for overtfitting and hard to understand by human. Generating decision rules containing more complicated conditions may improve quality and interpretability of a rule set. In this paper an algorithm taking a set of traditional rules and aggregating them in order to obtain a smaller set of more complex rules has been presented. As procedure uses convex hulls, it has been called Convex Hull-Based Iterative Aggregation Algorithm.

1 INTRODUCTION

Rules-based data models can be applied for the classification (Furnkranz, 1999), description (Fayad et al., 1996), or both the aims simultaneously. In the case of description or both description and classification, the possibility of interpretation of created rules-based data descriptions, thus the ability to understand and to use dependencies represented by rules, is their significant feature. Due to the specificity of the language of classification rules, defining simple rules-based descriptions is impossible for many problems (Fig. 1). It turned out that if the language of rules representation is slightly more complicated, simple and still intuitive descriptions can be found for many of the problems (Fig. 2). Therefore a lot of methods search for rules which premises contain linear combinations of conditional attributes. This enables to get so-called oblique descriptors. The majority of the methods make a decision trees induction (oblique decision trees) (Murthy et al., 1994), (Bennett, Blue, 1997), and then transform determined trees into rule sets. The algorithm ARED (Seunghyun, Ras, 2008) makes the rules induction with oblique descriptors. However the efficiency of the algorithm was not verified on a bigger number of datasets.

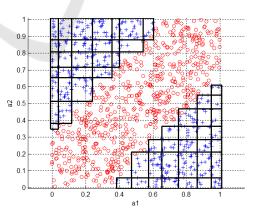


Figure 1: Example classification problem. Rules covering decision regions have been induced using JRip which is an implementation of the RIPPER algorithm (Cohen, 1995) contained in Weka software (Hall et al., 2009). One can see that the simple decision rule language is insufficient to model the data properly.

Recently, an increasing number of hybrid rules induction methods has been observed. Joining rules induction algorithms with support vector machines (SVM) is the most popular approach (Barakat, Bradley, 2006), (Martens et al., 2009), (Nunez et al., 2008). In the papers, SVM are usually used to concentrate training examples on boundaries of decision regions or to separate coherent regions in the features space, and then describe them by rules with higher dimensional curves (e.g. ellipsoids) in their premises.

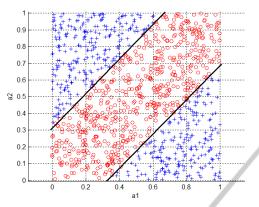


Figure 2: Using a more complicated description language allows to model classification problem nicely.

Each of the mentioned methods has advantages as well as disadvantages. The fact that diagonal or nonlinear descriptors occurring in rules apply all conditional attributes without offering any pruning strategies is the disadvantage in the majority of the methods. The pruning is meant as simplifying multidimensional expressions by eliminating conditional attributes forming them. Possibility of interpretation of unpruned rules is then strongly limited.

In the paper, a method of aggregating traditional classification rules obtained by any standard rule induction algorithm is presented. The procedure tries to join rules sequentially, by twos, finding a set of hyperplanes which bound a region covered by rules being joined.

The presented method is a generalization of rules joining algorithms which merge rules without changing their representation language (Sikora, 2005), (Latkowski, Mikołajczyk, 2004) or operate on a very specific representation (Pindur et al., 2004).

In the next part of the paper, an algorithm for rules aggregation and rules postprocessing, as well as the results of experiments are presented. The last chapter includes the summary of obtained results and directions of further works.

2 BASIC NOTIONS

Some basic notions used later in the paper have been introduced here.

Let's assume we have a set of objects O which can be divided into c disjoint classes labelled by elements of a set $\{1, 2, ..., c\}$. Information about assignment of a given object $o \in O$ to a decision class is included in a value of a decision variable y:

$$y: 0 \to \{1, 2, \dots, c\}.$$
 (1)

Let's additionally define a *d*-element set of attributes $A = \{a_1, a_2, ..., a_d\}$. To each object $o \in O$ we can assign a *d*-dimensional vector $x = description(o), x \in \mathbb{R}^d$ containing values of these attributes. One can see, that it has been assumed that all the attributes from *A* are continuous. This simplification is because the current version of the algorithm ignores categorical features in the aggregation process. However, a generalization for categorical attributes is possible and has been described later on.

A training set can be defined as follows:

$$T = (0, A \cup \{y\}).$$
 (2)

A classification task consists in finding a function ϕ which approximates the function y on the basis of a training set T (which is equivalent to generalising a knowledge represented by a training dataset).

Rules-based classifiers can be distinguished by classification strategies they exploit. Described algorithm is based on a rules hierarchy. This is because RIPPER algorithm (Cohen, 1995) which served to generate input rules uses this scheme as well. The idea is that rules are ordered. During the classification process the first rule covering an object being currently checked is picked. Hence, reordering rules affects classification performance (which is not the case in different aggregation strategies like voting rules).

A decision rule can be defined as follows:

$$r: \varphi \to \psi,$$
 (3)

where φ is a premise consisting of a conjunction of conditions (descriptors), and ψ is a conclusion (a class label). Majority of decision rules induction algorithms generates simple descriptors like $a_i \ge v_i$ or $a_i \le v_i$. Such conditions divide an input space into two parts by a hyperplane parallel to one of hyperplanes of the coordinate system. Aggregation algorithm covered in this paper generates oblique descriptors, which are linear combinations of attributes:

$$a_1a_1 + h_2a_2 + \dots + h_da_d + v \ge 0.$$
 (4)

Such descriptor divides a feature space by some hyperplane, not necessarily parallel to coordinate system hyperplanes. It is assumed, that inequalities in all descriptors have the same direction (one can always multiply expression by -1 to obtain this).

Several measures that reflect a quality of decision rules of the form (3) can be computed (Furnkranz, Flach, 2005), (Guillet, Hamilton, 2007), (Sikora, 2010). If the rule r is denoted as $\varphi \rightarrow \psi$, then $n_{\varphi} = n_{\varphi \psi} + n_{\varphi \to \psi} = |O_{\varphi}|$ is the number of objects that recognize the rule; $n_{\neg \varphi} = n_{\neg \varphi \psi} + n_{\neg \varphi \neg \psi} = |O_{\neg \varphi}|$ is the number of objects that do not recognize the rule; n_{ψ} = $n_{\omega \psi} + n_{\neg \omega \psi} = |O_{\psi}|$ is the number of objects that belong to the decision class described by the rule; $n_{\neg w}$ $n_{\varphi \to \psi} + n_{\neg \varphi \to \psi} = |O_{\neg \psi}|$ is the number of objects that do not belong to the decision class described by the rule; $n_{\varphi\psi} = |O_{\varphi} \cap O_{\psi}|$ is the number of objects that support the rule. Values $n_{\varphi \rightarrow \psi}$, $n_{\neg \varphi \rightarrow \psi}$, $n_{\neg \varphi \rightarrow \psi}$ are calculated similarly as $n_{\varphi\psi}$. It can be noticed that for the inequalities $l \leq n_{\omega w} \leq |O_w|$, any rule $\varphi \rightarrow \psi$ $0 \le n_{\varphi \neg \psi} \le |O_{\neg \psi}|$ hold. Hence, a quality measure is a function of two variables $n_{\phi\psi}$ and $n_{\phi\neg\psi}$, and can be defined as follows (Sikora, 2006):

$$q(\varphi \rightarrow \psi): \{1, \dots, |O_{\psi}|\} \times \{0, \dots, |O_{\neg \psi}|\} \rightarrow \boldsymbol{R}.$$
 (5)

Two basic quality measures are accuracy and coverage:

$$acc(\phi \rightarrow \psi) = n_{\phi\psi}/n_{\phi},$$

$$cov(\phi \rightarrow \psi) = n_{\phi\psi}/n_{\psi}.$$
(6)
(7)

The accuracy measure and two other measures are used for evaluation of joined rules in the aggregation algorithm. The first measure called RSS is empirical and enables to evaluate the accuracy and coverage of a rule simultaneously, taking into consideration examples distribution among the decision class indicated by the rule and the other decision classes.

$$q^{\text{RSS}}(\varphi \to \psi) = \frac{n_{\varphi\psi}}{n_{\varphi\psi} + n_{\varphi\psi}} + \frac{n_{\varphi\psi\psi}}{n_{\varphi\psi\psi} + n_{\varphi\psi\psi}} - 1.$$
(8)

Making an analysis of the formula (8) it can be noticed that the measure proposes the method of rule evaluation analogous to the method of classifiers sensitivity (first component of the sum) and specificity (second component of the sum) evaluation. The measure takes values from the interval [-1,1] and values equal to zero are achieved when a rule has the same accuracy as it implies from the positive and negative examples distribution in a training set.

The other measure used in a rule evaluation process is Laplace estimate (9).

$$q^{Laplace}(\varphi \to \psi) = \frac{n_{\varphi\psi} + 1}{n_{\varphi\psi} + n_{\varphi\neg\psi} + c}$$
(9)

The basic idea of the Laplace estimate is to assume that each rule covers a certain number of examples a priori. The estimate computes the accuracy, but starts to count covered positive or negative examples at a number greater than 0. The positive coverage of a rule is initialized with 1, while the negative coverage of a rule is initialized with a number of decision classes c.

3 CONVEX HULL-BASED ITERATIVE AGGREAGTION ALGORITHM

the following paragraph an algorithm for In aggregation decision rules has been presented. It has been called Convex Hull-Based Iterative Aggregation Algorithm. Iterative, because it sequentially tries to merge decision rules in order to model the problem more properly than the input rule set. Convex Hull-Based, because a procedure of aggregating two particular decision rules uses convex hulls. On the Fig. 3 one can see a full flowchart of the algorithm. In the sections below all steps of the procedure are covered in details. Example dataset from the Fig. 1 has been used as an illustration. One must keep in mind that it is assumed, that input rule set R contains rules corresponding to a single decision class.

3.1 Hyperrectangular Rules

The idea of the aggregation procedure is based on the assumption that a decision rule can be considered as some convex area in a feature space. The algorithm tries to merge these areas into larger ones. However, rules generated by traditional induction algorithms rarely determine convex areas to obtain this, a rule should by bounded from both sides in each dimension forming a hyperrectangle:

$$a_1 \in \langle l_1, h_1 \rangle \land \dots \land a_d \in \langle l_d, h_d \rangle \to \psi.$$
 (10)

This is why the first step of the algorithm consists in bringing an input rules into a hyperrectangular form. This can be done by adding missing descriptors with coordinates of the extreme points from a training set. It will not affect classification results of unseen data because these synthetically added descriptors will be removed after aggregation procedure has finished. In addition, hyperrectangular rules have a very nice feature - one can easily find vertices of the corresponding hyperrectangles. This is a crucial issue because in the following steps algorithm uses these vertices in the merging procedure. The implication of the proposed approach is that a rule rcan be represented in a dual way:

- as a set of hyperplanes H(r),
- as a set of vertices V(r).

Keeping integrity between the hyperplane representation used directly for classification and the vertex representation which takes part in the aggregation procedure is a very important task the algorithm must handle.

3.2 Aggregation Loop

As our aim is to reduce a number of rules in the entire rule set meeting some classification accuracy conditions, it must be decided which rules in which order should be picked for the aggregation procedure. The simplest approach is to use an exhaustive paradigm to check all possible rule permutations and pick the best one. However, the brute-force method is computationally too expensive to be used in practice. This is why some greedy heuristics for searching aggregation candidates has been proposed.

At the beginning we take the first rule as the current one and try to merge it with following rules (let's call those rules aggregation partners) creating a candidates for new rules. Method of joining two particular decision rules has been explained in the following section. For each candidate we check if it meets quality criterions.

Quality control consists of two elements. At first algorithm checks how quality of the rule candidate is related to the better rule from the pair being aggregated. If it is below some threshold, the candidate rejected. Parameter indicating is maximum drop of the quality is called ruleQualityDrop and is expressed in percentages. However, checking only this condition may result in an accumulation of the quality drops and big classification error of the final rule set. This is why an additional criterion has been introduced. Algorithm checks how aggregation of two particular rules affects classification accuracy obtained by the entire rule set. If an accuracy decrease exceeds rulesetAccuracyDrop parameter (which is given in percentages as well), the candidate is rejected. One must remember, that these two parameters are evaluated on a training set. Adjusting them gives full control on how the algorithm works. For example,

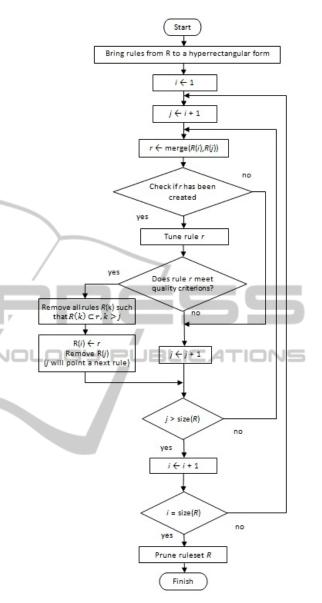


Figure 3: Flowchart of the Convex Hull-Based Iterative Aggregation Algorithm.

one can force the algorithm to operate in the way that classification accuracy obtained by a final rule set does not decrease (on a training set).

If quality criterions are fulfilled, a rule candidate is accepted. We replace the current rule with the new one, and remove the partner from the rule set. The algorithm also checks if the new rule covers some other rules which follow the partner and removes them if possible. There is no need to check rules preceding the partner because they are the ones that hasn't been aggregated with the current rule (so there is no possibility that they are covered by the new rule). After all partners have been checked, we change a current rule for the next one. Procedure stops when the current rule is the last one from the rule set.

3.3 Merging Two Decision Rules

This section covers in details how the algorithm merges two particular decision rules. Let's assume there are two rules r_i and r_j we would like to aggregate. Below steps of the basic aggregation procedure have been described. The modified version with some improvements has been introduced later.

- 1. Create a new rule r such that $V(r) = V(r_i) \cup V(r_j)$.
- 2. Calculate a *d*-dimensional convex hull of points belonging to the V(r) using Qhull algorithm (Barber et al., 1996). This algorithm returns a set F(r) describing all the facets of a convex hull. Each facet is represented by a set of *d* indices pointing some vertices from V(r).
- 3. For each face from F(r) calculate a corresponding hyperplane equation. As we know vertices belonging to each face, this can be done easily by solving a system of linear equations. Obtained hyperplanes are stored in H(r) set.

Important advantage of the method is that it can be generalized for categorical attributes easily. Merging descriptors $a_i \in X$ and $a_i \in Y$ simply produces $a_i \in X \cup Y$. Another feature of the procedure is that it is based only on rule vertices and does not use a training set.

However, the approach described above has its disadvantages. Most important one is that a number of vertices representing a rule grows exponentially with a number of dimensions. As hyperrectangular rules use all attributes in their premises, handling high dimensional feature spaces is computationally very expensive. This is why, a hull should be calculated only with respect to dimensions which are really used in aggregated rules. Therefore, for each input rule r, before bringing it to a hyperrectangular form, we store attributes present in a premise in a set A(r). Additional improvement is to limit maximum number of dimensions that can appear in a rule being created. There is an algorithm parameter called maxDim. If number of attributes in a rule candidate exceeds this parameter, a rule is not created. This allows user to control the complexity of obtained

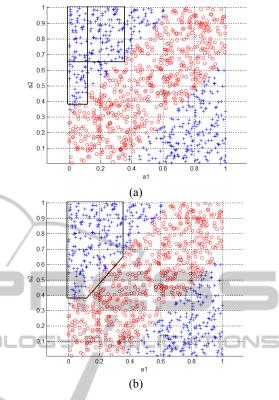


Figure 4: Two hyperrectangular rules separately (a) and after aggregation (b).

descriptors (for example, he can limit number of dimensions to 3 in order to visualise hyperplanes) or to speed-up the algorithm. The modified aggregation procedure of rules r_i and r_j is as follows:

- 1. Create a new rule *r* such that $V(r) = V(r_i) \cup V(r_j)$ and $A(r) = A(r_i) \cup A(r_j), |A(r)| = k$. If k > maxDim, reject *r* immediately and skip the following steps.
- 2. Calculate a k-dimensional convex hull of points from V(r) with respect to the attributes belonging to A(r). Following actions are the same as in the previous description.

The result of aggregation of two example rules has been presented on the Fig. 4.

3.4 Rule Tuning

One can see on the flowchart that just after creation of a new rule candidate there is a step called rule tuning. It comes out that sometimes a rule candidate may be further improved. This is very important from the point of view of the algorithm because quality influences if an aggregated rule is accepted. Hence, each candidate is tuned to obtain as good result as possible.

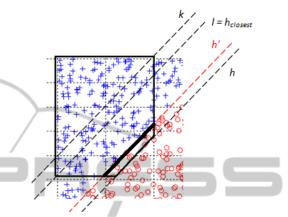
Rule tuning procedure has two phases. The first stage of tuning consists in joining coplanar facets and removing duplicated hyperplanes. As we mentioned before, in k-dimensional space each facet of a convex hull calculated by Qhull has k vertices. This means that, for example, in 3-dimensional space, we obtain triangular facets. Such approach may result in coplanar facets. For example cube contains 6 square facets but Qhull returns 12 triangle ones. In fact, as our input rules have a hyperrectangular form, this situation will happen very often. More the dimensions, more coplanar facets appear. This is why the algorithm must merge coplanar facets and remove duplicated hyperplane equations.

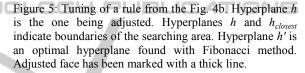
The second stage of tuning consists in adjusting hyperplanes equations to get the maximum quality of an aggregated rule. One can see on the Fig. 4 that the oblique line is not optimally situated (if we move it slightly upwards it will cover same number of positive examples and less negative examples). Hence, we introduce a technique of hyperplane adjusting. For simplicity it is assumed that only hyperplane translations are possible (no rotations are performed). Below we described adjustment steps for a rule r created as the result of aggregation of rules r_i and r_j .

- 1. Find a hyperplane to adjust $h \in H(r)$ such that $h \notin H(r_i) \land h \notin H(r_j)$ (there is no need to improve a boundary hyperplane or a hyperplane that could have already been improved in one of previous iterations).
- 2. Find a face f corresponding to the hyperplane h and all vertices from V(r) adjacent to the face f. A vertex is adjacent to the face f if it belongs to some face g adjacent to f and it does not belong to f.
- 3. Calculate equations of hyperplanes parallel to *h* going through vertices adjacent to *f* and choose the one closest to *h* (let's call it $h_{closest}$). Original hyperplane *h* and the closest one determine bounds of the searching area. This is to assure that no face vanishes in the adjustment procedure.
- 4. Use Fibonacci search technique (Ferguson, 1960) to find an optimal hyperplane h' parallel to h lying in the area bounded by h and $h_{closest}$ (as these hyperplanes are parallel, Fibonacci is used to find an optimal value of a free term). Assumed quality measure is used as a criterion function in searching.

Replace h with h' in H(r), calculate new positions of vertices forming face f using h' equation and equations of adjacent faces. This can be easily done by solving a system of linear equations.

An example of the adjustment procedure has been shown on the Fig. 5.





3.5 Rule Set Pruning

Rule set pruning is done at the very end of the whole procedure after the aggregation loop has finished. This phase consists of actions that either decrease rules quality (and should not be done in a tuning phase preventing from a premature rejection of candidates) or remove some data from a rule description so further aggregations become impossible.

There are two stages of pruning, both are repeated for all rules in a rule set. First one consists in eliminating hyperplanes that correspond to the boundaries of the domain. As it has been said before, the first step of the Iterative Aggregation Algorithm is transforming input rules to the hyperrectangular form. This means that some hyperplanes had to be synthetically added. After the entire aggregation procedure has finished, we can remove them.

The second stage of pruning consists in removing other hyperplanes that have no significant influence on classification results. Algorithm just iterates through all the hyperplanes and checks how deleting affects classifier performance. If a classification accuracy decrease evaluated on a training set does not exceed *pruningAccuracyDrop* parameter (given in percentages), a hyperplane is removed. Below one can see a comparison between some rule before and after pruning procedure.

```
Before pruning:
(-a1+0.602 >= 0) and
(-a1+0.432*a2+0.196 >= 0) and
(-a1+0.63*a2+0.022 >= 0) and
(-0.975*a1+a2-0.277 >= 0) and
(-0.491*a1+a2-0.329 >= 0) and
(-0.991*a2+1 >= 0) and
(+a2-0.348 >= 0) and
(+a1+0.01 >= 0) => class = positive
After pruning:
(-0.975*a1+a2-0.277 >= 0) =>
class = positive
```

As one can see, the reduction of descriptors number is significant.

Actions described here cannot be done in the rule tuning phase. This is because the hyperplane adjustment procedure uses equations of adjacent hyperplanes to calculate new positions of face vertices. If one removed some hyperplanes from H(r), it would become impossible.

Final results of the algorithm for the example input rule set has been shown on the Fig 6.

4 EXPERIMENTS

In order to evaluate results of the algorithm some experiments have been performed. Benchmark datasets have been chosen mainly from the UCI Machine Learning Repository. The only exceptions

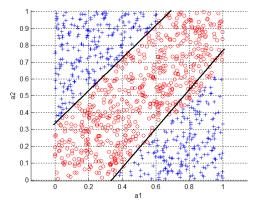


Figure 6: Final results of the Convex Hull-Based Iterative Aggregation Algorithm performed on the rule set presented on the Fig 1.

are synth2D (Fig. 1.) and synth3D datasets, which have been synthetically generated and sc503 which is the real dataset containing a microseismic hazard assessment in a coal mine. Input rules have been generated using JRip (Hall et al., 2009). Majority of the experiments have been run in 10-fold cross validation (only *segment* dataset has been tested in the train and test mode). Tests have been performed with four sets of parameters (from a very conservative to a very aggressive aggregation strategies) for the accuracy, RSS and Laplace quality measures. In each experiment it has been checked which measure leads to the best accuracy of an obtained rule set and the best reduction of a number of rules. Results are presented in the Table 1. One can see that accuracy is the best quality measure if we aim in the highest classification accuracy. If one would like to reduce a number of rules, he should pick RSS instead.

Table 1: Percentages indicate how often given quality measure leads to highest accuracy and highest rule reduction rate.

| | accuracy | RSS | Laplace |
|-------------------|----------|-----|---------|
| Highest accuracy | 41% | 29% | 30% |
| Highest reduction | 24% | 50% | 26% |
| rate | | | |

From all tested parameter sets we have chosen two which haves been called safe and aggressive aggregation strategies (see Table 2 for more detailed description). Tables 2a and 2b present results of classification for these parameter sets for accuracy and RSS quality measures respectively. This is due to fact that these measures gave the best results from the point of view of a classifier performance and a rule set reduction. Results for Laplace have been omitted, because they are located between accuracy and RSS. One can see that results for the safe strategy are similar for both quality measures. However, in the aggressive strategy RSS leads to lower accuracy and higher reduction of rule sets.

Below one can see a comparison between input and output rules for "R" decision class for the *balance-scale* data set. Aggressive parameter set has been used with hyperplane dimensionality limitation equal to 3. RSS has been set as a rule quality measure. Table 2: Results of classification for the example datasets. Input rule set, and rule sets obtained by the safe and aggressive aggregation strategies has been evaluated. Corresponding algorithm parameters are 20%-2%-1% and 20%-8%-2% (*ruleQualityDrop-rulsetAccuracyDrop-pruningAccuracyDrop*). To speed up the calculations maximum number of hyperplanes has been limited to 5. Accuracy (table a) and RSS (table b) have been used as quality measures.

| | Input rule set | | | Safe strategy | | | Aggressive strategy | | |
|---------------|-----------------|----------------|----------------|---------------|--------------|--------------|---------------------|--------------|--------------|
| Dataset | accuracy [%] | rules count | desc. count | Δ(acc) [%] | Δ(rc) [%] | Δ(dc) [%] | Δ(acc) [%] | Δ(rc) [%] | Δ(dc) [%] |
| vehicle | 67 | 16,7 | 42,1 | 2,3 | -7,8 | -32,5 | 0,6 | -9 | -37,5 |
| wine | 90,6 | 3,7 | 4,5 | 1,3 | 0 | -2,22 | 1,3 | 0 | -8,89 |
| glass | 65,4 | 7,2 | 14 | 1,3 | -2,8 | -7,86 | 1,5 | -4,2 | -12,1 |
| iris | 92,7 | 3,7 | 3,7 | 0,7 | -5,4 | -21,6 | 0,7 | -5,4 | -21,6 |
| australian | 84,1 | 4,7 | 8,7 | 1 | -4,3 | -14,9 | 1 | -36 | -67,8 |
| pima | 73,9 | 3,7 | 6,3 | -0,4 | -8,1 | -19 | -2 | -19 | -33,3 |
| balance-scale | 77,4 | 11,9 | 32,6 | 4,7 | -81 | -76,1 | 4,6 | -81 | -78,2 |
| ionosphere | 89,8 | 4,8 | 5,2 | 1 | -10 | -15,4 | -1 | -17 | -25 |
| heart-statlog | 74,4 | 4 | 6,1 | 3 | -40 | -37,7 | 3 | -43 | -42,6 |
| pendigits | 86,7 | 30,8 | 75,1 | -2 | -12 | -16,6 | -4 | -14 | -18,6 |
| ecoli | 83 | 9,5 | 16,7 | 0,7 | -6,3 | -17,4 | 0,4 | -7,4 | -24,6 |
| yeast | 57,8 | 16,7 | 39,3 | 1,5 | -14 | -36,9 | -0 | -22 | -47,8 |
| segment | 94,4 | 13 | 26 | -0,5 | -7,7 | -15,4 | -2 | -7,7 | -15,4 |
| synth2D | 95 | 10,9 | 19,8 | 2,3 | -72 | -86,4 | 1,7 | -72 | -87,9 |
| synth3D | 95,8 | 7,8 | 19,6 | -1 | -51 | -61,2 | -3 | -60 | -73,5 |
| sc503 | 90,2 | 4,1 | 6,2 | 0,6 | -22 | -30,6 | -1 | -27 | -40,3 |
| | · · · · · | | Average | 1,0 | -21,5 | -30,7 | 0,1 | -26,5 | -39,7 |

(a)

(b)

| | Input rule set | | | Safe strategy | | | Aggressive strategy | | |
|---------------|-----------------|-------|-------|---------------|--------------|--------------|---------------------|--------------|--------------|
| Dataset | accuracy [%] | rules | desc. | Δ(acc) [%] | Δ(rc) [%] | Δ(dc) [%] | Δ(acc) [%] | Δ(rc) [%] | Δ(dc) [%] |
| 1.1 | | count | count | | | | | | |
| vehicle | 67 | 16,7 | 42,1 | 2,6 | -16 | -37,3 | -3 | -23 | -48,5 |
| wine | 90,6 | 3,7 | 4,5 | 1,3 | 0 | -2,22 | 1,3 | 0 | -8,89 |
| glass | 65,4 | 7,2 | 14 | 0,6 | -4,2 | -9,29 | -4 | -11 | -20 |
| iris | 92,7 | 3,7 | 3,7 | 0,7 | -5,4 | -21,6 | 0,7 | -5,4 | -21,6 |
| australian | 84,1 | 4,7 | 8,7 | 1 | -6,4 | -18,4 | 1 | -36 | -67,8 |
| pima | 73,9 | 3,7 | 6,3 | -0,9 | -11 | -23,8 | -2 | -32 | -50,8 |
| balance-scale | 77,4 | 11,9 | 32,6 | 6,5 | -82 | -76,7 | 5,6 | -83 | -79,1 |
| ionosphere | 89,8 | 4,8 | 5,2 | 0 | -4,2 | -7,69 | -2 | -19 | -26,9 |
| heart-statlog | 74,4 | 4 | 6,1 | 3 | -40 | -37,7 | 3 | -43 | -42,6 |
| pendigits | 86,7 | 30,8 | 75,1 | -2 | -15 | -16,5 | -8 | -23 | -23,6 |
| ecoli | 83 | 9,5 | 16,7 | 0,7 | -11 | -20,4 | -2 | -17 | -32,3 |
| yeast | 57,8 | 16,7 | 39,3 | 1,2 | -28 | -43,5 | -1 | -38 | -60,1 |
| segment | 94,4 | 13 | 26 | -0,5 | -7,7 | -15,4 | -5 | -23 | -42,3 |
| synth2D | 95 | 10,9 | 19,8 | 2,2 | -72 | -86,9 | 2,1 | -72 | -87,9 |
| synth3D | 95,8 | 7,8 | 19,6 | -1 | -51 | -57,7 | -6 | -71 | -76,5 |
| sc503 | 90,2 | 4,1 | 6,2 | 0 | -27 | -33,9 | -1 | -27 | -38,7 |
| Average | | | 1,0 | -23,8 | -31,8 | -1,3 | -32,7 | -45,5 | |

```
Input rule set:
```

```
(right-weight >= 3) and (right-distance
 >= 3) and (left-weight <= 2) =>
    class=R
```

- (left-distance <= 2) and (right-weight
 >= 3) and (right-distance >= 3) =>
 class=R
- (left-weight <= 3) and (left-distance <= 2) and (right-weight >= 2) and (right-distance >= 2) => class=R
- (left-weight <= 1) and (left-distance <= 3) and (right-distance >= 3) => class=R
- (left-distance <= 1) and (right-weight
 >= 3) and (left-weight <= 4) =>
 class=R
- (left-weight <= 1) and (right-distance >= 2) and (right-weight >= 2) => class=R
- (right-distance >= 4) and (right-weight >= 4) and (left-weight <= 3) => class=R
- (right-weight >= 5) and (right-distance >= 4) and (left-weight <= 4) =>
- class=R
 (left-distance <= 3) and (right-weight
 >= 3) and (right-distance >= 4) =>
 class=R
- (left-distance <= 3) and (right-weight >= 4) and (right-distance >= 2) and (left-weight <= 3) => class=R
- (left-distance <= 1) and (right-weight >= 2) and (right-distance >= 2) => class=R
- (left-weight <= 3) and (left-distance <= 1) and (right-distance >= 4) => class=R

```
Output rule set:
```

- (-0.5*left-distance+1 >= 0) and (+0.5*right-weight-1 >= 0) => class=R
- (-0.5*left-weight+0.75*right-distance-1 >= 0) and (+0.5*right-weight-1 >= 0) => class=R
- (-0.333*left-distance+0.666*rightweight-1 >= 0) and (+0.5*rightdistance-1 >= 0) => class=R

```
(-0.333*left-distance+1 >= 0) and
  (+0.5*right-distance-1 >= 0) and
  (+0.25*right-weight-1 >= 0) =>
      class=R
(-left-distance+1 >= 0) => class=R
```

```
(-0.25*left-weight-0.25*left-
distance+1 >= 0) => class=R
```

One can see that rule reduction rate is significant. The appearance of oblique descriptors allowed to decrease the number of rules describing the training data set and to reflect better dependences occurring in the "R" decision class. Determined rules are consistent with the *balance scale* dataset specificity.

In the case of the 2D synthetic dataset rules such as in Fig. 6 were managed to obtain. It clearly shows that the almost perfect description of the dataset has been obtained. Obtaining the perfect one (as in the Fig. 2) would be possible in the case of rotation of one oblique descriptor by a certain angle. Such tuning procedure will be the subject of further works with the purpose of improving the algorithm.

5 CONCLUSIONS

The proposition of an algorithm for classification rules aggregation that enables to introduce oblique descriptors in rules premises is presented in the paper. The aim of the algorithm is rules aggregation in order to obtain less number of rules describing decision classes. At the same time, the decrease of a rules number should not influence negatively the generalization abilities of the rules-based classifier.

Research results presented in tables 2a and 2b show that the algorithm operates according to the assumptions. Reduction of a number of rules and descriptors occurring in their premises is the result of the algorithm. The decrease of descriptors number is obviously caused, among others, by the fact that descriptors of aggregated rules can be linear combinations of several attributes, so they are more complicated than input ones. However, as the example of the synthetic dataset presented in the previous section shows, the change of descriptors representation can be helpful in better understanding dependencies in data.

The procedure tries to join rules sequentially, by twos, finding a set of hyperplanes which limit a region covered by rules being joined. Boundary conditions added synthetically to premises of aggregated rules are used in the hyperplanes searching only.

The algorithm can be parameterized by using various rule quality measures and various threshold values connected with a quality of joined rules and their classification abilities. As tables 2a and 2b show, values of parameters are important for the efficiency of the algorithm.

Further works on improving the algorithm performance will concern developing more advanced tuning strategy of joined rules. Beside the method of a hyperplane translation, a strategy of its rotation around a given point by a given angle will be worked out. Probably, authors will take advantage of experiences described by Murthy et al. (1994).

The algorithm sources (written entirely in MATLAB) can be provided after sending a request to one of the authors.

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