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Learning Nondeterministic Classifiers

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Abstract

Nondeterministic classifiers are defined as those allowed to predict more than one class for some entries from an input space. Given that the true class should be included in predictions and the number of classes predicted should be as small as possible, these kind of classifiers can be considered as Information Retrieval (IR) procedures. In this paper, we propose a family of IR loss functions to measure the performance of nondeterministic learners. After discussing such measures, we derive an algorithm for learning optimal nondeterministic hypotheses. Given an entry from the input space, the algorithm requires the posterior probabilities to compute the subset of classes with the lowest expected loss. From a general point of view, nondeterministic classifiers provide an improvement in the proportion of predictions that include the true class compared to their deterministic counterparts; the price to be paid for this increase is usually a tiny proportion of predictions with more than one class. The paper includes an extensive experimental study using three deterministic learners to estimate posterior probabilities: a multiclass Support Vector Machine (SVM), a Logistic Regression, and a Naïve Bayes. The data sets considered comprise both UCI multi-class learning tasks and microarray expressions of different kinds of cancer. We successfully compare nondeterministic classifiers with other alternative approaches. Additionally, we shall see how the quality of posterior probabilities (measured by the Brier score) determines the goodness of nondeterministic predictions.

Keywords: nondeterministic, multiclassification, reject option, multi-label classification, posterior probabilities

1. Introduction

There are several learners that successfully solve classification tasks in which the number of classes is higher than two; see for instance Wu et al. (2004) and Lin et al. (2008). However, for each class C most classification errors frequently occur between small subsets of classes that are somehow similar to C, regardless of the approach used. This fact suggests that multiclass classifiers would increase in reliability if they were allowed to express their doubts whenever they were asked to classify some entries.

In this paper we explore how to learn classifiers with multiple outcomes, like nondeterministic automata; we shall call them *nondeterministic classifiers*. Since they return a set of values, these classifiers could be called *set-valued* classifiers. To fix ideas, let us consider a screening for a set of medical diseases (or other diagnostic situations); for some inputs, a nondeterministic classifier

would be able to predict not just one single disease, but a set of options. These multiple predictions will be provided to domain experts when the classifier is not sure enough to give a unique class. Thus nondeterministic predictions may discard some options and allow domain experts to make practical decisions. Even when the nondeterministic classifier returns most of the available classes for the representation of an entry, we can read that the learned hypothesis is acknowledging its ignorance about how to deal with that entry.

It is evident that nondeterministic classifiers will include *true* classes in their predictions more frequently than deterministic hypotheses: they only have one possibility to be right. In this sense, nondeterministic predictions are backed by greater reliability. To be useful, however, nondeterministic classifiers should not only predict a set of classes containing the correct or true one, but their prediction sets should also be as small as possible. Notice that these requirements are common in algorithms designed for *Information Retrieval*. In this case, the queries are the entries to be classified and the *Recall* and *Precision* are then applied to each prediction. Hence, the loss functions for nondeterministic classifiers can be built as combinations of IR measures, as F_{β} functions are.

Starting from the distribution of posterior probabilities of classes, given one entry, we present an algorithm that computes the subset of classes with the lowest expected loss. In the experiments reported at the end of the paper, we employed three deterministic learners that provide posterior probabilities: Support Vector Machines (SVM), Logistic Regression (LR), and Naïve Bayes (NB). We successfully compared the achievements of our nondeterministic classifiers with those obtained by other alternative approaches.

The paper is organized as follows. In the next section, we present an overview of related work on classifiers that return subsets of classes instead of a single class. The formal settings both for nondeterministic classifiers and their loss functions are presented in the third section. After that, in Section 4, we derive an algorithm to learn nondeterministic hypotheses. Then, we conclude the paper with a section in which we report an experimental study of their performance. In addition to the comparison mentioned above, we discuss the role played by the deterministic learner that provides posterior probabilities. We see that the quality of posterior probabilities determines the goodness of nondeterministic predictions. The data sets used are publicly available and, in addition to a group of data sets from the UCI Repository (Asuncion and Newman, 2007), they include a group of classification tasks of cancer patients from gene expressions captured by microarrays.

2. Related Work

Nondeterministic classifiers are somehow related to classifiers with *reject option* (Chow, 1970). In this approach, the entries that are likely to be misclassified are rejected, they are not classified and can be handled by more sophisticated procedures: a manual classification, for instance. The core assumption is that the cost of making a wrong decision is 1, while the cost of using the reject option is given by some d, 0 < d < 1. In this context, provided that posterior probabilities are exactly known, an optimal rejection rule can be devised (Chow, 1970; Bartlett and Wegkamp, 2008): an entry is rejected if the maximum posterior probability is less than a threshold. Notice that classifiers with reject option are a relaxed version of nondeterministic classifiers. Rejection is a nondeterministic classification that includes the complete set of classes. On the other hand, instead of avoiding difficult classifications, for each entry, nondeterministic classifiers adventure a set of possible classes, not necessarily the complete set.

However, predictors of more than one class are not completely new. Given an $\varepsilon \in [0,1]$, the so-called confidence machines make *conformal predictions* (Shafer and Vovk, 2008): they produce a set of labels containing the true class with a probability greater than $1 - \varepsilon$.

To the best of our knowledge, the most directly related work to the approach presented in this paper is that of Zaffalon (2002) and Corani and Zaffalon (2008a,b). In these papers, the authors describe the *Naïve Credal Classifier*, a set-valued classifier which is an extension of the Naïve Bayes classifier to imprecise probabilities. The Naïve Credal Classifier models prior ignorance about the distribution of classes by means of a set of prior densities (also called the prior *credal set*), which is turned into a set of posterior probabilities by element-wise application of Bayes' rule. The classifier returns all the classes that are *non-dominated* by any other class according to the posterior credal set.

Another learning task that is related to this paper is multi-label classification. However, training instances in multi-label tasks can belong to more than one class, while nondeterministic training sets are the same as those of standard classification. In Tsoumakas and Katakis (2007), the authors provide an in-depth description of multi-label classification, enumerate several methods and compare their performance using Information Retrieval measures. Some applications have likewise arisen within the context of hierarchical organization of biological objects: predicting gene functions (Clare and King, 2003), or mapping biological entities into ontologies (Kriegel et al., 2004).

The formal setting presented in this paper was previously introduced in Alonso et al. (2008). There, we dealt with an interesting application of nondeterministic classifiers, in which classes (or *ranks*, in that context) are linearly ordered. The aim was to predict the rank (in an ordered scale) of carcasses of beef cattle. This value determines, on the one hand, the prices to be obtained by carcasses and, on the other, the genetic value of animals in order to select studs for the next generation. In this application, nondeterministic classifiers return an interval of ranks. Interval predictions are useful even when the intervals comprise more than one rank. For instance, it is possible to reject an animal as a stud for the next generation when a prediction interval is included in the lowest part of the scale. However, if we need a unique rank, we may decide to appeal to an *actual* expert to resolve the ambiguity, an expensive classification procedure not always available in practice.

The novelty of this paper is that now we deal with a standard classification setting; that is, the sets of classes are not ordered. This fact is very important as the search for the optimal prediction leads to a dramatic difference in complexity. Thus, if k is the number of classes, the search in the ordinal case is just of order k^2 , while in the unordered case, at a first glance, the search is of order 2^k . However, the Theorem of Correctness of Algorithm 1 proves that this search can be accomplished in polynomial time.

Additionally, this paper reports an extensive experimental study. First, we test whether non-deterministic classifiers outperform *Naïve Credal Classifiers* and other alternative approaches. We then investigate the role played by the ingredients of nondeterministic classifiers.

3. Formal Presentation and Notation

Let X be an input space and $\mathcal{Y} = \{C_1, ..., C_k\}$ a finite set of classes. We consider a multiclassification task given by a training set $S = \{(x_1, y_1), ..., (x_n, y_n)\}$ drawn from an unknown distribution Pr(X, Y) from the product $X \times \mathcal{Y}$. Within this context, we define

Definition 1 A nondeterministic hypothesis is a function h from the input space to the set of nonempty subsets of \mathcal{Y} ; in symbols, if $\mathcal{P}(\mathcal{Y})$ is the set of all subsets of \mathcal{Y} ,

$$h: \mathcal{X} \longrightarrow \mathscr{P}(\mathcal{Y}) \setminus \{\varnothing\}.$$

The aim of such a learning task is to find a nondeterministic hypothesis h from a space \mathcal{H} that optimizes the *expected prediction performance (or risk)* on samples S' independently and identically distributed (i.i.d.) according to the distribution Pr(X,Y)

$$R^{\Delta}(h) = \int \Delta(h(x), y) d(Pr(x, y)),$$

where $\Delta(h(x), y)$ is a loss function that measures the penalty due to the prediction h(x) when the true value is y.

In nondeterministic classification, we would like to favor those decisions of h that contain the true classes, and a smaller rather than a larger number of classes. In other words, we interpret the output h(x) as an imprecise answer to a query about the right class of an entry $x \in \mathcal{X}$. Thus, nondeterministic classification can be seen as a kind of Information Retrieval task for each entry.

Performance in Information Retrieval is compared using different measures in order to consider different perspectives. The most frequently used measures are Recall (proportion of all relevant documents that are found by a search) and Precision (proportion of retrieved documents that are relevant). The harmonic average of the two amounts is used to capture the goodness of a hypothesis in a single measure. In the weighted case, the measure is called F_{β} . The idea is to measure a tradeoff between Recall and Precision.

For further reference, let us recall the formal definitions of these Information Retrieval measures. Thus, for a prediction of a nondeterministic hypothesis h(x) with $x \in \mathcal{X}$, and a class $y \in \mathcal{Y}$, we can compute the following contingency matrix, where $z \in \mathcal{Y}$,

in which each entry (a,b,c,d) is the number of times that the corresponding combination of memberships occurs. Notice that a can only be 1 or 0, depending on whether the class y is included in the prediction h(x) or not; b is the number of classes different from y included in h(x); c = 1 - a; and d is the number of classes different from y that are not included in h(x).

According to the matrix, Equation (1), if h is a nondeterministic hypothesis and $(x,y) \in X \times \mathcal{Y}$, we thus have the following definitions.

Definition 2 *The* Recall *in a* query (*i.e.*, *an entry x*) *is defined as the proportion of relevant classes* (y) *included in* h(x):

$$R(h(x), y) = \frac{a}{a+c} = a = 1_{y \in h(x)}.$$

Definition 3 The Precision is defined as the proportion of retrieved classes in h(x) that are relevant (y):

$$P(h(x), y) = \frac{a}{a+b} = \frac{1_{y \in h(x)}}{|h(x)|}.$$

h(x)	Precision	Recall	F_1	F_2
[1,2,3]	0.33	1	0.50	0.71
[1, 2]	0.50	1	0.67	0.83
[1]	1	1	1	1
[2, 3, 4]	0	0	0	0

Table 1: The *Precision*, *Recall*, F_1 , and F_2 for different predictions of a nondeterministic classifier h for an entry x with class 1, (y = 1)

In other words, given a hypothesis h, the *Precision* for an entry x, that is, P(h(x), y), is the probability of finding the true class (y) of the entry (x) by randomly choosing one of the classes of h(x).

Finally, the tradeoff is formalized by

Definition 4 *The* F_{β} *is defined, in general, by*

$$F_{\beta}(h(x), y) = \frac{(1+\beta^2)PR}{\beta^2 P + R} = \frac{(1+\beta^2)a}{(1+\beta^2)a + b + \beta^2 c}.$$
 (2)

Thus, for a nondeterministic classifier h and a pair (x, y),

$$F_{\beta}(h(x), y) = \begin{cases} \frac{1+\beta^2}{\beta^2 + |h(x)|} & \text{if} \quad y \in h(x) \\ 0 & \text{otherwise.} \end{cases}$$
 (3)

The most frequently used F-measure is F_1 . For ease of reference, let us state that

$$F_1(h(x), y) = \frac{2_{y \in h(x)}}{1 + |h(x)|}.$$

Notice that for deterministic classifiers, the accuracy is equal to *Recall*, *Precision*, and F_{β} given that |h(x)| = 1.

To illustrate the use of the F-measures of an entry, let us consider an example. If we assume that the true class of an entry x is 1, (y = 1), then, depending on the value of h(x), Table 1 reports the *Recall, Precision*, F_1 , and F_2 . We observe that the reward attached to a prediction containing the true class with another extra class ranges from 0.667 for F_1 to 0.833 for F_2 ; whereas the amounts are lower when the prediction includes 2 extra classes.

Once we have the definition of F_{β} for individual entries, it is straightforward to extend it to a test set. Hence, when S' is a test set of size n, the average loss on it will be computed by

$$R^{\Delta^{ND}}(h,S') = \frac{1}{n} \sum_{j=1}^{n} \Delta^{ND}(h(x'_j), y'_j) = \frac{1}{n} \sum_{j=1}^{n} \left(1 - F_{\beta}(h(x'_j), y'_j) \right)$$

$$= \frac{1}{n} \sum_{j=1}^{n} \left(1 - \frac{1 + \beta^2}{\beta^2 + |h(x'_j)|} 1_{y'_j \in h(x'_j)} \right). \tag{4}$$

The average *Recall* and *Precision* can be similarly defined. For ease of reference, let us remark that the *Recall* is the proportion of times that h(x') includes y' and is thus a generalization of the *deterministic accuracy*.

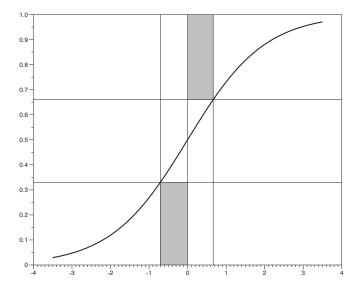


Figure 1: Conditional probabilities of class +1 given the discriminant value (horizontal axis) of entries $x \in \mathcal{X}$. Vertical bars separate the region where both classes $\{-1,+1\}$ have a probability of over 1/3

3.1 Nondeterministic Classification in a Binary Task

To complete this section, let us show what nondeterministic classifiers look like in the simplest case, which will be further developed in the following sections. Let us assume that in a binary classification task (the classes are codified by -1 and +1) we have a loss 1 for each false classification. On the other hand, we are allowed to predict both classes, in which case the loss will be 1/3: the F_1 for a classification of 2 classes containing the true one; see Table 1. The extension for dealing with F_{β} , with $\beta \neq 1$, is straightforward.

The optimum classifier will return only one class when it is sufficiently sure. In doubtful situations, however, the nondeterministic classifier should opt for predicting the 2 classes. This will be the case whenever the probability of error for both classes is higher than 1/3, since this is the loss for predictions of two classes; see Figure 1. Therefore, if we have the conditional probabilities of classes given the entries, the optimum classifier will be given by

$$h_{ND}(x) = \begin{cases} \{-1\} & \text{if } \eta(x) < 1/3\\ \{-1, +1\} & \text{if } 1/3 \le \eta(x) < 2/3\\ \{+1\} & \text{if } 2/3 \le \eta(x), \end{cases}$$
 (5)

where we are representing by $\eta(x)$ the posterior probability:

$$\eta(x) = Pr(class = +1|x).$$

Notice that Equation (5) is equivalent to the generalized Bayes discriminant function described in Bartlett and Wegkamp (2008) when the cost of using the reject option is calculated using the F_1 loss function.

Algorithm 1 The nondeterministic classifier nd^{\bullet} , an algorithm for computing the prediction with one or more classes for an entry x provided that the posterior probabilities of classes are given

```
Input: \left\{C_j: j=1,...,k \, sorted \, by \, Pr(C_j|x)\right\}
Input: \beta: trade-off between Recall and Precision
Initialize i=0,\,\Delta_0=1
repeat
i=i+1
\Delta_i=1-\frac{1+\beta^2}{\beta^2+i}\sum_{j=1}^i Pr(C_j|x)
until ((i==k) \text{ or } (\Delta_{i-1} \leq \Delta_i)
if (\Delta_{i-1} \leq \Delta_i) then
return \left\{C_j: j=1,...,i-1\right\}
else
return \left\{C_j: j=1,...,k\right\}
end if
```

4. Nondeterministic Classification Using Multiclass Posterior Probabilities

In the general multiclass setting presented at the beginning of Section 3, let x be an entry of the input space \mathcal{X} and let us now assume that we know the conditional probabilities of classes given the entry, $Pr(C_j|x)$. Additionally, we shall assume that the classes are ordered according to these probabilities. In this context, we wish to define the

$$h(x) = Z \subset \mathcal{Y} = \{C_1, \dots, C_k\}$$

that minimizes the risk defined in Equation (1) when we use the nondeterministic loss given by F_{β} , (Equations 2, 3, and 4). We shall prove that such an h(x) can be computed by Algorithm 1, which does not need to search through all non-empty subsets of \mathcal{Y} .

Theorem 1 (Correctness). If the conditional probabilities $Pr(C_j|x)$ are known, Algorithm 1 returns the nondeterministic prediction for h(x) that minimizes the risk given by the loss $1 - F_{\beta}$.

Proof To minimize the risk, Equation (1), it suffices to compute

$$\Delta_{x}(Z) = \sum_{y \in \mathcal{Y}} \Delta^{ND}(Z, y) Pr(y|x), \tag{6}$$

with $Z \subset \{C_1, \dots, C_k\}$. Then, we only have to define

$$h(x) = argmin\{\Delta_x(Z) : Z \subset \{C_1, \dots, C_k\}\}.$$

The proof has two parts. First, we shall see that if h(x) has r classes, then those are the r classes with the highest probabilities; bearing in mind that classes are ordered, $h(x) = Z_r = \{C_j : j = 1, ..., r\}$. For this purpose, we need to see that any other subset of r classes will increase the loss due to Z_r . This is a consequence of the following.

The value of Equation (6) for Z_r is Δ_r in Algorithm 1. In fact, with the complementary probability of $\sum_{i=1}^r Pr(C_i|x)$, we expect a loss of 1: the *true* class will not be one of the r first classes. On

the other hand, with this sum of probabilities, the *true* class will be in h(x), and therefore the loss will be 1 minus the F_{β} of the prediction $h(x) = \{C_j : j = 1,...,r\}$:

$$\Delta_{x}(C_{j}: j = 1, ..., r) = \left(1 - \sum_{j=1}^{r} Pr(C_{j}|x)\right) + \left(\sum_{j=1}^{r} Pr(C_{j}|x)\right) \left(1 - \frac{1 + \beta^{2}}{\beta^{2} + r}\right)$$

$$= 1 - \frac{1 + \beta^{2}}{\beta^{2} + r} \sum_{j=1}^{r} Pr(C_{j}|x)$$

$$= \Delta_{r}.$$

Notice that for any other subset of r classes, we could achieve a similar expression simply by modifying the set of posterior probabilities of the last sum. Therefore, to minimize the value of Equation (6) with r classes, we need those with the highest probability.

In the second step, we only have to show that the index r returned by the Algorithm is the right one. We shall see that the search for the best r can be accomplished in linear time, as in the Algorithm. In fact, we shall establish that when the Algorithm reaches the number of classes with which the loss increases, adding further classes will only increase the loss. In symbols, we shall prove that

$$\Delta_r \leq \Delta_{r+1} \Rightarrow \Delta_{r+1} \leq \Delta_{r+2}$$
.

To do so, we shall next express the exit condition of the loop $\Delta_r \leq \Delta_{r+1}$ when $(r+1) \leq k$ in a different way. The following expressions are equivalent:

$$\Delta_{r} \leq \Delta_{r+1}$$

$$\frac{1+\beta^{2}}{\beta^{2}+r} \sum_{j=1}^{r} Pr(C_{j}|x) \geq \frac{1+\beta^{2}}{\beta^{2}+r+1} \sum_{j=1}^{r+1} Pr(C_{j}|x)$$

$$(\beta^{2}+r+1) \sum_{j=1}^{r} Pr(C_{j}|x) \geq (\beta^{2}+r) \sum_{j=1}^{r+1} Pr(C_{j}|x)$$

$$\sum_{j=1}^{r} Pr(C_{j}|x) \geq (\beta^{2}+r) Pr(C_{r+1}|x).$$
(7)

Therefore, if $\Delta_r \leq \Delta_{r+1}$ and $(r+1) \leq k$, then

$$Pr(C_{r+1}|x) + \sum_{j=1}^{r} Pr(C_j|x) \ge (\beta^2 + r)Pr(C_{r+1}|x) + Pr(C_{r+1}|x).$$

However, bearing in mind that the classes are ordered, we have that $Pr(C_{r+1}|x) \ge Pr(C_{r+2}|x)$, and using Equation (7), we conclude that

$$\sum_{i=1}^{r+1} Pr(C_j|x) \ge (\beta^2 + r + 1) Pr(C_{r+2}|x) \Leftrightarrow \Delta_{r+1} \le \Delta_{r+2}.$$

4.1 Corollaries

In order to draw some practical consequences, let us reword the previous Theorem. It states that the optimum classification for an input x is the set of r classes with the highest posterior probabilities, where r is the lowest integer that fulfills

$$\sum_{j=1}^{r} Pr(C_j|x) \ge (\beta^2 + r) Pr(C_{r+1}|x), \tag{8}$$

or the set of all classes when this condition is not fulfilled by any r. Expressed in this way, it is straightforward to see that for two classes, with $\beta = 1$, Algorithm 1 coincides with the rule defined in Equation (5).

Additionally, we would like to underscore that Equation (8) hinders the use of naïve *thresholds* to compute nondeterministic predictions. Thus, a nondeterministic classifier that always predicts the top r classes for a constant value r is not a correct option. Equation (8) shows that r, at least, depends on the input x.

Moreover, we should not search for a threshold λ to return, for all inputs, the first r classes whose sum of probabilities is above λ :

$$\sum_{j=1}^{r} Pr(C_j|x) \ge \lambda. \tag{9}$$

Note that given a λ value in [0,1], Equation (9) straightforwardly gives rise to a nondeterministic classifier as follows. For each input x, if the set of classes is ordered according to their posterior probabilities, we define

$$h_{\lambda}(x) = \left\{ C_1, \dots, C_r : \sum_{j=1}^r \Pr(C_j | x) \ge \lambda \quad \& \quad \sum_{j=1}^{r-1} \Pr(C_j | x) < \lambda \right\}. \tag{10}$$

Again, the right-hand side of Equation (8) shows that the threshold (λ) would depend on the number of classes predicted, the probability of the first class excluded from the prediction, and the parameter β : the trade-off between *Precision* and *Recall*. The idea behind Equation (8) is that, once we have decided to include the top r classes, to add the $(r+1)^{th}$ class we should guarantee that $Pr(C_{r+1}|x)$ is not much smaller than the sum of probabilities of the top r classes.

However, it may be argued that the inaccuracy of posterior probabilities would partially invalidate the preceding theoretical discussion. In fact, posterior probabilities are not known in practice: they are estimated by algorithms that frequently try to optimize the classification accuracy of a hypothesis that returns the class with the highest probability. In other words, probabilities are discriminant values instead of thorough descriptions of the distribution of classes in a learning task. Therefore, in the experiments reported at the end of the paper, we shall consider the classifiers defined by Equation (10) as a possible alternative method to the nondeterministic classifier of Algorithm 1.

5. Experimental Results

In this section we report the results of a set of experiments conducted to evaluate the proposals of this paper. The next subsection describes the settings used in the experiments: deterministic learners, data sets, procedures to set parameters, and methods to estimate the scores.

Data sets	#classes	#samples	#features
Z00	7	101	16
iris	3	150	4
glass	6	214	9
ecoli	8	336	7
balance scale	3	625	4
vehicle	4	846	18
vowel	11	990	11
contraceptive	3	1473	9
yeast	10	1484	8
car	4	1728	6
image	7	2310	19
waveform	3	5000	40
landsat	6	6435	36
letter recognition	26	20000	16

Table 2: Description of the data sets downloaded from the UCI repository. The classes are not linearly separable

We have two goals here. On the one hand, we compare our approach with two alternative methods. The comparison will first be established with a state-of-the-art set-valued algorithm, the Naïve Credal Classifier (NCC) (Zaffalon, 2002; Corani and Zaffalon, 2008a,b). This algorithm is an extension of the traditional Naïve Bayes classifier towards imprecise probabilities and is designed to return robust set-valued (nondeterministic) classifications. We show that our method can improve the performance of NCC. We then contrast our method with an implementation of Equation (10); once again our proposals outperform this alternative way to learn nondeterministic classifiers.

On the other hand, we analyze the influence of a number of factors related to nondeterministic learners. We accordingly discuss how the scores of a nondeterministic learner are affected by the quality of posterior probabilities. We see that the performance of a nondeterministic classifier is highly correlated with the accuracy of its deterministic counterpart. The section ends with a study of the meaning of the parameter β .

5.1 Experimental Settings

We used three different methods for learning posterior probabilities in order to build nondeterministic classifiers. First, we employed the Naïve Bayes (NB) used by NCC as its deterministic counterpart (Corani and Zaffalon, 2008b). The second deterministic learner was a multiclass SVM; the implementation used was libsvm (Wu et al., 2004) with the linear kernel. Last, we employed the logistic regression (LR) of Lin et al. (2008). It should be noted that we are not only using the multiclass classifiers learned by SVM or LR. Primarily, we apply the mechanisms that provide posterior probabilities from their outputs.

For each of these learners, we built nd^d , where d stands for the name of the deterministic counterpart, nb, svm or lr. Recall that nd^d is the implementation of Algorithm 1 that aims to optimize F_1 ; that is, $\beta = 1$.

Data sets	#classes	#samples	#features	Original source	Used in
brain	5	42	5597	Pomeroy et al. (2002)	[1]
nci	9	60	7131	Ross et al. (2000)	[1, 3, 4]
lung 6	6	70	16387	Tamayo et al. (2007)	
leukemia 3	3	72	12582	Armstrong et al. (2002)	[2]
lung 4	4	82	9036	Tamayo et al. (2007)	
lung 11	11	89	4459	Tamayo et al. (2007)	
tumors 11	11	174	12533	Su et al. (2001)	[1, 2]
tumors 14	14	190	16063	Ramaswamy et al. (2001)	[1, 2, 4]
lung 16	16	201	493	Tamayo et al. (2007)	
leukemia 7	7	327	12558	Yeoh et al. (2002)	[2]

Table 3: Description of cancer microarray data sets used in the experiments including the original sources and papers from which they are taken. For the sake of brevity, we have denoted the papers as follows: [1] Tibshirani and Hastie (2007), [2] Tan et al. (2005), [3] Staunton et al. (2001), [4] Yeung and Bumgarner (2003)

In the experiments that follow, we used two kinds of data sets. First, we considered data sets downloaded from the UCI repository (Asuncion and Newman, 2007), all of which have more examples than attributes. We included all the data sets that fulfill the following rules: continuous or ordinal attribute values, no more than 40 attributes and no more than 20000 examples. The intention was to consider small data sets that are not linearly separable. Additionally, we excluded those learning tasks with missing values or in which every deterministic learner considered (NB, SVM, LR) achieves a proportion of successful classifications of over 95%; otherwise nondeterministic learners would be too similar to their deterministic counterpart. A description of the group of data sets considered can be found in Table 2.

We then evaluated the performance on learning tasks in which the aim was to classify cancer patients from gene expressions captured by microarrays. Unlike the first package of data sets, all the classes are now linearly separable given the dimensions of the input space and the number of entries. Table 3 shows the details of these data sets.

Every table of scores (Tables 4, 5, 6, 7, 8) is devoted to reporting the experimental results achieved in one of the kinds of data sets by one of the deterministic learners and by two nondeterministic algorithms that are to be compared. All the tables have a similar layout. First, they contain the scores of the deterministic learner d: the F_1 (or accuracy or Recall), and the Brier score, a measure for the quality of posterior probabilities (Brier, 1950; Yeung et al., 2005), computed by means of

$$BS = \frac{1}{2n} \sum_{i=1}^{n} \sum_{j=1}^{k} ([y_i = C_j] - Pr(C_j | x_i))^2.$$

Then we report, for each nondeterministic learner, the F_1 , Precision, Recall, and the average number of classes predicted (|h(x)|). All the scores were estimated by means of a 5-fold cross validation repeated 2 times. We did not use the 10-fold procedure, since in certain data sets there are too few examples in some of the classes.

Following Demšar (2006), we used the Wilcoxon signed ranks test to compare the performance of two classifiers when the measurements are F_1 , *Precision*, *Recall*, or the average |h(x)|. Unless

	N	'B		Λ	ICC		$ $ nd^{nb}			
Data set	F_1	BS	F_1	P	R	h(x)	F_1	P	R	h(x)
ZOO	95.0	0.03	92.3	90.5	100.0	1.496	95.2	94.3	97.0	1.055
iris	93.3	0.05	92.9	92.5	94.0	1.037	93.9	93.5	94.7	1.023
glass	68.0	0.22	69.2	66.5	76.6	1.321	70.7	67.6	77.6	1.253
ecoli	83.5	0.12	82.0	81.0	85.7	1.240	84.4	82.2	88.7	1.136
balance	73.9	0.16	76.0	74.2	79.7	1.132	79.9	74.9	90.1	1.370
vehicle	60.8	0.30	60.9	59.8	63.4	1.103	63.3	60.2	69.6	1.241
vowel	62.1	0.25	64.6	62.6	69.8	1.296	65.5	60.9	75.5	1.429
contra	50.0	0.30	50.3	50.1	50.6	1.013	56.6	47.9	74.6	1.670
yeast	58.1	0.28	58.4	58.2	59.0	1.037	60.8	54.4	74.3	1.500
car	86.8	0.11	87.3	87.0	87.8	1.017	83.4	76.6	98.0	1.487
image	90.9	0.08	91.4	90.5	94.8	1.195	91.2	90.9	92.0	1.026
waveform	80.1	0.17	80.1	80.0	80.4	1.007	80.9	80.0	82.5	1.051
landsat	82.0	0.17	82.0	81.6	83.1	1.058	82.1	81.9	82.4	1.011
letter	73.9	0.19	74.6	74.2	75.8	1.081	74.8	73.3	78.0	1.166

Table 4: Scores obtained by Naïve Bayes, the Naïve Credal Classifier and nondeterministic classifiers on UCI data sets using a 5-fold cross validation repeated 2 times. For ease of reading, F_1 , Precision (P), and Recall (R) are expressed as percentages. The best nondeterministic F_1 for each data set is boldfaced

explicitly stated, we use the expression statistically *significant differences* to mean that p < 0.01. Additionally, in order to provide a quick view of the order of magnitude of the scores, we have boldfaced the best nondeterministic F_1 score for each data set.

To select the regularization parameter, C, for SVM and LR, we used a 2-fold cross validation repeated 5 times performed on training sets. We searched within $C \in [10^{-2}, \dots, 10^{2}]$.

5.2 Nondeterministic Classifiers vs. Naïve Credal Classifiers

In this subsection, we compare our nondeterministic learner with NCC (Corani and Zaffalon, 2008b), a state-of-the-art set-valued (nondeterministic) algorithm. In order to ensure a fair comparison, our approach uses the Naïve Bayes (NB) employed by NCC as its deterministic counterpart. Table 4 reports the scores of NB, NCC and our algorithm nd^{nb} .

The nondeterministic nd^{nb} is significantly (remember that we are using Wilcoxon tests) better than NCC both in Recall and F_1 . Moreover, nd^{nb} wins in 12 out of 14 data sets in F_1 , and in 11 out of 14 in Recall. However, the scores in Precision and size of predictions are more balanced; the differences are not significant. In Precision, NCC wins in 5 cases, loses in 8, and there is 1 tie situation. The size scores are favorable to NCC in 8 out of 14 data sets.

To complete the comparison, we should discuss the results achieved on high dimensional data sets (Table 3). Nevertheless, we do not show the scores on each data set. The characteristics of these tasks are not appropriate for Naïve Bayes (a large number of attributes with a small number of examples); therefore, the posterior probabilities of NB are poor (they are significantly worse than those achieved by SVM and LR) and this affects the performance of our nondeterministic algorithm and NCC. Our method tends to be almost deterministic, the average value for the size of predictions is |h(x)| = 1.008. This is not optimal, as we shall see later, but it is acceptable behavior. However,

	SV	[/] M	nd_{λ}^{svm}			nd ^{svm}				
Data set	F_1	BS	F_1	P	\tilde{R}	h(x)	F_1	P	R	h(x)
ZOO	94.0	0.08	38.9	24.6	100.0	4.390	94.2	92.4	98.0	1.134
iris	96.0	0.02	83.2	74.8	100.0	1.510	97.6	96.7	99.3	1.053
glass	61.7	0.26	63.7	53.3	85.0	1.711	63.0	55.9	77.3	1.484
ecoli	86.5	0.11	75.1	66.3	97.2	1.854	87.4	85.0	92.3	1.152
balance	91.7	0.06	83.8	77.5	98.7	1.528	91.3	89.0	98.1	1.272
vehicle	79.8	0.13	79.6	71.0	97.8	1.576	82.5	77.9	92.0	1.297
vowel	82.0	0.15	66.3	55.0	97.5	2.313	82.9	78.8	91.5	1.288
contra	51.3	0.29	55.9	48.3	71.3	1.599	57.7	46.7	83.1	1.960
yeast	59.0	0.27	60.6	50.4	82.2	1.817	62.4	53.4	81.6	1.706
car	85.3	0.11	82.8	76.1	97.3	1.475	85.6	83.0	90.8	1.169
image	95.9	0.03	84.8	79.1	99.8	1.579	96.1	95.3	97.9	1.058
waveform	86.4	0.10	85.7	80.0	97.1	1.343	87.6	81.5	91.8	1.126
landsat	86.8	0.09	84.4	78.4	97.6	1.453	87.8	85.7	91.9	1.139
letter	85.8	0.11	71.0	64.3	98.2	2.949	86.3	76.7	91.0	1.186

Table 5: Scores obtained by SVM learners on UCI data sets using a 5-fold cross validation repeated 2 times. For ease of reading, F_1 , Precision (P), and Recall (R) are expressed as percentages. The best nondeterministic F_1 for each data set is boldfaced

		R		1	id_{λ}^{lr}		$ $ nd^{lr}			
Data set	F_1	BS	F_1	P	\tilde{R}	h(x)	F_1	P	R	h(x)
ZOO	95.0	0.04	91.0	88.4	97.0	1.252	95.4	95.0	96.0	1.045
iris	96.7	0.05	74.4	61.7	100.0	1.767	94.4	92.2	99.0	1.137
glass	60.3	0.27	61.5	49.3	86.0	1.844	63.0	51.8	85.5	1.774
ecoli	87.5	0.11	76.9	68.3	96.1	1.668	87.0	84.4	92.1	1.173
balance	86.7	0.11	88.9	87.4	92.6	1.185	88.7	87.7	90.9	1.136
vehicle	77.0	0.16	74.8	64.9	95.3	1.674	79.2	74.1	89.7	1.342
vowel	57.9	0.30	54.1	41.5	83.5	2.226	57.8	48.6	79.7	1.908
contra	50.8	0.29	55.9	47.7	72.3	1.644	58.0	47.4	82.5	1.928
yeast	58.4	0.28	59.4	49.0	80.9	1.818	61.0	52.2	79.9	1.713
car	80.9	0.13	80.7	74.1	95.0	1.482	82.0	78.9	88.5	1.215
image	88.4	0.11	72.3	60.9	98.7	1.915	88.0	85.1	93.8	1.196
waveform	86.5	0.10	81.8	72.8	99.6	1.536	87.4	82.9	96.4	1.272
landsat	77.7	0.18	68.6	58.1	93.8	1.940	76.6	71.7	86.9	1.387
letter	71.8	0.24	49.3	36.5	90.7	3.253	70.3	64.9	82.5	1.556

Table 6: Scores obtained by LR learners on UCI data sets using a 5-fold cross validation repeated 2 times. For ease of reading, F_1 , Precision (P), and Recall (R) are expressed as percentages. The best nondeterministic F_1 for each data set is boldfaced

the scores of *NCC* on these data sets are inadmissible; their classifiers predict almost all classes for every example, their average values are: $F_1 = 25.73$, P = 15.39, R = 100, and |h(x)| = 8.58.

In fact, the behavior of *NCC* is difficult to predict, sometimes it is almost a deterministic classifier, whereas in other tasks the number of classes predicted by *NCC* is very high. Moreover, its degree of nondeterminism is not related to the difficulty of the learning task. When the accuracy of the deterministic classifiers decreases, the average number of classes predicted would be expected

	SV	VM	nd_{λ}^{svm}			nd ^{svm}				
Data set	F_1	BS	F_1	P	\tilde{R}	h(x)	F_1	P	R	h(x)
brain	81.8	0.15	59.2	44.9	97.5	2.504	82.9	78.0	93.8	1.401
nci	48.3	0.35	42.9	33.2	68.3	2.492	47.7	41.4	65.0	2.167
lung 6	72.1	0.21	65.7	57.9	85.7	1.907	73.0	70.4	78.6	1.221
leukemia 3	94.5	0.04	75.1	64.5	100.0	1.862	95.7	94.9	97.3	1.049
lung 4	87.1	0.11	73.9	63.0	96.9	1.743	87.3	85.3	91.4	1.122
lung 11	58.4	0.31	49.3	36.5	84.2	2.656	60.4	53.8	78.0	1.903
tumors 11	89.6	0.13	30.6	19.1	99.7	6.135	88.9	87.1	92.8	1.199
tumors 14	70.0	0.26	45.0	35.3	95.0	4.550	66.5	60.2	84.7	2.021
lung 16	84.8	0.17	25.0	14.5	100.0	7.440	87.3	83.1	95.8	1.266
leukemia 7	92.0	0.07	70.1	59.9	99.4	2.216	92.1	90.6	95.1	1.090

Table 7: Scores obtained by SVM learners on cancer microarray data sets using a 5-fold cross validation repeated 2 times. For ease of reading, F_1 , Precision (P), and Recall (R) are expressed as percentages. The best nondeterministic F_1 for each data set is boldfaced

	L	R	nd_{λ}^{lr}			nd^{lr}				
Data set	F_1	BS	F_1	P	\tilde{R}	h(x)	F_1	P	R	h(x)
brain	86.8	0.11	86.3	82.7	94.0	1.274	86.1	84.5	89.2	1.106
nci	55.8	0.33	56.7	55.8	58.3	1.142	57.8	56.7	60.0	1.158
lung 6	70.7	0.22	74.3	72.5	77.9	1.150	73.3	72.1	75.7	1.107
leukemia 3	97.3	0.02	92.2	88.8	100.0	1.258	97.7	97.3	98.7	1.028
lung 4	88.9	0.10	88.9	86.4	93.9	1.153	88.8	87.8	90.8	1.061
lung 11	69.0	0.24	69.1	65.2	77.5	1.354	68.9	65.7	75.9	1.316
tumors 11	94.8	0.05	89.5	85.9	99.1	1.421	93.7	93.0	95.1	1.057
tumors 14	75.3	0.18	76.8	73.9	83.2	1.337	76.8	75.2	80.3	1.145
lung 16	88.1	0.10	88.3	86.0	93.0	1.157	88.4	87.4	90.3	1.060
leukemia 7	91.9	0.07	90.6	87.9	96.3	1.202	91.9	91.4	93.1	1.040

Table 8: Scores obtained by LR learners on cancer microarray data sets using a 5-fold cross validation repeated 2 times. For ease of reading, F_1 , Precision (P), and Recall (R) are expressed as percentages. The best nondeterministic F_1 for each data set is boldfaced

to increase. However the correlation between the accuracy of NB and |h(x)| of NCC is 0.24. In the case of nd^{nb} , this correlation is -0.75: negative and quite high.

5.3 Comparing nd with Another Alternative Method

In accordance with the discussion in Section 4.1, we shall now compare the nondeterministic classifiers learned by Algorithm 1 with the alternative classifier defined in Equation (10) that uses a threshold λ for the sum of posterior probabilities. The comparison will be established with posterior probabilities provided by *SVM* and *LR* given that both outperform the accuracy achieved by Naïve Bayes classifiers in the data sets used in these experiments. The λ nondeterministic classifiers will be denoted by nd_{λ}^{d} , where d stands for the deterministic counterpart.

To select the parameter λ , we use a grid search employing a 2-fold cross validation repeated 5 times, aiming to optimize F_1 . The searching space depends on the learning task S. If the proportion of successful classifications for deterministic classifiers, the accuracy, is a, then we search within

 $\lambda \in [a_0, a_1, \dots, a_5]$; six options distribute from a to 0.99. In symbols, $a_0 = a, a_5 = 0.99$, and $a_{i+1} - a_i = \frac{0.99 - a}{5}$.

In UCI data sets, Tables 5 and 6, nd^{svm} and nd^{lr} win the corresponding nd_{λ} in 13 out of 14 data sets in F_1 and Precision. In Recall we have the opposite situation; λ classifiers win in 13 out of 14 cases. Moreover, λ classifiers always predict more classes than nd^{svm} and nd^{lr} . In other words, λ classifiers predict more classes than necessary. All differences are significant. Thus, our nd classifiers are better than those computed with the λ parameter.

In cancer microarray data, Tables 7 and 8, nd^{svm} always wins in F_1 , Precision, and average |h(x)|; while nd^{svm} always loses in Recall. All differences are again significant. However, when posterior probabilities are provided by LR, the differences are not significant in F_1 , although nd^{lr} has 5 wins, 1 tie and 4 losses; in Precision and average size of predictions the differences are significant in favor of nd^{lr} . Furthermore, as usual, the Recall is significantly higher for λ classifiers.

The conclusion is that λ classifiers seem to need more classes in their predictions than nd classifiers. In fact, Equation (9) only considers the *Recall*. In practice, this means more *Recall*, but less *Precision* and F_1 . Therefore, to optimize the F_1 measure, in an experimental environment, Equation (8) is more adequate than Equation (9), as we have conjectured theoretically in Section 4.1.

5.4 The Importance of Posterior Probabilities

The objective of this subsection is to experimentally investigate the degree of dependency between nondeterministic scores and the accuracy of posterior probabilities. In this study we again employ *SVM* and *LR* with the collection of data sets detailed in Tables 2 and 3.

Let us first consider the set of UCI data sets. Comparing the results in Tables 5 and 6, it can be seen that the scores of nd^{lr} are significantly worse than those of nd^{svm} in F_1 , Precision, Recall (p < 0.03), and in average size of predictions. The general message is that nd^{lr} include unnecessary classes in their predictions. The base posterior probabilities seem to be the cause of this behavior: the Brier score of LR is significantly worse than that of SVM.

On the other hand, the scores obtained with cancer microarray data sets are shown in Tables 7 and 8. The characteristics of UCI and microarray data sets are quite different, and this affects the performance of classifiers. The main difference is that LR now has a significantly better Brier score than SVM. Moreover, the nd^{lr} algorithm achieves better results than nd^{svm} . The differences are significant in F_1 , Precision, $Recall\ (p < 0.02)$, and average |h(x)|. Yet again, inferior posterior probabilities seem to be responsible for the inclusion of unnecessary classes in nondeterministic predictions.

In the preceding discussion of the scores achieved by nondeterministic learners, we found significant differences when the Brier scores of the deterministic counterparts presented significant differences. In fact, the scores of a learner built with Algorithm 1 depend on the quality of the posterior probabilities supplied by the corresponding deterministic learner. It seems plausible to draw the conclusion that the better the posterior probabilities, the better the nondeterministic scores. In order to quantify this statement, we compared deterministic Brier scores with nondeterministic F_1 , Recall, and Precision values; see Figure 2. We separated the scores achieved by UCI and cancer data sets and included the scores of nd^{nb} in UCI data sets. Similar results would be achieved if we compared nondeterministic scores with deterministic accuracy.

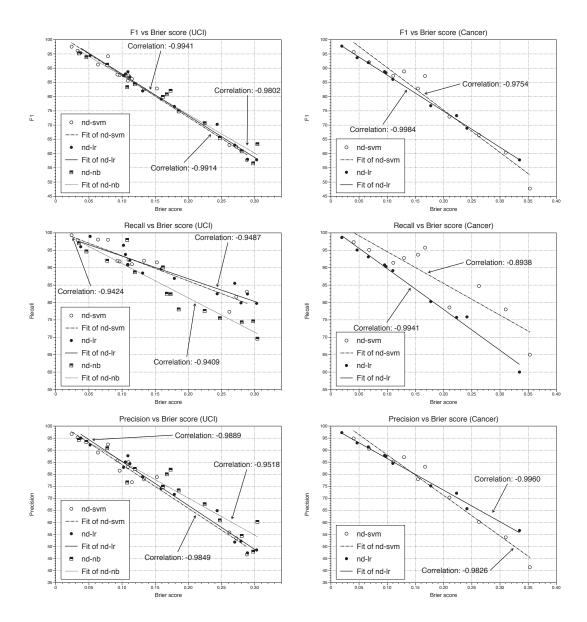


Figure 2: Correlation between Brier scores and F_1 , Recall, and Precision. The left column shows the results with UCI data sets, while the right column uses cancer data sets. Similar results would be achieved if we compared nondeterministic scores with deterministic accuracy

We observed that the correlations between the Brier scores of deterministic learners and non-deterministic scores (F_1 , Recall, and Precision) are very high: their absolute values are in all cases greater than 0.89. Therefore, in order to choose a nondeterministic approach in a practical application, given a data set, it would be recommendable to first analyze the Brier score of different deterministic learners.

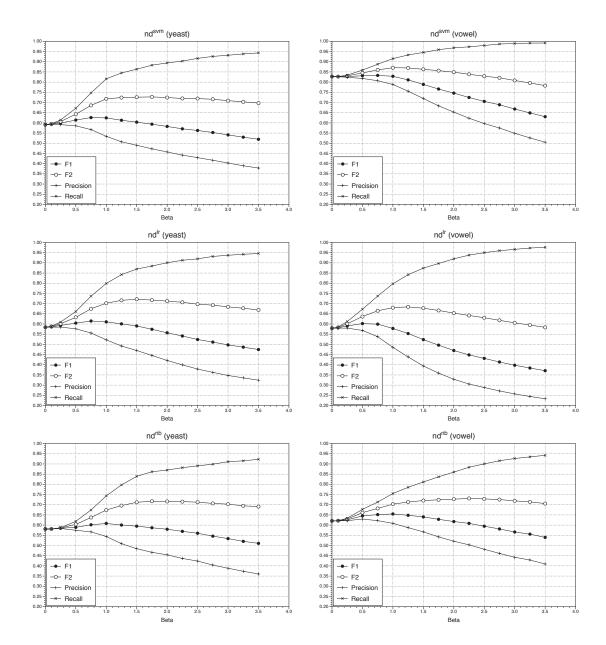


Figure 3: Evolution of F_1 , F_2 , *Precision* and *Recall* on two UCI data sets (*yeast* and *vowel*) for different β values and for the nondeterministic learners generated by *SVM*, *LR*, and *NB*

5.5 The Meaning of β

In this subsection, we analyze from the point of view of the user the role played by the parameter β in Algorithm 1. Its theoretical aim is to control the size of predictions: as the β value increases, the size of predictions will become bigger and therefore the *Recall* scores will be higher; see Equation (8). The problem is that it is not always of interest to increase *Recall* values, since that would worsen F_1 scores: adding more classes in predictions increases incorrect answers.

In Figure 3 we show the evolution of F_1 , F_2 , *Precision* and *Recall* on two UCI data sets (*yeast* and *vowel*) for different β values and for the nondeterministic learners generated by *SVM*, *LR*, and

NB. Quite similar graphs could have been generated for the other data sets used in the experiments reported in this section.

Initially, $\beta=0$ makes the nondeterministic classifiers deterministic. Therefore, the scores represented in the left-hand side of all the graphs in Figure 3 are all the same: the accuracy of the deterministic classifier. As β values become higher, the *Recall* increases and the *Precision* decreases. The main goal of the learning method proposed here is to look for a tradeoff of these measures that is determined by β , a user-modifiable parameter.

In practice, the value of β that the classifier must aim to optimize should be fixed by an expert in the field of application in which the classifier is going to be employed. The kind of decisions that one would like to take from nondeterministic classifications must be considered.

It can be observed in the graphs in Figure 3 that the best scores in F_1 are not always achieved for $\beta = 1$. With small values of β , F_1 increases. However, when some point near 1 is exceeded, the F_1 score of the nondeterministic learner typically falls below the accuracy of the corresponding deterministic learner. Nonetheless, optimal values are frequently reached around the *nominal* value: $\beta = 1$ (or 2 respectively). Slight improvements can be achieved in F_1 (in general F_β) if we use a grid search for β values to be used in Algorithm 1.

6. Conclusions

We have studied classifiers that are allowed to predict more than one class for entries from an input space: nondeterministic or set-valued classifiers. Using a clear analogy with Information Retrieval, we have proposed a family of loss functions based on F_{β} measures. After discussing such measures, we derived an algorithm to learn optimal nondeterministic hypothesis. Given an entry from the input space, the algorithm requires the posterior probabilities to compute the subset of classes with the lowest expected loss.

The paper includes a set of experiments carried out on two collections of data sets. The first one was downloaded from the UCI repository, the classes of which are not linearly separable. The second group is formed by data sets whose input spaces represent microarray expressions of different kinds of cancer, the classes of which are separable.

Using these benchmarks, we first compared nondeterministic learners obtained from a Naïve Bayes with those learned by a state-of-the-art set-valued (nondeterministic) algorithm, the Naïve Credal Classifier (NCC) (Zaffalon, 2002; Corani and Zaffalon, 2008a,b), an extension of the traditional Naïve Bayes classifier designed to return robust set-valued classifications. We showed that, using the loss measures defined in this paper, our method can improve the performance of NCC. Additionally, an important advantage of our nondeterministic classifiers over NCC is that we can control the degree of nondeterministic behavior. We can regulate the number of classes predicted by fixing the F_{β} to be optimized: as β is higher (the weight of Recall is increased in the harmonic average F_{β}), the size of our predictions grows (see Section 5.5). However the nondeterministic behavior of NCC is quite difficult to predict.

In addition to Naïve Bayes, we used a multiclass SVM and a Logistic Regression. With the posterior probabilities provided by these deterministic learners, we built another alternative method to predict more than one class: the set of classes which the highest posterior probabilities summing more than a threshold λ . We also found that the classifiers built with our algorithm outperform this option based on a threshold.

On the other hand, in the experiments reported in this paper, we studied the role of the deterministic learners that explicitly provide posterior probabilities. We found that the better the posterior probabilities, the better the nondeterministic classifiers. In fact we obtained very high correlations between the Brier scores of deterministic probabilities and the F_1 , *Precision* and *Recall* values of their nondeterministic counterparts.

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References

- J. Alonso, J. J. del Coz, J. Díez, O. Luaces, and A. Bahamonde. Learning to predict one or more ranks in ordinal regression tasks. *Proceedings of the European Conference on Machine Learning* and Principles and Practice of Knowledge Discovery in Databases (ECML PKDD'08), LNAI 5211, pages 39–54. Springer, 2008.
- S.A. Armstrong, J.E. Staunton, L.B. Silverman, R. Pieters, M.L. den Boer, M.D. Minden, S.E. Sallan, E.S. Lander, T.R. Golub, and S.J. Korsmeyer. MLL translocations specify a distinct gene expression profile that distinguishes a unique leukemia. *Nature Genetics*, 30(1):41–47, 2002.
- A. Asuncion and D.J. Newman. UCI machine learning repository. *School of Information and Computer Sciences. University of California, Irvine, California, USA*, 2007.
- P.L. Bartlett and M.H. Wegkamp. Classification with a reject option using a hinge loss. *Journal of Machine Learning Research*, 9:1823–1840, 2008.
- G.W. Brier. Verification of forecasts expressed in terms of probability. *Monthly Weather Rev*, 78: 1–3, 1950.
- C. Chow. On optimum recognition error and reject tradeoff. *IEEE Transactions on Information Theory*, 16(1):41–46, 1970.
- A. Clare and R.D. King. Predicting gene function in Saccharomyces cerevisiae. *Bioinformatics*, 19 (2):42–49, 2003.
- G. Corani and M. Zaffalon. Learning reliable classifiers from small or incomplete data sets: The Naive Credal Classifier 2. *Journal of Machine Learning Research*, 9:581–621, 2008a.
- G. Corani and M. Zaffalon. JNCC2: The java implementation of Naive Credal Classifier 2. *Journal of Machine Learning Research (Machine Learning Open Source Software)*, 9:2695–2698, 2008b.
- J. Demšar. Statistical comparisons of classifiers over multiple data sets. *Journal of Machine Learning Research*, 7:1–30, 2006.

- H.P. Kriegel, P. Kroger, A. Pryakhin, and M. Schubert. Using support vector machines for classifying large sets of multi-represented objects. *Proc. 4th SIAM Int. Conf. on Data Mining*, pages 102–114, 2004.
- C-J. Lin, R. C. Weng, and S. S. Keerthi. Trust region newton method for logistic regression. *Journal of Machine Learning Research*, 9(Apr):627–650, 2008.
- S. L. Pomeroy, P. Tamayo, M. Gaasenbeek, L. M. Sturla, M. Angelo, M. E. McLaughlin, J. Y. H. Kim, L. C. Goumnerova, P. M. Black, C. Lau, J. C. Allen, D. Zagzag, J. M. Olson, T. Curran, C. Wetmore, J. A. Biegel, T. Poggio, S. Mukherjee, R. Rifkin, A. Califano, G. Stolovitzky, D. N. Louis, J. P. Mesirov, E. S. Lander, and T. R. Golub. Prediction of central nervous system embryonal tumour outcome based on gene expression. *Nature*, 415(6870):436–442, 2002.
- S. Ramaswamy, P. Tamayo, R. Rifkin, S. Mukherjee, C.H. Yeang, M. Angelo, C. Ladd, M. Reich, E. Latulippe, J.P. Mesirov, et al. Multiclass cancer diagnosis using tumor gene expression signatures. *Proceedings of the National Academy of Sciences (PNAS)*, 98(26):15149–15154, 2001.
- D.T. Ross, U. Scherf, M.B. Eisen, C.M. Perou, C. Rees, P. Spellman, V. Iyer, S.S. Jeffrey, M. Van de Rijn, M. Waltham, et al. Systematic variation in gene expression patterns in human cancer cell lines. *Nature Genetics*, 24(3):227–234, 2000.
- G. Shafer and V. Vovk. A tutorial on conformal prediction. *Journal of Machine Learning Research*, 9:371–421, 2008.
- J.E. Staunton, D.K. Slonim, H.A. Coller, P. Tamayo, M.J. Angelo, J. Park, U. Scherf, J.K. Lee, W.O. Reinhold, J.N. Weinstein, et al. Chemosensitivity prediction by transcriptional profiling. Proceedings of the National Academy of Sciences (PNAS), 98(19):10787–10792, 2001.
- A.I. Su, J.B. Welsh, L.M. Sapinoso, S.G. Kern, P. Dimitrov, H. Lapp, P.G. Schultz, S.M. Powell, C.A. Moskaluk, H.F. Frierson, and G. M. Hampton. Molecular classification of human carcinomas by use of gene expression signatures. *Cancer Research*, 61(20):7388–7393, 2001.
- P. Tamayo, D. Scanfeld, B.L. Ebert, M.A. Gillette, C.W.M. Roberts, and J.P. Mesirov. Metagene projection for cross-platform, cross-species characterization of global transcriptional states. *Proceedings of the National Academy of Sciences (PNAS)*, 104(14):5959–5964, 2007.
- A.C. Tan, D.Q. Naiman, L. Xu, R.L. Winslow, and D. Geman. Simple decision rules for classifying human cancers from gene expression profiles. *Bioinformatics*, 21(20):3896–3904, 2005.
- R. Tibshirani and T. Hastie. Margin trees for high-dimensional classification. *Journal of Machine Learning Research*, 8:637–652, 2007.
- G. Tsoumakas and I. Katakis. Multi-label classification: An overview. *International Journal of Data Warehousing and Mining*, 3(3):1–13, 2007.
- T.-F. Wu, C.-J. Lin, and R. C. Weng. Probability estimates for multi-class classification by pairwise coupling. *Journal of Machine Learning Research*, 5:975–1005, August 2004.

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- E.J. Yeoh, M.E. Ross, S.A. Shurtleff, W.K. Williams, D. Patel, R. Mahfouz, F.G. Behm, S.C. Raimondi, M.V. Relling, A. Patel, C. Cheng, D. Campana, D. Wilkins, X. Zhou, J. Li, H. Liu, C.-H. Pui, W. E. Evans, C. Naeve, L. Wong, and J. R. Downing. Classification, subtype discovery, and prediction of outcome in pediatric acute lymphoblastic leukemia by gene expression profiling. *Cancer Cell*, 1(2):133–143, 2002.
- K.Y. Yeung and R.E. Bumgarner. Multiclass classification of microarray data with repeated measurements: application to cancer. *Genome Biology*, 4(12):R83, 2003.
- K.Y. Yeung, R.E. Bumgarner, and A.E. Raftery. Bayesian model averaging: development of an improved multiclass, gene selection and classification tool for microarray data. *Bioinformatics*, 21(10):2394–2402, 2005.
- M. Zaffalon. The Naïve Credal Classifier. *Journal of Statistical Planning and Inference*, 105(1): 5–21, 2002.