An Empirical Study of Lazy Multilabel Classification Algorithms

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Abstract. Multilabel classification is a rapidly developing field of machine learning. Despite its short life, various methods for solving the task of multilabel classification have been proposed. In this paper we focus on a subset of these methods that adopt a lazy learning approach and are based on the traditional k-nearest neighbor (kNN) algorithm. Two are our main contributions. Firstly, we implement BRkNN, an adaptation of the kNN algorithm for multilabel classification that is conceptually equivalent to using the popular Binary Relevance problem transformation method in conjunction with the kNN algorithm, but much faster. We also identify two useful extensions of BRkNN that improve its overall predictive performance. Secondly, we compare this method against two other lazy multilabel classification methods, in order to determine the overall best performer. Experiments on different real-world multilabel datasets, using a variety of evaluation metrics, expose the advantages and limitations of each method with respect to specific dataset characteristics.

1 Introduction

Traditional single-label classification is concerned with learning from a set of examples that are associated with a single label λ from a set of disjoint labels L, |L| > 1. If |L| = 2, then the learning task is called *binary* classification, while if |L| > 2, then it is called *multi-class* classification. In *multilabel* classification, each example is associated with a set of labels $Y \subseteq L$.

Multilabel classification methods can be categorized into two different groups [1]: i) problem transformation methods, and ii) algorithm adaptation methods. The first group of methods are algorithm independent. They transform the multilabel classification task into one or more single-label classification, regression or label ranking tasks. The second group of methods extend specific learning algorithms in order to handle multilabel data directly.

In this paper we focus on lazy multilabel classification methods of both categories that are based on the k Nearest Neighbor (kNN) algorithm. Among the strong points of these methods is that their time complexity scales linearly with respect to |L|. Furthermore, their main computationally intensive operation is the calculation of nearest neighbors, which is actually independent of |L|.

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Two are our main contributions in this work. Firstly, we implement BRkNN, an adaptation of the kNN algorithm for multilabel classification that is conceptually equivalent to using the popular Binary Relevance problem transformation method in conjunction with the kNN algorithm, but |L| times faster. We also identify two useful extensions of BRkNN that improve its overall predictive performance. Secondly, we compare this method against two other lazy multilabel classification methods, in order to determine the overall best performer.

The rest of this paper is structured as follows. Section 2 presents the BRkNN method and its extensions. Section 3 presents the setup of the experimental work and Section 4 discusses the results. Finally, Section 5 concludes this work.

2 BRkNN and Extensions

Binary Relevance (BR) is the most widely-used problem transformation method for multilabel classification. It learns one binary classifier $h_{\lambda} : X \to \{\neg \lambda, \lambda\}$ for each different label $\lambda \in L$. BR transforms the original data set into |L| data sets D_{λ} that contain all examples of the original data set, labeled as λ if the labels of the original example contained λ and as $\neg \lambda$ otherwise. It is the same solution used in order to deal with a multi-class problem using a binary classifier, commonly referred to as one-against-all or one-versus-rest.

BRkNN is an adaptation of the kNN algorithm that is conceptually equivalent to using BR in conjunction with the kNN algorithm. Therefore, instead of implementing BRkNN, we could have utilized existing implementations of BR [2] and kNN [3]. However, the problem in pairing BR with kNN is that it will perform |L| times the same process of calculating the k nearest neighbors. To avoid these redundant time-intensive computations, BRkNN extends the kNN algorithm so that independent predictions are made for each label, following a single search of the k nearest neighbors. This way BRkNN is |L| times faster than BR plus kNN during testing, a fact that could be crucial in domains with a large set of labels and requirements for low response times. BRkNN was implemented within the MULAN multilabel classification software [2].

We propose two extensions to the basic BRkNN algorithm. Both are based on the calculation of *confidence* scores for each label $\lambda \in L$ from BRkNN. The confidence for a label can be easily obtained by considering the percentage of the k nearest neighbors that include it. Formally, let $Y_j, j = 1 \dots k$, be the label sets of the k nearest neighbors of a new instance x. The confidence c_{λ} of a label $\lambda \in L$ is equal to:

$$c_{\lambda} = \frac{1}{k} \sum_{j=1}^{k} I_{Y_j}(\lambda)$$

where $I_{Y_j} : L \to \{0, 1\}$ is a function that outputs 1 if its input label λ belongs to set Y_j and 0 otherwise, called *indicator function* in set theory.

The first extension of BRkNN, called BRkNN-a, checks whether BRkNN outputs the empty set, due to none of the labels $\lambda \in L$ being included in at least

half of the k nearest neighbors. If this condition holds, then it outputs the label with the highest confidence. It so deals with a general disadvantage of BR, that has not been raised in the past: as each label is independently predicted in BR, there exists a possibility that the empty set is given as the overall output. We hypothesize that better results will be obtained through the proposed extension that outputs the most probable label when this phenomenon arises.

The second extension of BR*k*NN, called BR*k*NN-b calculates the average size s of the label sets of the k nearest neighbors at a first step, $s = \frac{1}{k} \sum_{j=1}^{k} |Y_j|$, and then outputs the [s] (nearest integer of s) labels with the highest confidence.

3 Experimental Setup

3.1 Datasets

We experiment with 3 datasets from 3 different application domains: The biological dataset *yeast* [4] is concerned with protein function classification. The image dataset *scene* [5] is concerned with semantic indexing of still scenes. The music dataset *emotions* [6] is concerned with the classification of songs according to the emotions they evoke.

Table 1 shows certain standard statistics of these datasets, such as the number of examples in the train and test sets, the number of numeric and discrete attributes and the number of labels, along with multilabel data statistics, such as the number of distinct label subsets, the label cardinality and the label density [1]. Label cardinality is the average number of labels per example, while label density is the same number divided by |L|.

Dataset	Examples	Attri Numeric	butes Discrete	Labels	Distinct Subsets	Label Cardinality	Label Density
scene	2712	294	0	6	15	1.074	0.179
emotions	593	72	0	6	27	1.868	0.311
yeast	2417	103	0	14	198	4.327	0.302

Table 1. Standard and multilabel statistics for the data sets used in the experiments

3.2 Evaluation Methodology

We perform two sets of experiments. In the first one, we compare BRkNN to its extensions. In the second one, we compare the best version of BRkNN in each dataset to two other lazy multilabel classification methods, LPkNN and MLkNN, in order to make a final recommendation.

LPkNN is simply the pairing of the Label Powerset (LP) problem transformation method [2] with the kNN algorithm. LP considers each different subset of Lthat appears in the training set as a different label of a single-label classification task. LPkNN has not been discussed in the related literature to the best of our knowledge. MLkNN [7] is another adaptation of the kNN algorithm for multilabel data. What mainly differentiates this method from BRkNN is the use of prior and posterior probabilities which are directly estimated from the training set based on frequency counting. We implemented MLkNN in Java within the MULAN multilabel classification software [2] for the purposes of this study.

Each method was executed with a varying number of nearest neighbors. Specifically, the parameter k ranged from 1 to 30. The performance of each method for each k was evaluated using 10-fold cross-validation, in order to obtain an accurate performance estimate. In each fold, the following metrics were calculated [2], and eventually averaged over all folds:

- Example-based. Hamming loss, accuracy, F-measure and subset accuracy
- Label-based. Micro and macro version of F-measure

4 Experimental Results

4.1 Do the Proposed Extensions Improve BRkNN?

In this subsection we investigate whether BRkNN-a and BRkNN-b improve the performance of BRkNN. Table 2 reports the average performance of the three algorithms across all 30 values of the k parameter for each dataset. It presents results for all evaluation metrics mentioned in Section 3.2. The best result on each metric and dataset is shown with bold typeface. The last line contains for each algorithm the number of metrics for which it achieves the best result, while within parentheses there is the number of metrics for which BRkNN-a and BRkNN-b are better than the base BRkNN algorithm.

The results show that both extensions outperform the base BRkNN method in more than half of the 6 metrics on all datasets. BRkNN-a outperforms BRkNNin 6, 5 and 5 out of the 6 metrics in the scene, emotions and yeast datasets respectively. BRkNN-b outperforms BRkNN in 6, 4 and 4 out of the 6 metrics in the scene, emotions and yeast datasets respectively. These two pieces of evidence strongly support that both BRkNN-a and BRkNN-b are beneficial extensions.

Studying the performance of the algorithms at each individual dataset, we notice that BRkNN-a dominates in scene and emotions, while BRkNN-b dominates

	scene				emotions	3	yeast		
metric	base	ext-a	$\operatorname{ext-b}$	base	ext-a	ext-b	base	ext-a	ext-b
Hamming loss	0.0950	0.0938	0.0941	0.1976	0.1982	0.2175	0.1974	0.1975	0.2082
accuracy	0.6256	0.7226	0.7218	0.5215	0.5441	0.5430	0.5062	0.5080	0.5346
F-measure	0.6386	0.7392	0.7381	0.6275	0.6576	0.6590	0.5777	0.5795	0.6652
subset accuracy	0.5993	0.6889	0.6886	0.2895	0.2971	0.2759	0.1958	0.1959	0.1766
micro F-measure	0.6964	0.7296	0.7284	0.6499	0.6577	0.6509	0.6374	0.6380	0.6567
macro F-measure	0.6955	0.7363	0.7349	0.6224	0.6303	0.6294	0.3926	0.3931	0.4261
#wins (#better)	0	6(6)	0(6)	1	4(5)	1(4)	1	1(5)	4 (4)

Table 2. Experimental results of BRkNN, BRkNN-a and BRkNN-b on all datasets, averaged for all k

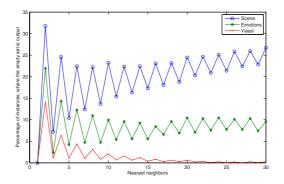


Fig. 1. Percentage of new instances, where BRkNN outputs the empty set (y axis), with respect to the number of nearest neighbors (k) (x axis) for all datasets

in yeast. This performance pattern correlates with the cardinality of the datasets, which is 1.074, 1.868 and 4.327 for the scene, emotions and yeast dataset respectively (see Table 3.1). Actually, it is natural for datasets of low cardinality, such as scene and emotions, to favor BRkNN-a over BRkNN, because the probability that the latter outputs the empty set increases in such datasets. This is clearly shown in Figure 1, which plots the percentage of the instances, where BRkNN outputs the empty set, for various values of the k parameter. BRkNN-a deals with exactly this problem of BRkNN. On the other hand, BRkNN-b works better in datasets with larger cardinality, as it includes a mechanism to predict the number of true labels associated with a new instance.

4.2 Comparison of BRkNN, LPkNN and MLkNN

Table 3 reports the average performance of the three algorithms across all 30 values of the k parameter for each dataset. It presents results for all evaluation metrics mentioned in Section 3.2. The best result on each metric and dataset is shown with bold typeface. The last line contains for each algorithm the number of metrics for which it achieves the best result.

	scene			emotions			yeast		
metric	BR-a	LP	ML	BR-a	LP	ML	BR-b	LP	ML
Hamming loss	0.0938	0.0955	0.0884	0.1982	0.2094	0.2003	0.2082	0.2143	0.1950
accuracy	0.7226	0.7181	0.6720	0.5441	0.5600	0.5233	0.5346	0.5280	0.5105
F-measure	0.7392	0.7343	0.6944	0.6576	0.6662	0.6352	0.6652	0.6375	0.5823
subset accuracy	0.6889	0.6854	0.6272	0.2971	0.3287	0.2780	0.1766	0.2452	0.1780
micro F-measure	0.7296	0.7249	0.7316	0.6577	0.6649	0.6509	0.6567	0.6415	0.6422
macro F-measure	0.7363	0.7323	0.7341	0.6303	0.6505	0.6110	0.4261	0.4322	0.3701
#wins	4	0	2	1	5	0	3	2	1

Table 3. Experimental results of best version of BRkNN, LPkNN and MLkNN with normalization on all datasets, averaged for all k

We notice that BRkNN-a and LPkNN dominate in the scene and emotions datasets respectively, while in the yeast dataset there is no clear winner. However BRkNN-b performs better in most measures, followed by LPkNN and finally MLkNN. There is no apparent explanation on why LPkNN performs better in the emotions dataset. We notice in Table 3.1 that this dataset has the highest label density, while the scene dataset where LPkNN has the worst performance has the lowest label density. However we cannot safely argue that high density datasets lead to improved performance of the LPkNN algorithm.

5 Conclusions

This paper has studied how the k Nearest Neighbor (kNN) algorithm is used for the classification of multilabel data. It presented BRkNN, an efficient implementation of the pairing of BR with kNN, along with two interesting extensions. Experimental results indicated that the proposed extensions are in the right direction. In addition, the paper compared experimentally BRkNN with two other methods (LPkNN and MLkNN) and reached to some interesting conclusions as to what kind of evaluation metrics and what kind of datasets are well-suited to the different methods.

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