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Pattern Recognition 00 (2011) 1–33

Pattern
Recognition

An extensive experimental comparison of methods for multi-label learning

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Abstract

Multi-label learning has received significant attention in the research community over the past few years: this has resulted in the development of a variety of multi-label learning methods. In this paper, we present an extensive experimental comparison of 12 multi-label learning methods using 16 evaluation measures over 11 benchmark datasets. We selected the competing methods based on their previous usage by the community, the representation of different groups of methods and the variety of basic underlying machine learning methods. Similarly, we selected the evaluation measures to be able to assess the behavior of the methods from a variety of view-points. In order to make conclusions independent from the application domain, we use 11 datasets from different domains. Furthermore, we compare the methods by their efficiency in terms of time needed to learn a classifier and time needed to produce a prediction for an unseen example. We analyze the results from the experiments using Friedman and Nemenyi tests for assessing the statistical significance of differences in performance. The results of the analysis show that for multi-label classification the best performing methods overall are random forests of predictive clustering trees (RF-PCT) and hierarchy of multi-label classifiers (HOMER), followed by binary relevance (BR) and classifier chains (CC). Furthermore, RF-PCT exhibited the best performance according to all measures for multi-label ranking. The recommendation from this study is that when new methods for multi-label learning are proposed, they should be compared to RF-PCT and HOMER using multiple evaluation measures.

Keywords: multi-label ranking, multi-label classification, comparison of multi-label learning methods

1. Introduction

The problem of single-label classification is concerned with learning from examples, where each example is associated with a single label λ_i from a finite set of disjoint labels $\mathcal{L} = \{\lambda_1, \lambda_2, \dots, \lambda_Q\}$, $Q > 1$. For $Q > 2$, the learning problem is referred to as *multi-class classification*. On the other hand, the task of learning a mapping from an example $\mathbf{x} \in \mathcal{X}$ (\mathcal{X} denotes the domain of examples) to a set of labels $\mathcal{Y} \subseteq \mathcal{L}$ is referred to as a *multi-label classification*. In contrast to multi-class classification, alternatives in multi-label classification are not assumed to be mutually exclusive: multiple labels may be associated with a single example, i.e., each example can be a member of more than one class. Labels in the set \mathcal{Y} are called relevant, while the labels in the set $\mathcal{L} \setminus \mathcal{Y}$ are irrelevant for a given example.

Besides the concept of multi-label classification, multi-label learning introduces the concept of *multi-label ranking* [1]. Multi-label ranking can be considered as a generalization of multi-class classification, where instead of predicting

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only a single label (the top label), it predicts the ranking of all labels. In other words, multi-label ranking is understood as learning a model that associates a query example \mathbf{x} both with a ranking of the complete label set and a bipartition of this set into relevant and irrelevant labels.

The issue of learning from multi-label data has recently attracted significant attention from many researchers, motivated by an increasing number of new applications. The latter include semantic annotation of images and video (news clips, movies clips), functional genomics (gene and protein function), music categorization into emotions, text classification (news articles, web pages, patents, emails, bookmarks, ...), directed marketing and others. In the last few years, several workshops have been organized and journal special issues edited covering the topic of multi-label learning.

In recent years, many different approaches have been developed to solving multi-label learning problems. Tsoumakas and Katakis [2] summarize them into two main categories: a) algorithm adaptation methods, and b) problem transformation methods. Algorithm adaptation methods extend specific learning algorithms to handle multi-label data directly. Examples include lazy learning [3] [4] [5], neural networks [6] [7], boosting [8] [9], classification rules [10], decision trees [11] [12] etc. Problem transformation methods, on the other hand, transform the multi-label learning problem into one or more single-label classification problems. The single-label classification problems are solved with a commonly used single-label classification approach and the output is transformed back into a multi-label representation. A common approach to problem transformation is to use class binarization methods, i.e., decompose the problem into several binary sub-problems that can then be solved by using a binary base classifier. The simplest strategies in the multi-label setting are the one-against-all and one-against-one strategies, also referred to as the binary relevance method [2] and pair-wise method [13] [14] respectively.

In this study, we extend this categorization of multi-label methods with a third group of methods, namely, ensemble methods. This group of methods consists of methods that use ensembles to make multi-label predictions and their base classifiers belong to either problem transformation or algorithm adaptation methods. Methods that belong to this group are RAKEL [15], ensembles of classifier chains (ECC) [16], random forests of predictive clustering trees [17] [18] and random forests of multi-label C4.5 trees [11].

As new methods for multi-label learning are proposed, they are experimentally compared to existing methods. The typical experimental evaluation compares the proposed method to a few existing ones on a few datasets. The methods are compared on performance in terms of one or a few error metrics and the comparison typically shows that the proposed method outperforms the other methods on some of the considered datasets and metrics. It is worth noting that a significant number of metrics has also been proposed for evaluating the performance of multi-label methods, which can concern the classification or ranking variant of the problem.

The number of proposed methods, datasets and metrics for multi-label learning constantly increases. As the research area of multi-label learning matures, there is a strong need for a comprehensive overview of methods and metrics. The need for a wider, extensive, and un-biased experimental comparison of multi-label learning methods is even stronger. It is this need that we address with the present paper.

In this study, we experimentally evaluate 12 methods for multi-label learning using 16 evaluation measures over 11 benchmark datasets. The multi-label methods comprise 3 algorithm adaptation methods, 5 problem transformation methods and 4 ensemble methods. The benchmark datasets are from five application domains: 2 from image classification, 1 from gene function prediction, 6 from text classification, 1 from music classification and 1 from video classification. The predictive performance of the methods is assessed using 6 example-based measures, 6 label-based measures and 4 ranking-based measures. Furthermore, we assess the efficiency of the methods by measuring their training and testing times. The large number of methods, datasets and evaluation measures is enabling us to draw some more general conclusions and to perform an unbiased assessment of the predictive performance of the multi-label methods.

The results from our extensive experimental evaluation will facilitate further research on multi-label learning as follows. First, this study will provide the research community with a better insight about the predictive performance of the methods currently available in the literature. Second, this study will identify a few methods that should be further used by the research community as benchmarks to compete against when proposing new methods. Third, this study uses a diverse collection of publicly available datasets that can be reused by other researchers as benchmark datasets for multi-label learning. Finally, this study will highlight the advantages of certain methods for certain types of datasets.

The remainder of this paper is organized as follows. Section 2 defines the tasks of multi-label classification

and label ranking and surveys the related work. The state-of-the-art methods for multi-label learning used in the experimental evaluation are presented in Section 3. Section 4 describes the multi-label problems, the evaluation measures and the experimental setup, while Section 5 presents and discusses the experimental results. Finally, the conclusions are given in Section 6.

2. Background

In this section, we present the task of multi-label learning and methods for solving it. We begin by a formal definition of the task of multi-label learning. We then present an overview of the methods for multi-label learning.

2.1. The task of multi-label learning

Multi-label learning is concerned with learning from examples, where each example is associated with multiple labels. These multiple labels belong to a predefined set of labels. Depending on the goal, we can distinguish two types of tasks: multi-label classification and multi-label ranking. In the case of multi-label classification, the goal is to construct a predictive model that will provide a list of relevant labels for a given, previously unseen example. On the other hand, the goal in the task of multi-label ranking is to construct a predictive model that will provide, for each unseen example, a list of preferences (i.e., a ranking) of the labels from the set of possible labels.

We define the task of multi-label learning as follows:

Given:

- An example space \mathcal{X} that consists of tuples of values of primitive data types (boolean, discrete or continuous), i.e., $\forall \mathbf{x}_i \in \mathcal{X}, \mathbf{x}_i = (x_{i1}, x_{i2}, \dots, x_{iD})$, where D is the size of the tuple (or number of descriptive attributes),
- a label space $\mathcal{L} = \{\lambda_1, \lambda_2, \dots, \lambda_Q\}$ which is a tuple of Q discrete variables (with values 0 or 1),
- a set of examples E , where each example is a pair of tuples from the example and label space, respectively, i.e., $E = \{(\mathbf{x}_i, \mathcal{Y}_i) | \mathbf{x}_i \in \mathcal{X}, \mathcal{Y}_i \in \mathcal{L}, 1 \leq i \leq N\}$ and N is the number of examples of E ($N = |E|$), and
- a quality criterion q , which rewards models with high predictive accuracy and low complexity.

If the task at hand is multi-label classification, then the goal is to

Find: a function $h: \mathcal{X} \rightarrow 2^{\mathcal{L}}$ such that h maximizes q .

On the other hand, if the task is multi-label ranking, then the goal is to

Find: a function $f: \mathcal{X} \times \mathcal{L} \rightarrow \mathcal{R}$, such that f maximizes q , where \mathcal{R} is the ranking of the labels for a given example.

2.2. An overview of methods for multi-label learning

Tsoumakas and Katakis [2] have presented the first overview of methods for multi-label learning where the methods for multi-label learning are divided into two categories: algorithm adaptation and problem transformation methods. There, three problem transformation methods were evaluated on a small empirical study (three datasets). In this study, we perform an extensive experimental evaluation of 12 methods for multi-label learning over 11 benchmark multi-label datasets using 16 evaluation measures. Furthermore, besides the two categories of methods for multi-label learning, we introduce a third category: ensemble methods. In the remainder of this section, we present the three categories of methods for multi-label learning: algorithm adaptation, problem transformation and ensemble methods.

2.2.1. Algorithm adaptation methods

The multi-label methods that adapt, extend and customize an existing machine learning algorithm for the task of multi-label learning are called algorithm adaptation methods. Here, we present multi-label methods proposed in the literature that are based on the following machine learning algorithms: boosting, k-nearest neighbors, decision trees and neural networks. The extended methods are able to directly handle multi-label data.

Boosting ADABOOST.MH and ADABOOST.MR [8] are two extensions of ADABOOST for multi-label data. While ADABOOST.MH is designed to minimize Hamming loss, ADABOOST.MR is designed to find a hypothesis which ranks the correct labels at the top. Furthermore, ADABOOST.MH can also be combined with an algorithm for producing alternating decision trees [9]. The resulting multi-label models of this combination can be interpreted by humans.

k-Nearest neighbors Several variants for multi-label learning (ML-kNN) of the popular k Nearest Neighbors (kNN) lazy learning algorithm have been proposed [3] [4] [5]. The retrieval of the k nearest neighbors is the same as in the traditional k -NN algorithm. The main difference is the determination of the label set of a test example. Typically, these algorithms use prior and posterior probabilities of each label within the k nearest neighbors. Cheng et al. [19] have proposed a hybrid method that uses logistic regression and k-nearest neighbors.

Decision trees Clare et al. [11] adapted the C4.5 algorithm for multi-label data (ML-C4.5) by modifying the formula for calculating entropy. Blockeel et al. [12] proposed the concept of predictive clustering trees (PCTs). PCTs have been used for predicting tuples of variables, predicting time series and predicting classes organized into a hierarchy or a directed acyclic graph. However, they can also be used in the context of multi-label learning, where each label is a component of the target tuple.

Neural networks Neural networks have also been adapted for multi-label classification [6] [7]. BP-MLL [7] is an adaptation of the popular back-propagation algorithm for multi-label learning. The main modification to the algorithm is the introduction of a new error function that takes multiple labels into account.

Support vector machines Elisseeff and Weston [20] have proposed a ranking approach for multi-label learning that is based on SVMs. The cost function they use is the average fraction of incorrectly ordered pairs of labels.

2.2.2. Problem transformation methods

The problem transformation methods are multi-label learning methods that transform the multi-label learning problem into one or more single-label classification or regression problems. For smaller single-label problems, there exists a plethora of machine learning algorithms. Problem transformation methods can be grouped into three categories: binary relevance, label power-set and pair-wise methods.

Binary relevance methods The simplest strategy for problem transformation is to use the one-against-all strategy to convert the multi-label problem into several binary classification problems. This approach is known as the binary relevance method (BR) [2]. A method closely related to the BR method is the Classifier Chain method (CC) proposed by Read et al. [16]. This method involves Q binary classifiers linked along a chain. Godbole et al. [21] present algorithms which extend the SVM binary classifiers along two dimensions: training set extension and improvement of margin. With the first approach, the training set is extended with the predictions of the binary classifiers and then a new set of binary classifiers is trained on the extended dataset. For the second extension, Godbole et al. remove very similar negative training examples and remove the negative training examples of a complete class that are similar to the positive class.

Label power-set methods A second problem transformation method is the label combination method, or label power-set method (LP), which has been the focus of several recent studies [15] [22] [2]. The basis of these methods is to combine entire label sets into atomic (single) labels to form a single-label problem (i.e., single-class classification problem). For the single-label problem, the set of possible single labels represents all distinct label subsets from the original multi-label representation. In this way, LP based methods directly take into account the label correlations. However, the space of possible label subsets can be very large. To resolve this issue, Read [23] has developed a pruned problem transformation (PPT) method, that selects only the transformed labels that occur more than a predefined number of times. Another label power-set method is HOMER [24], which first constructs a hierarchy of the multiple labels and then constructs a classifier for the label sets in each node of the hierarchy.

Pair-wise methods A third problem transformation approach to solving the multi-label learning problem is pair-wise or round robin classification with binary classifiers [13] [14]. The basic idea here is to use $Q \cdot (Q - 1) / 2$ classifiers covering all pairs of labels. Each classifier is trained using the samples of the first label as positive examples and the samples of the second label as negative examples. To combine these classifiers, the pairwise classification method naturally adopts the majority voting algorithm. Given a test example, each classifier predicts (i.e., votes for) one of the two labels. After the evaluation of all $Q \cdot (Q - 1) / 2$ classifiers, the labels are ordered according to their sum of votes. A label ranking algorithm is then used to predict the relevant labels for each example. Besides majority voting in CLR, Park et al. [25] propose a more effective voting algorithm. It computes the class with the highest accumulated voting mass, while avoiding the evaluation of all possible pairwise classifiers. Mencia et al. [26] adapted the QWeighted approach to multi-label learning (QWML).

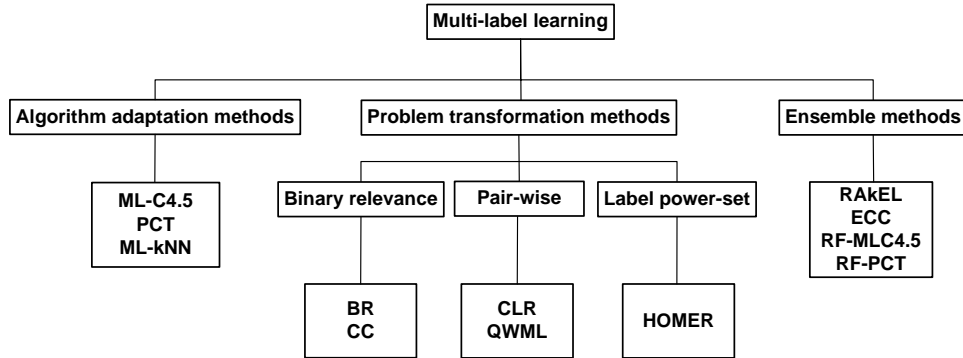


Figure 1: The multi-label learning methods used in this study divided into groups as discussed in the related work section above.

2.2.3. Ensemble methods

The ensemble methods for multi-label learning are developed on top of the common problem transformation or algorithm adaptation methods. The most well known problem transformation ensembles are the RAKEL system by Tsoumakas et al. [15], ensembles of pruned sets (EPS) [27] and ensembles of classifier chains (ECC) [16].

RAKEL constructs each base classifier by considering a small random subset of labels and learning a single-label classifier for the prediction of each element in the power-set of this subset. EPS uses pruning to reduce the computational complexity of label power-set methods, and an example duplication method to reduce the error rate as compared to label power-set and other methods. This method proved to be particularly competitive in terms of efficiency.

ECC are ensemble methods that have classifier chains (CC) as base classifiers. The final prediction is obtained by summing the predictions by label and then applying threshold for selecting the relevant labels. Note that binary methods are occasionally referred to as ensemble methods because they involve multiple binary models. However, none of these models is multi-label itself and therefore we use the term ensemble strictly in the sense of an ensemble of multi-label methods.

Algorithm adaptation ensemble methods are the ensembles whose base classifiers are themselves algorithm adaptation methods. An example of an algorithm adaptation ensemble method are the ensembles of predictive clustering trees (PCTs) [18]. These ensembles use PCTs for predicting tuples of variables as base classifiers. Each base classifier makes a multi-label prediction and then these predictions are combined by using some voting scheme (e.g., majority or probability distribution voting).

3. Methods for multi-label learning

In this section, we briefly introduce the state-of-the-art methods for multi-label learning that are used in this study. Figure 1 depicts how these methods are divided into groups using the categorization scheme from the related work section. In this study, we use one label power-set method, two binary relevance and two pair-wise transformation methods, two algorithm adaptation methods, and four ensemble methods. Moreover, one of the ensemble methods is label power-set based, while the other methods are algorithm adaptation based.

We also divide the used multi-label learning approaches based on the type of basic machine learning algorithm they use. The methods use three types of base algorithms: SVMs, decision trees and k-nearest neighbors. We show this categorization in Figure 2.

3.1. Binary relevance methods

Binary relevance (BR) [2] is the well known one-against-all strategy. It addresses the multi-label learning problem by learning one classifier for each label, using all the examples labeled with that label as positive examples and all remaining examples as negative. When making a prediction, each binary classifier predicts whether its label is

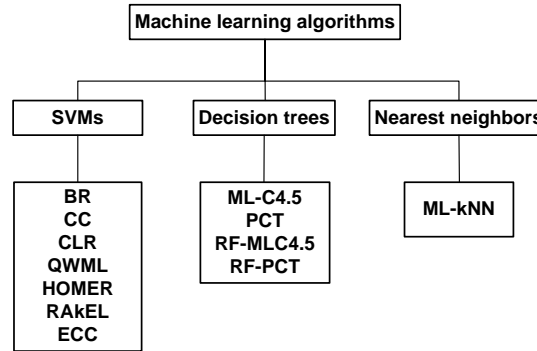


Figure 2: The multi-label learning methods used in this study divided into groups based on the base machine learning algorithm they use.

relevant for the given example or not, resulting in a set of relevant labels. In the ranking scenario, the labels are ordered according to the probability associated to each label by the respective binary classifier.

The *Classifier chaining method (CC)* [16] involves Q binary classifiers as in BR. Classifiers are linked along a chain where the i -th classifier deals with the binary relevance problem associated with label $\lambda_i \in L$, ($1 \leq i \leq Q$). The feature space of each link in the chain is extended with the 0/1 label associations of all previous links. The ranking and the prediction of the relevant labels in the CC method are the same as in the BR method.

3.2. Pair-wise methods

Calibrated label ranking (CLR) [25] is a technique for extending the common pair-wise approach to multi-label learning. It introduces an artificial (calibration) label λ_0 , which represents the split-point between relevant and irrelevant labels. The calibration label λ_0 is assumed to be preferred over all irrelevant labels, but all relevant labels are preferred over it. It is represented by the binary relevance classifiers, that are introduced as pair-wise classifiers in the context of pair-wise learning. At prediction time (majority voting is usually used), one will get a ranking over $Q + 1$ labels (the Q original labels plus the calibration label λ_0). CLR is considered a combination of multi-label classification and ranking.

The *Quick Weighted voting method* for multi-class classification, proposed by Park et al. [25], is a variant of the CLR method that introduces a more effective voting strategy than the majority voting used by the CLR method. Quick weighted voting exploits the fact that during voting some classes can be excluded from the set of possible top rank classes early in the process, when it becomes clear that even if they reach the maximal voting mass in the remaining evaluations they can not exceed the current maximum. Pairwise classifiers are selected depending on a voting loss value, which is the number of votes that a class has not received. The voting loss starts with a value of zero and increases monotonically with the number of performed preference evaluations. The class with the current minimal loss is the best candidate for the top ranked class. If all preferences involving this class have been evaluated (and it still has the lowest loss), it can be concluded that no other class can achieve a better ranking. Thus, the quick weighted algorithm always focuses on classes with low voting loss. The adaptation of quick weighted algorithm for multi-label learning (QWML) [26] is done by repeating the process while all relevant labels are not determined, i.e., until the returned label is the artificial label, which means that all remaining labels will be considered irrelevant.

3.3. Label power-set method

Hierarchy Of Multi-label classifiERs (HOMER) [24] is an algorithm for effective and computationally efficient multi-label learning in domains with a large number of labels. HOMER constructs a hierarchy of multi-label classifiers, each one dealing with a much smaller set of labels compared to Q (the total number of labels) and a more balanced example distribution. This leads to improved predictive performance and also to linear training and logarithmic testing complexities with respect to Q . One of the main processes within HOMER is the even distribution of a set

of labels into k disjoint subsets so that similar labels are placed together and dissimilar apart. The best predictive performance is reported using a balanced k means algorithm customized for HOMER [24]. HOMER is a computationally efficient multi-label classification method, specifically designed for large multi-label datasets.

3.4. Algorithm adaptation methods

Multi-Label C4.5 (ML-C4.5) [11] is an adaptation of the well known C4.5 algorithm for multi-label learning by allowing multiple labels in the leaves of the tree. Clare et al. [11] modified the formula for calculating entropy (see Equation 1) for solving multi-label problems. The modified entropy sums the entropies for each individual class label. The key property of ML-C4.5 is its computational efficiency.

$$\text{entropy}(E) = - \sum_{i=1}^N (p(c_i) \log p(c_i) + q(c_i) \log q(c_i)) \quad (1)$$

where E is the set of examples, $p(c_i)$ is the relative frequency of class label c_i and $q(c_i) = 1 - p(c_i)$.

Predictive clustering trees (PCTs) [12] are decision trees viewed as a hierarchy of clusters: the top-node corresponds to one cluster containing all data, which is recursively partitioned into smaller clusters while moving down the tree. PCTs are constructed using a standard top-down induction of decision trees algorithm, where the variance and the prototype function can be instantiated according to the task at hand. Namely, PCTs can handle several types of structured outputs: tuples of continuous or discrete variables, time series, classes organized into a hierarchy, tuples of time series and tuples of hierarchies [18]. For the task of predicting tuples of discrete variables, the variance function is computed as the sum of the *Gini indices* [28] of the variables from the target tuple, i.e., $\text{Var}(E) = \sum_{i=1}^T \text{Gini}(E, Y_i)$, $\text{Gini}(E, Y_i) = 1 - \sum_{j=1}^{C_i} p_{c_{ij}}$, where T is the number of target attributes, c_{ij} is the j -th class of target attribute Y_i and C_i is the number of classes of target attribute Y_i . The prototype function returns a vector of probabilities that an example belongs to a given class for each variable from the target tuple. In the case of multi-label learning, it returns a vector of probabilities that an example is labeled with a given label.

Multi-Label k-Nearest Neighbors (ML-kNN) [3] is an extension of the popular k-nearest neighbors (kNN) algorithm. Firstly, for each test example, its k nearest neighbors in the training set are identified. Then, according to statistical information gained from the label sets of these neighboring examples, i.e., the number of neighboring examples belonging to each possible label, the maximum *a posteriori* principle is used to determine the label set for the test example.

3.5. Ensemble methods

The *RANdom k-labELsets (RAkEL)* [15] is an ensemble method for multi-label classification. It draws m random subsets of labels with size k from all labels \mathcal{L} and trains a label power-set classifier using each set of labels. A simple voting process determines the final set of labels for a given example. In this way, the proposed algorithm aims to take into account label correlations using single-label classifiers that are applied on subtasks with a manageable number of labels and adequate number of examples per label.

Ensembles of Classifier Chains (ECC) [16] are an ensemble multi-label classification technique that uses classifier chains as a base classifier. ECC trains m CC classifiers C_1, C_2, \dots, C_m . Each C_k is trained with a random chain ordering (of \mathcal{L}) and a random subset of X . Hence each C_k model is likely to be unique and able to give different multi-label predictions. These predictions are summed per label so that each label receives a number of votes. A threshold is used to select the most popular labels which form the final predicted multi-label set.

Random Forest of Predictive Clustering Trees (RF-PCT) [17] [18] and *Random Forest of ML-C4.5 (RFML-C4.5)*¹ are ensembles that use PCTs and ML-C4.5 trees, respectively, as base classifiers. The diversity among the base classifiers is obtained by using bagging, and additionally by changing the feature set during learning [29]. More precisely, at each node in the decision trees, a random subset of the input attributes is taken, and the best feature is selected from this subset. The number of attributes that are retained is given by a function f of the total number of input attributes x (e.g., $f(x) = 1$, $f(x) = \sqrt{x}$, $f(x) = \lfloor 0.1 \cdot x + 1 \rfloor$, $f(x) = \lfloor \log_2(x) + 1 \rfloor \dots$). The predictions of the base classifiers are then combined using some voting scheme (typically, majority or probability distribution vote).

¹We have implemented the random forest of ML-C4.5 trees within the MULAN library for multi-label learning.

4. Experimental design

In this section, we present the experimental design used to compare the methods for multi-label learning. We first shortly describe the benchmark multi-label datasets. We then give a short overview of the evaluation measures typically applied to assess the predictive performance of methods for multi-label learning. Next, we present the specific setup and the instantiation of the parameters for the used methods for multi-label learning. Finally, we present the procedure for statistical evaluation of the experimental results.

4.1. Datasets

We use eleven different multi-label classification benchmark problems. Parts of the selected problems were used in various studies and evaluations of methods for multi-label learning. In the process of selection of problems, we opted to include benchmark datasets with different scale and from different application domains. Table 1 presents the basic statistics of the datasets. We can note that the datasets vary in size: from 391 up to 60000 training examples, from 202 up to 27856 testing examples, from 72 up to 2150 features, from 6 to 983 labels, and from 1.07 to 19.02 average number of labels per example (i.e., label cardinality [30]). From the literature, these datasets come pre-divided into training and testing parts: Thus, in the experiments, we use them in their original format. The training part usually comprises around 2/3 of the complete dataset, while the testing part the remaining 1/3 of the dataset.

Table 1: Description of the benchmark problems in terms of application domain (*domain*), number of training (*#tr.e.*) and test (*#t.e.*) examples, the number of features (*D*), the total number of labels (*Q*) and label cardinality (*l_c*). The problems are ordered by their overall complexity roughly calculated as $\#tr.e. \times D \times Q$.

	<i>domain</i>	<i>#tr.e.</i>	<i>#t.e.</i>	<i>D</i>	<i>Q</i>	<i>l_c</i>
emotions [31]	multimedia	391	202	72	6	1.87
scene [32]	multimedia	1211	1159	294	6	1.07
yeast [20]	biology	1500	917	103	14	4.24
medical [16]	text	645	333	1449	45	1.25
enron [33]	text	1123	579	1001	53	3.38
corel5k [34]	multimedia	4500	500	499	374	3.52
tmc2007 [35]	text	21519	7077	500	22	2.16
mediamill [36]	multimedia	30993	12914	120	101	4.38
bibtex [37]	text	4880	2515	1836	159	2.40
delicious [24]	text	12920	3185	500	983	19.02
bookmarks [37]	text	60000	27856	2150	208	2.03

The datasets come from three domains: biology, multimedia and text categorization. From the biological domain, we have the *yeast* dataset [20]. It is a widely used dataset, where genes are instances in the dataset and each gene can be associated with 14 biological functions (labels).

The datasets that belong to the multimedia domain are: *emotions*, *scene*, *corel5k* and *mediamill*. *Emotions* [31] is a dataset where each instance is a piece of music. Each piece of music can be labelled with six emotions: sad-lonely, angry-aggressive, amazed-surprised, relaxing-calm, quiet-still, and happy-pleased. *Scene* [32] is a widely used scene classification dataset. Each scene can be annotated in the following six contexts: beach, sunset, field, fall-foliage, mountain, and urban. The *Corel5k* [34] data set contains Corel images that are segmented using normalized cuts. The segmented regions are then clustered into 499 bins, which are further used to describe the images. Each image can be then assigned several of the 374 possible labels. *Mediamill* [36] originates from the 2005 NIST TRECVID challenge dataset,² which contains data about annotated videos. The label space is represented by 101 "annotation concepts", such as explosion, aircraft, face, truck, urban, etc.

The domain of text categorization is represented with 6 datasets: *medical*, *enron*, *tmc2007*, *bibtex*, *delicious* and *bookmarks*. *Medical* [16] is a dataset used in the Medical Natural Language Processing Challenge³ in 2007. Each

²<http://www.science.uva.nl/research/mediamill/challenge/>

³<http://www.computationalmedicine.org/challenge/>

instance is a document that contains brief free-text summary of a patient symptom history. The goal is to annotate each document with the probable diseases from the International Classification of Diseases (ICD-9-CM) [38]. *Enron* [33] is a dataset that contains the e-mails from 150 senior Enron officials. The e-mails were categorized into several categories developed by the UC Berkeley Enron Email Analysis Project⁴. The labels can be further grouped into four categories: coarse genre, included/forwarded information, primary topics, and messages with emotional tone. *Tmc2007* [35] contains instances of aviation safety reports that document problems that occurred during certain flights. The labels represent the problems being described by these reports. We use a reduced version of this dataset with the top 500 attributes selected, same as Tsoumakas et al. [15]. *Delicious*, *bibtex* and *bookmarks* are used for automatic tag suggestion. *Delicious* [24] contains web pages and their tags. The web pages are taken from the del.icio.us social bookmarking site⁵. Note that the label space is greater than the size of the input space ($Q > D$) for this dataset. *Bibtex* [37] contains metadata for bibtex items, such as the title of the paper, the authors, book title, journal volume, publisher, etc., while *bookmarks* [37] contains metadata for bookmark items, such as the URL of the web page, an URL hash, a description of the web page, etc.

4.2. Evaluation measures

Performance evaluation for multi-label learning systems differs from that of classical single-label learning systems. In any multi-label experiment, it is essential to include multiple and contrasting measures because of the additional degrees of freedom that the multi-label setting introduces. In our experiments, we used various evaluation measures that have been suggested by Tsoumakas et al. [30]. Figure 3 depicts a categorization of the used evaluation measures. Furthermore, we evaluate the algorithms by their efficiency. Namely, we measure the time needed to construct the predictive models (*training time*) and the time needed to obtain a prediction for an unseen example (*testing time*).

The evaluation measures of predictive performance are divided into two groups: *bipartitions-based* and *rankings-based*. The bipartitions-based evaluation measures are calculated based on the comparison of the predicted relevant labels with the ground truth relevant labels. This group of evaluation measures is further divided into *example-based* and *label-based*. The example-based evaluation measures are based on the average differences of the actual and the predicted sets of labels over all examples of the evaluation dataset. The label-based evaluation measures, on the other hand, assess the predictive performance for each label separately and then average the performance over all labels. In our experiments, we used six example-based evaluation measures (*Hamming loss*, *accuracy*, *precision*, *recall*, *F₁ score* and *subset accuracy*) and six label-based evaluation measures (*micro precision*, *micro recall*, *micro F₁*, *macro precision*, *macro recall* and *macro F₁*). Note that these evaluation measures require predictions stating that a given label is present or not (binary 1/0 predictions). However, most predictive models predict a numerical value for each label and the label is predicted as present if that numerical value exceeds some pre-defined threshold τ . The performance of the predictive model thus directly depends on the selection of an appropriate value of τ . To this end, we applied a threshold calibration method by choosing the threshold that minimizes the difference in label cardinality between the training data and the predictions for the test data [16].

The ranking-based evaluation measures compare the predicted ranking of the labels with the ground truth ranking. We used four ranking-based measures: *one-error*, *coverage*, *ranking loss* and *average precision*. A detailed description of the evaluation measures is given in Appendix A.

4.3. Experimental setup

The comparison of the multi-label learning methods was performed using the implementations in the following machine learning systems: MULAN⁶ library under the machine learning framework WEKA [39], MEKA⁷ extension for the WEKA framework and CLUS⁸ system for predictive clustering. The MULAN library was used for BR, CLR, QWML, HOMER, ML-C4.5, RFML-C4.5, ML-kNN and RAKEL; the MEKA environment was used for CC and ECC and the CLUS system for PCT and RF-PCT. All experiments were performed on a server with an Intel Xeon processor at 2.50GHz on 64GB of RAM with the Fedora 14 operating system. In the remainder of this section, we first state the base classifiers that were used for the multi-label methods and then the parameter instantiations of the methods.

⁴http://bailando.sims.berkeley.edu/enron_email.html

⁵<http://del.icio.us>

⁶<http://mulan.sourceforge.net/>

⁷<http://meka.sourceforge.net/>

⁸<http://clus.sourceforge.net>

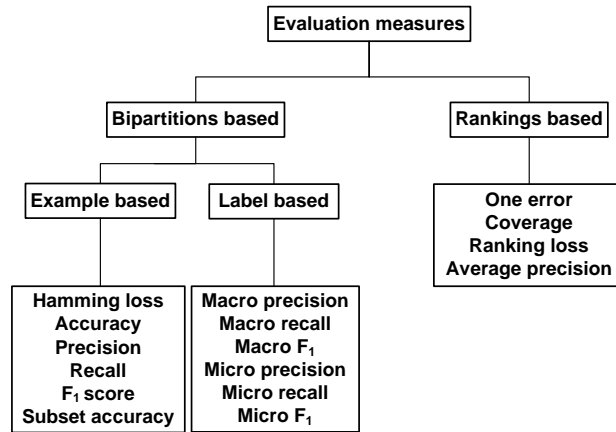


Figure 3: Categorization of evaluation measures used to assess the predictive performance of methods for multi-label learning.

4.3.1. Base classifiers

The methods used in this study use two types of base classifiers for solving the partial binary classification problems in all problem transformation methods and the ensemble methods: SVMs and decision trees (see Figure 2). For training the SVMs, we used the implementation from the LIBSVM library [40]. In particular, we used SVMs with a radial basis kernel for all problem transformation methods and RAKEL and ECC. The kernel parameter γ and the penalty C , for each combination of dataset and method, were determined by 10-fold cross validation using only the training sets. The exception to this is the ensemble method RAKEL where the kernel parameter γ and the penalty C were determined by 5-fold cross validation for the *tmc2007* and *mediamill* datasets because of its computational complexity. The values $2^{-15}, 2^{-13}, \dots, 2^1, 2^3$ were considered for γ and $2^{-5}, 2^{-3}, \dots, 2^{13}, 2^{15}$ for the penalty C . After determining the best parameters values for each method on every dataset, the classifiers were trained using all available training examples and were evaluated by recognizing all test examples from the corresponding dataset.

We used two implementations of decision trees: ML-C4.5 from MULAN and PCTs from CLUS. The ML-C4.5 and PCT as predictive models were pruned using a pruning method. ML-C4.5 uses a post pruning strategy based on a confidence factor, while PCTs use a pre-pruning strategy based on the F-test (whether a given split significantly reduces the variance). On the other hand, when they were used as base classifiers in the ensembles (RFML-C4.5 and RF-PCT), the trees were fully grown [41].

4.3.2. Parameter instantiation

The parameters of the methods were instantiated following the recommendations from the literature. In particular, for the ensemble methods based on decision trees (RFML-C4.5 and RF-PCT), the number of models (classifiers) used in the ensemble was 100 as suggested by Bauer and Kohavi [41]. For the size of the feature subsets needed for construction of the base classifiers for RFML-C4.5, we selected the $f(x) = \lfloor \log_2(x) + 1 \rfloor$ as recommended by Breiman [29], while for RF-PCT we selected $f(x) = \lfloor 0.1 \cdot x + 1 \rfloor$ as recommended by Kocev [18]. The number of models in the ECC method was set to 10 as proposed by Read et al. [16]. Next, the number of models in RAKEL was set to $\min(2 \cdot Q, 100)$ (Q is the number of labels) for all datasets [15], except for the *mediamill*, *delicious* and *bookmarks* datasets, where this parameter was set to 10 as a result of the memory requirements of this method. Besides the number of base classifiers, RAKEL requires one additional parameter: the size of the label-sets k . For each dataset, this parameter was set to half the number of labels ($Q/2$). Tsoumakas et al. [15] and Read et al. [16] have shown that this is a reasonable choice, since it provides a balance between computational complexity and predictive performance.

The ML-C4.5 method uses sub-tree raising as a post-pruning strategy with a pruning confidence set to 0.25. Furthermore, the minimal number of examples in the leaves in each model of the RFML-C4.5 was set to 10. PCTs use a pre-pruning strategy that employs the F-test to determine whether a given split results in a significant reduction of variance. The significance level for the F-test was automatically selected from a predefined list of significance levels using 3-fold cross-validation. The number of neighbors in the ML-kNN method for each dataset was determined

from the values 6 to 20 with step 2. HOMER also requires one additional parameter to be configured: the number of clusters. For this parameter, five different values (2-6) were considered in the experiments [24] and we report the best results.

4.4. Statistical evaluation

To assess whether the overall differences in performance across the ten different approaches are statistically significant, we employed the corrected Friedman test [42] and the post-hoc Nemenyi test [43] as recommended by Demšar [44]. The Friedman test is a non-parametric test for multiple hypotheses testing. It ranks the algorithms according to their performance for each dataset separately, thus the best performing algorithm gets the rank of 1, the second best the rank of 2, etc. In case of ties, it assigns average ranks. Then, the Friedman test compares the average ranks of the algorithms and calculates the Friedman statistic χ_F^2 , distributed according to the χ_F^2 distribution with $k - 1$ degrees of freedom (k being the number of algorithms). Iman and Davenport [45] have shown that the Friedman statistic is undesirably conservative and derive a corrected F-statistic that is distributed according to the F-distribution with $k - 1$ and $(k - 1) \cdot (N - 1)$ degrees of freedom (N being the number of datasets).

If a statistically significant difference in the performance is detected, then next step is a post-hoc test to detect between which algorithms those differences appear. The Nemenyi test is used to compare all the classifiers to each other. In this procedure, the performance of two classifiers is significantly different if their average ranks differ by more than some critical distance. The critical distance depends on the number of algorithms, the number of datasets and the critical value (for a given significance level - p) that is based on the Studentized range statistic and can be found in statistical textbooks (e.g., see [46]).

We present the results from the Nemenyi post-hoc test with average rank diagrams [44]. These are given in Figures 4 to 7, and Figure B.8 to B.11. A critical diagram contains an enumerated axis on which the average ranks of the algorithms are drawn. The algorithms are depicted along the axis in such a manner, that the best ranking ones are at the right-most side of the diagram. The lines for the average ranks of the algorithms that do not differ significantly (at the significance level of $p = 0.05$) are connected with a line.

For the larger datasets, several algorithms did not construct a predictive model within one week under the available resources⁹. These occurrences are marked as DNF (Did Not Finish) in the tables with the results. Considering this, we perform the statistical analysis twice. For the first analysis, we use only the datasets for which all the methods finished and provided results (eight datasets). For the second analysis, we penalize the algorithms that do not finish by assigning them the lowest value (i.e., the lowest rank value for the given algorithm-dataset pair) for each evaluation measure.

5. Results and discussion

In this section, we present the results from the experimental evaluation. For each type of evaluation measure, we present and discuss the critical diagrams from the tests for statistical significance using the datasets on which all algorithms provided predictive models. We give complete results over all evaluation measures and all critical diagrams (including those for all datasets) in Appendix B.

5.1. Results on the example-based measures

The example-based evaluation measures include Hamming loss, Accuracy, Precision, Recall, F_1 score and Subset accuracy. The results of the statistical evaluation are given in Figure 4, while the complete results are given in Tables B.2 through B.8, and Figure B.8. Considering the results that include the datasets for which all algorithms finished (Figure 4), we can make several conclusions. The first conclusion that draws our attention is that HOMER performs best as evaluated by *recall*, while RF-PCT performs best according to *precision*. This means that the predictions made by HOMER are more complete: the original relevant labels were correctly predicted as relevant labels (small number of false negatives results in high *recall*). However, the lower *precision* means that besides the labels that were originally relevant, HOMER predicts non-relevant labels as relevant (larger number of false positives results

⁹The experiments were performed on a server running Linux, with two Intel Quad-Core Processors running at 2.5GHz and 64GB of RAM.

in low *precision*). The situation is somewhat reversed when looking at the predictions from RF-PCT. The predictions of RF-PCT are more exact: the labels predicted as relevant were truly relevant in the original examples (small number of false positives results in high *precision*). However, RF-PCT is leaving out some of the relevant labels when making predictions (larger number of false negatives results in high *recall*).

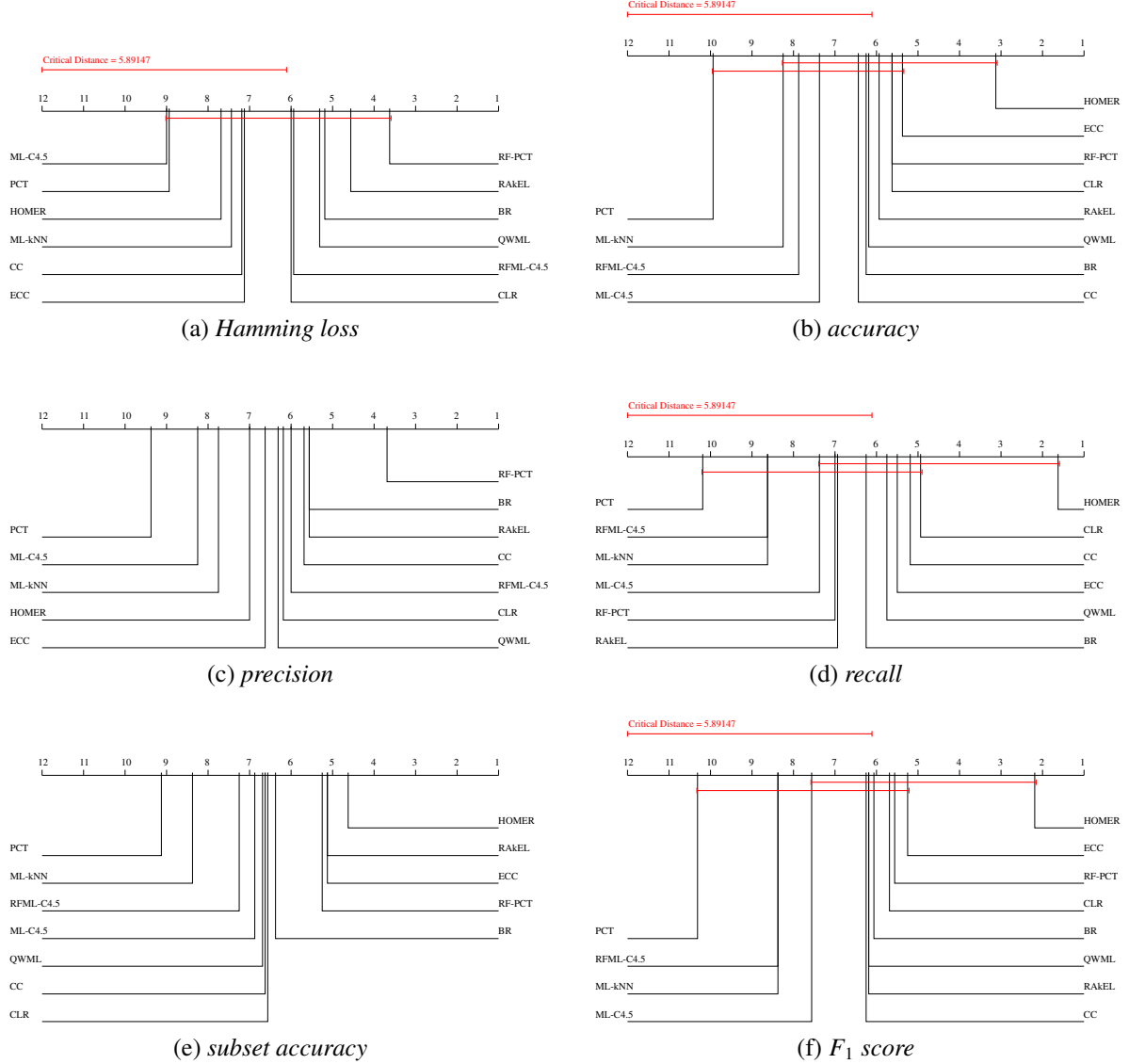


Figure 4: The critical diagrams for the example-based evaluation measures: The results from the Nemenyi post-hoc test at 0.05 significance level on the datasets for which all algorithms provided results. For *precision* and *subset accuracy* the differences are not statistically significant according to the Friedman test (see Table B.8), thus we show only the average ranks of the algorithms.

We further analyze the performance of the methods across all six evaluation measures: the best performing methods on all measures are either RF-PCT or HOMER. We can further note that RF-PCT is the best performing method, closely followed by HOMER, BR and CC. The RF-PCT method performs best according to *subset accuracy*, *Hamming loss* and *precision*, second best according to *accuracy*. HOMER is the best performing as evaluated by *accuracy*, *recall* and *F₁ score*, and second best according to *subset accuracy*. The HOMER method has poor performance as evaluated by *Hamming loss* and *precision* (10-th and 9-th position on the critical diagrams, respectively), while on

these two measures RF-PCT performs the best. We hypothesize that the low performance of HOMER according to *Hamming loss* is because the procedure for construction of HOMER's hierarchical structure does not optimize *Hamming loss*.

The differences in predictive performance are rarely significant at the significance level of 0.05. HOMER and RF-PCT are often significantly better than single PCT, or single ML-C4.5 trees. From an ensemble learning point of view, this means that the RF-PCTs lift the predictive performance of a single PCT even when the target concept is a set of labels, similarly as for simple regression and classification. On the other hand, the increase in predictive performance is not constraint for the ECC and RF-MLC4.5 ensembles: according to some evaluation measure, the single models perform even better on average than the corresponding ensembles. For this we have two hypotheses: first, CC are stable classifiers and ensemble can't much improve over their predictive performance. Second, RF-MLC4.5 did not perform competitively because it selects feature subsets with a logarithmic size compared to the complete set of features. Considering that the domains we used in this study (and other multi-label domains) have a large number of features (typically larger than 500), the logarithmic function under-samples the feature space and is missing some useful information that can contribute to better classification.

We next focus the discussion on the different types of base machine learning algorithms. First, we can note that the multi-label variant of k-nearest neighbors (ML-kNN) performs poor by across all evaluation measures. Next, the SVM-based methods perform better for the smaller datasets, while tree-based methods for the larger datasets. This is because the Gaussian kernel can handle very well the smaller number of examples: when the number of examples increases, the performance of the kernel approaches the performance of a linear kernel. Furthermore, the SVM-based methods are better for the domains with larger numbers of features. For instance, in text classification an example is a document typically represented as a bag-of-words, where each feature can play a crucial role in making a correct prediction. The SVMs exploit the information from all the features, while the decision trees use only a (small) subset of features and may miss some crucial information.

Finally, we discuss the addition of all datasets in the statistical analysis (shown in Figure B.8). This analysis shows that RF-PCT and BR have improved predictive performance at the expense of the methods that did not finish. RF-PCT is again the best performing method overall, followed by BR, HOMER and CC. As for the subset of datasets, RF-PCT is best according to *Hamming loss*, *precision* and *subset accuracy*, while HOMER is best according to *accuracy*, F_1 *score* and *recall*. RF-PCT is second best on *accuracy* and F_1 *score* and has an improved performance according to *recall*. Furthermore, RF-PCT is statistically significantly better than HOMER according to *Hamming loss*.

5.2. Results on the label-based measures

The label-based evaluation measures include *micro precision*, *micro recall*, *micro F_1* , *macro precision*, *macro recall* and *macro F_1* . The results from the statistical evaluation are given in Figure 5, while complete results are given in Tables B.9 through B.15, and Figure B.9. First, we focus on the results and the statistical analysis on the datasets for which all methods have finished. As for the example-based measures, the best performing methods are RF-PCT, HOMER, BR and CC. HOMER performs best according to four evaluation measures: *macro F_1* , *macro recall*, *micro F_1* and *micro recall* and it performs worst of all methods according to *micro precision* and 8-th according to *macro precision*. RF-PCT is the best performing using *micro precision* and *macro precision* and, according to the critical diagrams, performs statistically significantly better than HOMER.

We next discuss the performance of the ensembles and the single models. Again, as for the example-based measures, RF-PCT is better than single PCT over all evaluation measures. On the other hand, this is not the case for RF-MLC4.5 and ECC. The reasons for this are the same as for the example-based measures: CC is a stable classifier and the logarithmic size of the feature subset for RF-MLC4.5 is under-sampling the feature space.

The behavior of the base machine learning algorithms remains the same as for the example-based measures. ML-kNN again has very poor predictive performance across all evaluation measures. SVMs are better for the smaller datasets and decision trees for the larger datasets.

Finally, the addition of the datasets for which some (but not all) of the methods finished did not change the results much (compare Figure 5 with Figure B.9). The updated results improved the average performance of RF-PCT, BR and CC at the expense of the methods not able to produce results. However, the relative average performance remained the same as for the subset of datasets: HOMER is best according to *macro F_1* , *macro recall*, *micro F_1* and *micro recall*, while RF-PCT is best according to *micro precision* and *macro precision*. Moreover, RF-PCT is statistically significantly better than HOMER according to *micro precision* and *macro precision*.

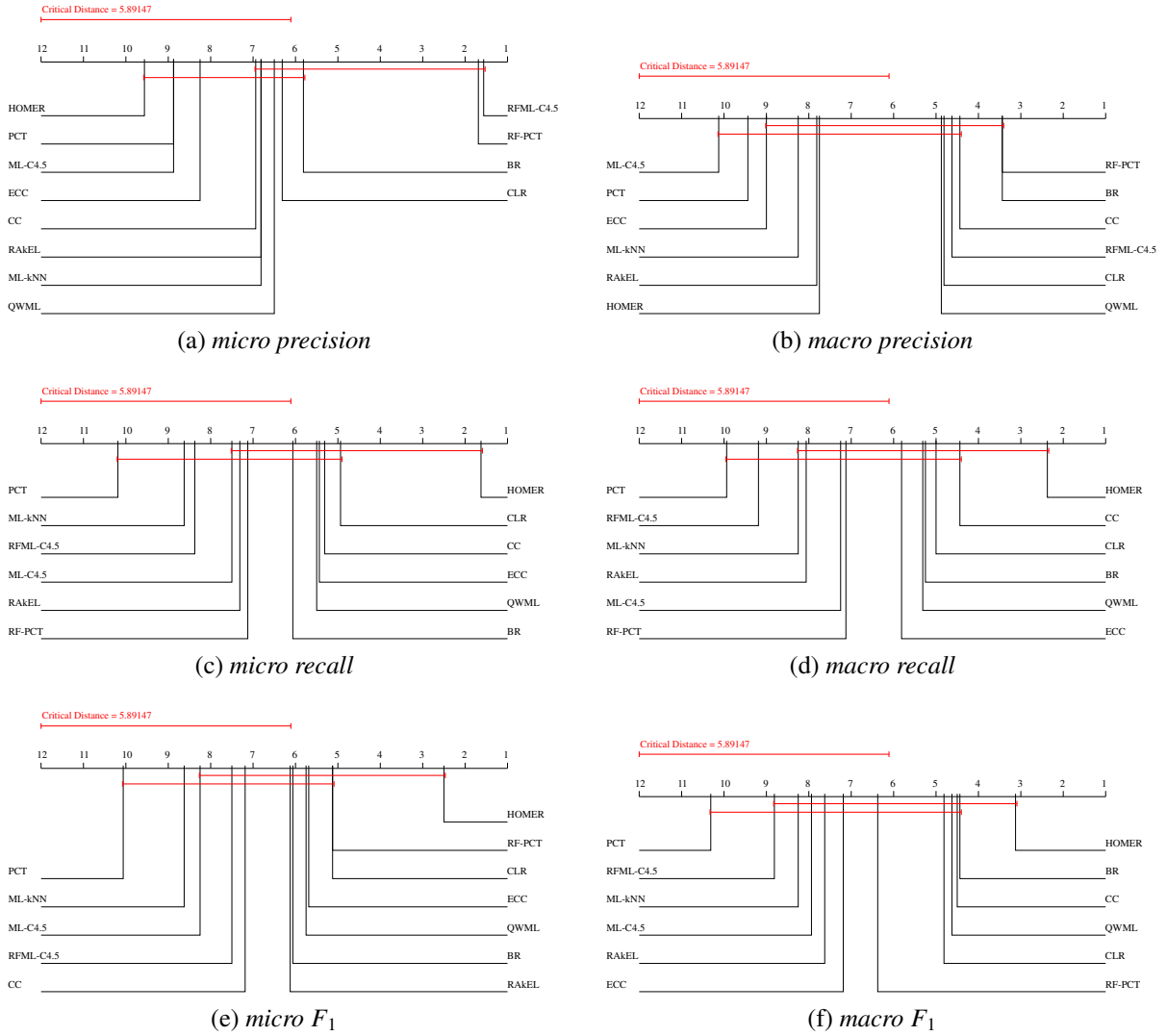


Figure 5: The critical diagrams for the label-based evaluation measures: The results from the Nemenyi post-hoc test at 0.05 significance level on the datasets for which all algorithms provided results.

5.3. Results on the ranking-based measures

The ranking-based measures include *one-error*, *ranking loss*, *coverage* and *average precision*. The results from the statistical evaluation are given in Figure 6, while the complete results are given in Tables B.16 through B.20, and Figure B.10. We first focus on the results for the datasets for which all methods have finished. The best performing method is RF-PCT, followed by BR, CC and CLR. RF-PCT is the best performing method on all four evaluation measures, BR is second best on three measures (*one-error*, *ranking loss* and *average precision*) and third on *coverage*. Considering ensemble learning, the ensembles based on decision trees perform better than the corresponding single models. However, the ensembles of CC perform worse than a single CC. The ranking-based measures indicate that the SVM-based methods perform better for smaller datasets, while tree-based measures perform better on larger datasets.

Let us further compare RF-PCT with HOMER using the ranking measures. The statistical evaluation of the performance reveals that RF-PCT is statistically significantly better than HOMER at the significance level of 0.05 according to *coverage* and *ranking loss* (see Figure 6). Furthermore, the statistical evaluation using all datasets (Figure B.10) shows that RF-PCT is statistically significantly better than HOMER on all four evaluation measures.

The other results from the statistical analysis using all datasets are similar to the results on the subset of datasets.

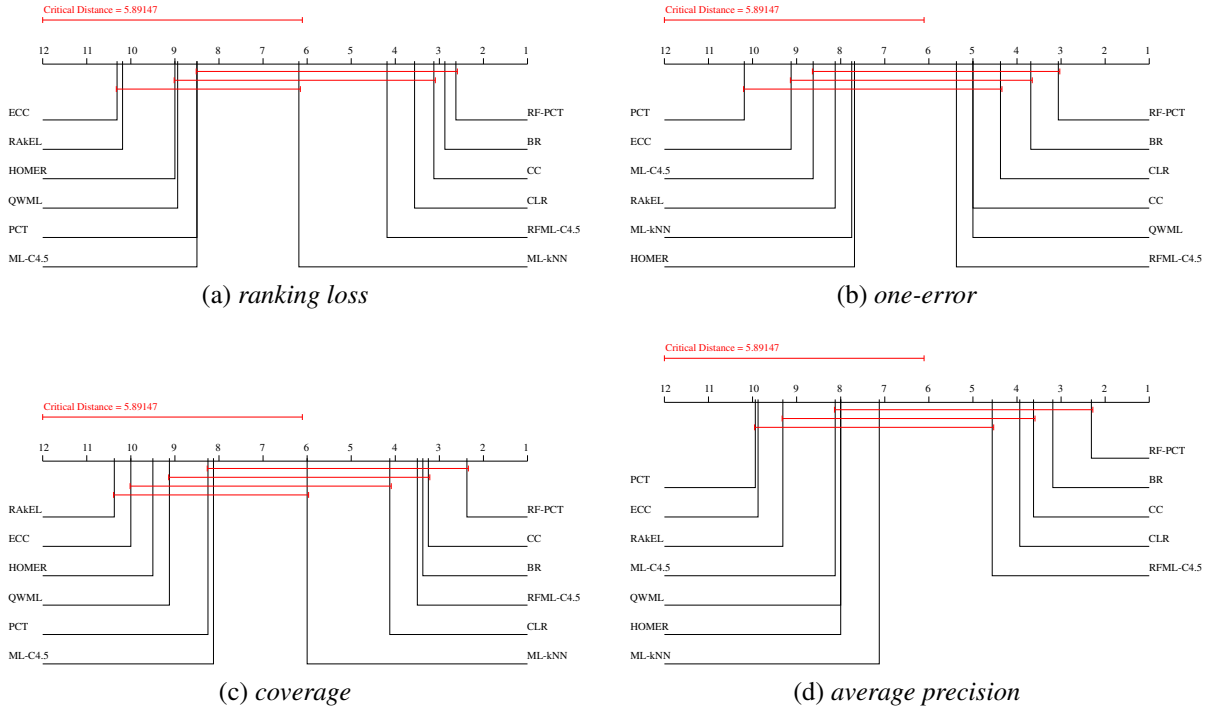


Figure 6: The critical diagrams for the ranking-based evaluation measures: The results from the Nemenyi post-hoc test at 0.05 significance level on the datasets for which all algorithms provided results.

5.4. Results on the efficiency measures

We finally discuss the efficiency of the proposed methods in terms of training and testing time. The results are given in Figures 7 and B.11, and Tables B.21 through B.23. They show that the tree-based methods are more efficient than the SVM-based methods. Namely, PCT is the most efficient method, followed by ML-C4.5 and ML-kNN. PCTs are faster to construct than ML-C4.5 because of the pruning strategy they employ: the former used pre-pruning and the latter post-pruning.

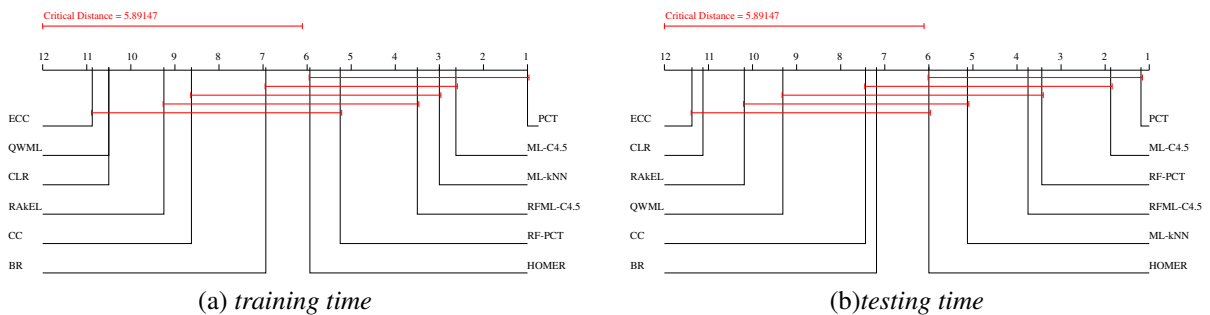


Figure 7: The critical diagrams for the efficiency measures: The results from the Nemenyi post-hoc test at 0.05 significance level on the datasets for which all algorithms provided results.

We further discuss the methods that exhibited the best predictive performance according to the other evaluation measures: RF-PCT and HOMER. RF-PCT is better than HOMER on the time needed to produce a prediction for an

unseen example (*testing time*) and on the time needed for learning a classifier (*training time*) for both analyses: the first includes only the datasets with complete results and the second includes all datasets. Moreover, HOMER did not produce results for the *bookmarks* dataset.

6. Conclusions

In this study, we present an extensive experimental evaluation of methods for multi-label learning. The topic of multi-label learning has lately received significant research effort. It has also attracted much attention from the research community, in the form of journal special issues and workshops at major conferences. This has resulted in a variety of methods for addressing the task of multi-label learning. However, a wider experimental comparison of these methods is still lacking in the literature.

We evaluate the most popular methods for multi-label learning using a wide range of evaluation measures on a variety of datasets. Below we explain the dimensions of the extensive experimental evaluation. First, we selected 12 multi-label methods that were recently proposed in the literature. The selected methods are divided in three main groups: algorithm adaptation (3 methods), problem transformation (5 methods) and ensembles (4 methods). The methods use three types of basic machine learning algorithms: SVMs (7 methods), decision trees (4 methods) and k-nearest neighbors (1 method). Second, we used 16 different evaluation measures that are typically used in the context of multi-label learning. The variety of evaluation measures is necessary to provide a view on algorithm performance from different perspectives. The evaluation measures are divided in three groups: example-based (6 measures), label-based (6 measures) and ranking-based (4 measures). Furthermore, we assess the efficiency of the methods by measuring the time needed to learn the classifier and the time needed to produce a prediction for an unseen example. Third, we evaluate the methods on 11 multi-label benchmark datasets from 5 application domains: text classification (6 datasets), image classification (2 datasets), gene function prediction (1 dataset), music classification (1 dataset) and video classification (1 dataset). We then analyze the results from the experiments using Friedman and Nemenyi tests for assessing the statistical significance of the differences in performance. We present the results from the statistical tests using critical diagrams.

The results of the experimental comparison revealed that the best performing methods are RF-PCT and HOMER, followed by BR and CC. For each performance measure, the best algorithm was either RF-PCT or HOMER. The example-based measures, which are most widely used for multi-label classification, show that RF-PCT is best according to precision and has average performance on recall. On the other hand, HOMER is best according to recall, while having poor performance on precision. This means that the predictions from RF-PCT are more exact than the ones from HOMER, while the predictions from HOMER are more complete than the ones from RF-PCT. Considering the basic machine learning algorithms underlying the compared approaches, the SVM based methods are better on datasets with a large number of features and a smaller number of examples, since they can exploit the information from all of the features, while the decision trees exploit only a subset of the features.

The label-based measures showed behavior similar to that of the example-based measures. However, the gap between HOMER and RF-PCT on recall and precision is now much bigger. Namely, RF-PCT is statistically significantly better than HOMER on the two precision-based measures.

The ranking-based measures offer a different perspective on the results. RF-PCT was the best performing method, followed by CC and BR. On these measures HOMER exhibited poor performance. RF-PCT is statistically significantly better than HOMER on two evaluation measures (coverage and ranking loss) using the datasets for which there are results from all methods and statistically significantly better on all four evaluation measures using all datasets. Furthermore, these measures emphasize the advantages of the SVM-based methods on the smaller datasets and the tree-based methods on the larger datasets.

Considering efficiency, the tree-based methods are generally faster to train a classifier and produce a prediction for an unseen example than the SVM-based methods. We further compare the efficiency of the RF-PCT and HOMER methods. The results show that RF-PCT is faster than HOMER on testing time (on average 194.6 times) and on training time (on average 8.4 times).

The experimental comparison can be extended by including more methods for multi-label learning. For example, one can also include bagging of PCTs: ensemble method that has competitive performance to random forests of PCTs [18]. The comparison can be also extended by including other evaluation measures. One evaluation measure that can

be easily adapted for multi-label setting is the precision-recall curve (and the area under the precision-recall curve thereof) [47]. This will offer a better insight to the trade-off between the precision and recall performance of a given method for multi-label learning.

All in all, the final recommendation considering the performance and the efficiency of the evaluated methods is that RF-PCT, HOMER, BR and CC should be used as benchmark methods for multi-label learning. Over all evaluation measures these methods performed best. Furthermore, RF-PCT and HOMER exhibited the best predictive performance and better efficiency than the rest of the methods.

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Appendix A. Evaluation measures

In this section, we present the measures that are used to evaluate the predictive performance of the compared methods in our experiments. In the definitions below, \mathcal{Y}_i denotes the set of true labels of example \mathbf{x}_i and $h(\mathbf{x}_i)$ denotes the set of predicted labels for the same examples. All definitions refer to the multi-label setting.

Appendix A.1. Example based measures

Hamming loss evaluates how many times an example-label pair is misclassified, i.e., label not belonging to the example is predicted or a label belonging to the example is not predicted. The smaller the value of $hamming_loss(h)$, the better the performance. The performance is perfect when $hamming_loss(h) = 0$. This metric is defined as:

$$hamming_loss(h) = \frac{1}{N} \sum_{i=1}^N \frac{1}{Q} |h(\mathbf{x}_i) \Delta \mathcal{Y}_i| \quad (\text{A.1})$$

where Δ stands for the symmetric difference between two sets, N is the number of examples and Q is the total number of possible class labels.

Accuracy for a single example \mathbf{x}_i is defined by the Jaccard similarity coefficients between the label sets $h(\mathbf{x}_i)$ and \mathcal{Y}_i . Accuracy is micro-averaged across all examples.

$$accuracy(h) = \frac{1}{N} \sum_{i=1}^N \frac{|h(\mathbf{x}_i) \cap \mathcal{Y}_i|}{|h(\mathbf{x}_i) \cup \mathcal{Y}_i|} \quad (\text{A.2})$$

Precision is defined as:

$$precision(h) = \frac{1}{N} \sum_{i=1}^N \frac{|h(\mathbf{x}_i) \cap \mathcal{Y}_i|}{|h(\mathbf{x}_i)|} \quad (\text{A.3})$$

Recall is defined as:

$$recall(h) = \frac{1}{N} \sum_{i=1}^N \frac{|h(\mathbf{x}_i) \cap \mathcal{Y}_i|}{|h(\mathbf{x}_i)|} \quad (\text{A.4})$$

F_1 **score** is the harmonic mean between precision and recall and is defined as:

$$F_1 = \frac{1}{N} \sum_{i=1}^N \frac{2 \times |h(\mathbf{x}_i) \cap \mathcal{Y}_i|}{|h(\mathbf{x}_i)| + |\mathcal{Y}_i|} \quad (\text{A.5})$$

F_1 is an example based metric and its value is an average over all examples in the dataset. F_1 reaches its best value at 1 and worst score at 0.

Subset accuracy or classification accuracy is defined as follows:

$$subset_accuracy(h) = \frac{1}{N} \sum_{i=1}^N I(h(\mathbf{x}_i) = \mathcal{Y}_i) \quad (\text{A.6})$$

where $I(true) = 1$ and $I(false) = 0$. This is a very strict evaluation measure as it requires the predicted set of labels to be an exact match of the true set of labels.

Appendix A.2. Label based measures

Macro precision (precision averaged across all labels) is defined as:

$$macro_precision = \frac{1}{Q} \sum_{j=1}^Q \frac{tp_j}{tp_j + fp_j} \quad (\text{A.7})$$

where tp_j , fp_j are the number of true positives and false positives for the label λ_j considered as a binary class.

Macro recall (recall averaged across all labels) is defined as:

$$macro_recall = \frac{1}{Q} \sum_{j=1}^Q \frac{tp_j}{tp_j + fn_j} \quad (\text{A.8})$$

where tp_j , fp_j are defined as for the macro precision and fn_j is the number of false negatives for the label λ_j considered as a binary class.

Macro F_1 is the harmonic mean between precision and recall, where the average is calculated per label and then averaged across all labels. If p_j and r_j are the precision and recall for all $\lambda_j \in h(\mathbf{x}_i)$ from $\lambda_j \in \mathcal{Y}_i$, the macro F_1 is

$$macro_F_1 = \frac{1}{Q} \sum_{j=1}^Q \frac{2 \times p_j \times r_j}{p_j + r_j} \quad (\text{A.9})$$

Micro precision (precision averaged over all the example/label pairs) is defined as:

$$micro_precision = \frac{\sum_{j=1}^Q tp_j}{\sum_{j=1}^Q tp_j + \sum_{j=1}^Q fp_j} \quad (\text{A.10})$$

where tp_j , fp_j are defined as for macro precision.

Micro recall (recall averaged over all the example/label pairs) is defined as:

$$micro_recall = \frac{\sum_{j=1}^Q tp_j}{\sum_{j=1}^Q tp_j + \sum_{j=1}^Q fn_j} \quad (\text{A.11})$$

where tp_j and fn_j are defined as for macro recall.

Micro F_1 is the harmonic mean between micro precision and micro recall. Micro F_1 is defined as:

$$micro_F_1 = \frac{2 \times micro_precision \times micro_recall}{micro_precision + micro_recall} \quad (\text{A.12})$$

Appendix A.3. Ranking based measures

One error evaluates how many times the top-ranked label is not in the set of relevant labels of the example. The metric $one_error(f)$ takes values between 0 and 1. The smaller the value of $one_error(f)$, the better the performance. This evaluation metric is defined as:

$$one_error(f) = \frac{1}{N} \sum_{i=1}^N \left[\left[\arg \max_{\lambda \in \mathcal{Y}} f(\mathbf{x}_i, \lambda) \right] \notin \mathcal{Y}_i \right] \quad (\text{A.13})$$

where $\lambda \in \mathcal{L} = \{\lambda_1, \lambda_2, \dots, \lambda_Q\}$ and $\llbracket \pi \rrbracket$ equals 1 if π holds and 0 otherwise for any predicate π . Note that, for single-label classification problems, the One Error is identical to ordinary classification error.

Coverage evaluates how far, on average, we need to go down the list of ranked labels in order to cover all the relevant labels of the example. The smaller the value of $coverage(f)$, the better the performance.

$$coverage(f) = \frac{1}{N} \sum_{i=1}^N \max_{\lambda \in \mathcal{Y}_i} rank_f(\mathbf{x}_i, \lambda) - 1 \quad (\text{A.14})$$

where $rank_f(\mathbf{x}_i, \lambda)$ maps the outputs of $f(\mathbf{x}_i, \lambda)$ for any $\lambda \in \mathcal{L}$ to $\{\lambda_1, \lambda_2, \dots, \lambda_Q\}$ so that $f(\mathbf{x}_i, \lambda_m) > f(\mathbf{x}_i, \lambda_n)$ implies $rank_f(\mathbf{x}_i, \lambda_m) < rank_f(\mathbf{x}_i, \lambda_n)$. The smallest possible value for $coverage(f)$ is l_c , i.e., the label cardinality of the given dataset.

Ranking loss evaluates the average fraction of label pairs that are reversely ordered for the particular example given by:

$$ranking_loss(f) = \frac{1}{N} \sum_{i=1}^N \frac{|D_i|}{|\mathcal{Y}_i| |\bar{\mathcal{Y}}_i|} \quad (\text{A.15})$$

where $D_i = \{(\lambda_m, \lambda_n) | f(\mathbf{x}_i, \lambda_m) \leq f(\mathbf{x}_i, \lambda_n), (\lambda_m, \lambda_n) \in \mathcal{Y}_i \times \bar{\mathcal{Y}}_i\}$, while $\bar{\mathcal{Y}}$ denotes the complementary set of \mathcal{Y} in \mathcal{L} . The smaller the value of $ranking_loss(f)$, the better the performance, so the performance is perfect when $ranking_loss(f) = 0$.

Average Precision is the average fraction of labels ranked above an actual label $\lambda \in \mathcal{Y}_i$ that actually are in \mathcal{Y}_i . The performance is perfect when $avg_precision(f) = 1$; the larger the value of $avg_precision(f)$, the better the performance. This metric is defined as:

$$avg_precision(f) = \frac{1}{N} \sum_{i=1}^N \frac{1}{|\mathcal{Y}_i|} \sum_{\lambda \in \mathcal{Y}_i} \frac{|\mathcal{L}_i|}{rank_f(\mathbf{x}_i, \lambda)} \quad (\text{A.16})$$

where $\mathcal{L}_i = \{\lambda' | rank_f(\mathbf{x}_i, \lambda') \leq rank_f(\mathbf{x}_i, \lambda), \lambda' \in \mathcal{Y}_i\}$ and $rank_f(\mathbf{x}_i, \lambda)$ is defined as in coverage above.

Appendix B. Complete results from the experimental evaluation

In this section, we present the complete results from the experimental evaluation. We present the results based on the evaluation measures. We first present the results for the example-based evaluation measures. We then show the results for label-based evaluation measures. We next give the results for ranking-based evaluation measure. Finally, we present the efficiency of the methods by their training and testing times.

Appendix B.1. Results on the example-based evaluation measures

Table B.2: The performance of the multi-label learning approaches in terms of the *precision* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.550	0.551	0.538	0.548	0.509	0.606	0.577	0.502	0.564	0.580	0.625	0.644
scene	0.718	0.758	0.714	0.711	0.746	0.592	0.565	0.661	0.768	0.770	0.403	0.565
yeast	0.722	0.727	0.719	0.718	0.663	0.620	0.705	0.732	0.715	0.667	0.738	0.744
medical	0.211	0.217	0.695	0.697	0.762	0.797	0.285	0.575	0.730	0.662	0.284	0.635
enron	0.703	0.464	0.650	0.624	0.616	0.623	0.415	0.587	0.708	0.652	0.690	0.709
corel5k	0.042	0.042	0.329	0.326	0.317	0.005	0.000	0.035	0.000	0.002	0.018	0.030
tmc2007	0.941	0.944	0.937	0.937	0.926	0.146	0.659	0.738	0.928	0.872	0.874	0.977
mediamill	0.731	0.741	0.201	0.203	0.597	0.056	0.694	0.724	0.705	0.690	0.765	0.772
bibtex	0.515	0.508	0.488	0.496	0.472	0.123	0.140	0.254	DNF	0.324	0.159	0.292
delicious	0.443	0.399	DNF	DNF	0.369	0.001	0.001	0.424	DNF	DNF	0.472	0.512
bookmarks	DNF	DNF	DNF	DNF	DNF	0.271	0.133	0.218	DNF	DNF	0.182	0.218

Table B.3: The performance of the multi-label learning approaches in terms of the *recall* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.409	0.397	0.410	0.429	0.775	0.703	0.534	0.377	0.491	0.533	0.545	0.582
scene	0.711	0.726	0.712	0.709	0.744	0.582	0.539	0.655	0.740	0.771	0.388	0.541
yeast	0.591	0.600	0.601	0.600	0.714	0.608	0.490	0.549	0.615	0.673	0.491	0.523
medical	0.735	0.754	0.795	0.801	0.760	0.740	0.227	0.547	0.679	0.642	0.251	0.599
enron	0.497	0.507	0.557	0.453	0.610	0.487	0.229	0.358	0.469	0.560	0.398	0.452
corel5k	0.055	0.056	0.264	0.264	0.250	0.002	0.000	0.014	0.000	0.001	0.005	0.009
tmc2007	0.928	0.934	0.929	0.929	0.943	0.111	0.478	0.664	0.880	0.903	0.677	0.920
mediamill	0.450	0.424	0.101	0.101	0.563	0.052	0.379	0.470	0.353	0.372	0.456	0.476
bibtex	0.373	0.378	0.364	0.366	0.389	0.111	0.046	0.132	DNF	0.187	0.060	0.167
delicious	0.155	0.157	DNF	DNF	0.303	0.001	0.001	0.112	DNF	DNF	0.176	0.160
bookmarks	DNF	DNF	DNF	DNF	DNF	0.244	0.137	0.207	DNF	DNF	0.181	0.208

Table B.4: The performance of the multi-label learning approaches in terms of the *Hamming loss* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAkEL	ECC	RFML-C4.5	RF-PCT
emotions	0.257	0.256	0.257	0.254	0.361	0.247	0.267	0.294	0.282	0.281	0.198	0.189
scene	0.079	0.082	0.080	0.081	0.082	0.141	0.129	0.099	0.077	0.085	0.116	0.094
yeast	0.190	0.193	0.190	0.191	0.207	0.234	0.219	0.198	0.192	0.207	0.205	0.197
medical	0.077	0.077	0.017	0.012	0.012	0.013	0.023	0.017	0.012	0.014	0.022	0.014
enron	0.045	0.064	0.048	0.048	0.051	0.053	0.058	0.051	0.045	0.049	0.047	0.046
corel5k	0.017	0.017	0.012	0.012	0.012	0.010	0.009	0.009	0.009	0.009	0.009	0.009
tmc2007	0.013	0.013	0.014	0.014	0.015	0.093	0.075	0.058	0.021	0.026	0.037	0.011
mediamill	0.032	0.032	0.043	0.043	0.038	0.044	0.034	0.031	0.035	0.035	0.030	0.029
bibtex	0.012	0.012	0.012	0.012	0.014	0.016	0.014	0.014	DNF	0.013	0.014	0.013
delicious	0.018	0.018	DNF	DNF	0.022	0.019	0.019	0.018	DNF	DNF	0.018	0.018
bookmarks	DNF	DNF	DNF	DNF	DNF	0.009	0.009	0.009	DNF	DNF	0.009	0.009

Table B.5: The performance of the multi-label learning approaches in terms of the *accuracy* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAkEL	ECC	RFML-C4.5	RF-PCT
emotions	0.361	0.356	0.361	0.373	0.471	0.536	0.448	0.319	0.419	0.432	0.488	0.519
scene	0.689	0.723	0.686	0.683	0.717	0.569	0.538	0.629	0.734	0.735	0.388	0.541
yeast	0.520	0.527	0.524	0.523	0.559	0.480	0.440	0.492	0.531	0.546	0.453	0.478
medical	0.206	0.211	0.656	0.658	0.713	0.730	0.228	0.528	0.673	0.611	0.250	0.591
enron	0.446	0.334	0.459	0.388	0.478	0.418	0.196	0.319	0.428	0.462	0.374	0.416
corel5k	0.030	0.030	0.195	0.195	0.179	0.002	0.000	0.014	0.000	0.001	0.005	0.009
tmc2007	0.891	0.899	0.889	0.889	0.888	0.110	0.436	0.574	0.852	0.808	0.663	0.914
mediamill	0.403	0.390	0.095	0.095	0.413	0.052	0.354	0.421	0.337	0.349	0.423	0.441
bibtex	0.348	0.352	0.334	0.338	0.330	0.108	0.046	0.129	DNF	0.186	0.060	0.166
delicious	0.136	0.137	DNF	DNF	0.207	0.001	0.001	0.102	DNF	DNF	0.151	0.146
bookmarks	DNF	DNF	DNF	DNF	DNF	0.237	0.133	0.202	DNF	DNF	0.176	0.204

Table B.6: The performance of the multi-label learning approaches in terms of the F_1 score measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAkEL	ECC	RFML-C4.5	RF-PCT
emotions	0.469	0.461	0.465	0.481	0.614	0.651	0.554	0.431	0.525	0.556	0.583	0.611
scene	0.714	0.742	0.713	0.710	0.745	0.587	0.551	0.658	0.754	0.771	0.395	0.553
yeast	0.650	0.657	0.655	0.654	0.687	0.614	0.578	0.628	0.661	0.670	0.589	0.614
medical	0.328	0.337	0.742	0.745	0.761	0.768	0.253	0.560	0.704	0.652	0.267	0.616
enron	0.582	0.484	0.600	0.525	0.613	0.546	0.295	0.445	0.564	0.602	0.505	0.552
corel5k	0.047	0.048	0.293	0.292	0.280	0.003	0.000	0.021	0.000	0.001	0.008	0.014
tmc2007	0.934	0.939	0.933	0.933	0.934	0.126	0.554	0.699	0.904	0.887	0.763	0.948
mediamill	0.557	0.539	0.134	0.135	0.579	0.054	0.490	0.570	0.471	0.483	0.572	0.589
bibtex	0.433	0.434	0.417	0.421	0.426	0.117	0.069	0.174	DNF	0.237	0.087	0.212
delicious	0.230	0.225	DNF	DNF	0.343	0.001	0.001	0.017	DNF	DNF	0.256	0.244
bookmarks	DNF	DNF	DNF	DNF	DNF	0.257	0.135	0.213	DNF	DNF	0.181	0.213

Table B.7: The performance of the multi-label learning approaches in terms of the *subset accuracy* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAkEL	ECC	RFML-C4.5	RF-PCT
emotions	0.129	0.124	0.144	0.149	0.163	0.277	0.223	0.084	0.208	0.168	0.272	0.307
scene	0.639	0.685	0.633	0.630	0.661	0.533	0.509	0.573	0.694	0.665	0.372	0.518
yeast	0.190	0.239	0.195	0.192	0.213	0.158	0.152	0.159	0.201	0.215	0.129	0.152
medical	0.000	0.000	0.486	0.480	0.610	0.646	0.177	0.462	0.607	0.526	0.216	0.538
enron	0.149	0.000	0.117	0.097	0.145	0.140	0.002	0.062	0.136	0.131	0.124	0.131
corel5k	0.000	0.000	0.010	0.012	0.002	0.000	0.000	0.000	0.000	0.001	0.008	0.000
tmc2007	0.772	0.787	0.767	0.768	0.765	0.078	0.215	0.305	0.734	0.608	0.421	0.816
mediamill	0.080	0.080	0.044	0.044	0.053	0.049	0.065	0.110	0.060	0.065	0.104	0.122
bibtex	0.194	0.202	0.183	0.186	0.165	0.095	0.004	0.056	DNF	0.109	0.011	0.098
delicious	0.004	0.006	DNF	DNF	0.001	0.001	0.001	0.003	DNF	DNF	0.018	0.007
bookmarks	DNF	DNF	DNF	DNF	DNF	0.209	0.129	0.187	DNF	DNF	0.167	0.189

Table B.8: The p-values of the assessment of performance of the multi-label learning approaches by the Friedman test using the example-based evaluation measures. *Subset* shows the calculated p-values for the datasets on which all algorithms finished. *All* shows the calculated p-values for all datasets including those which did not finished.

	All	Subset
Hamming loss	0.0895	0.047
Accuracy	0.077	0.037
Precision	0.117	0.19
Recall	$5 \cdot 10^{-4}$	$7.9 \cdot 10^{-5}$
F_1 score	$5 \cdot 10^{-3}$	0.0013
Subset accuracy	0.555	0.343

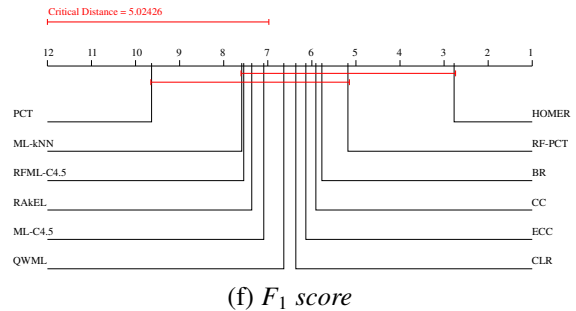
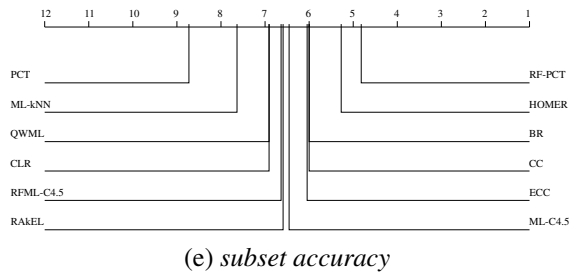
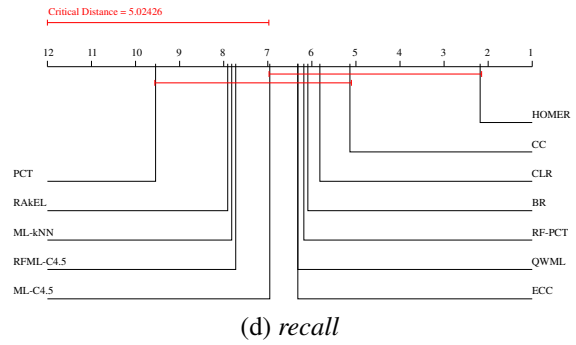
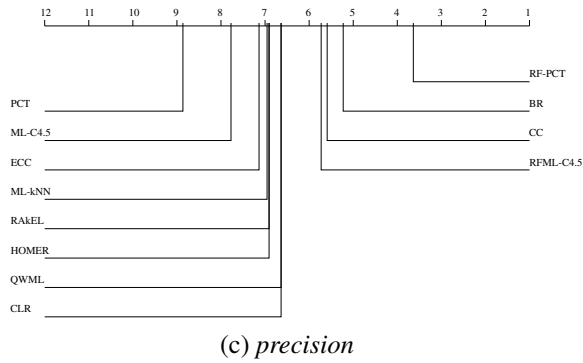
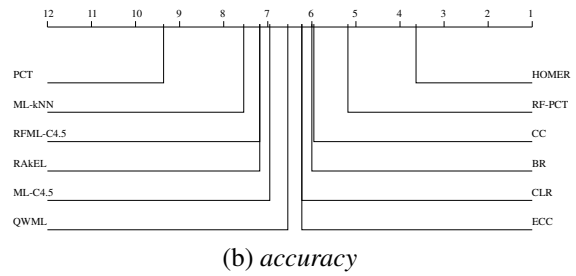
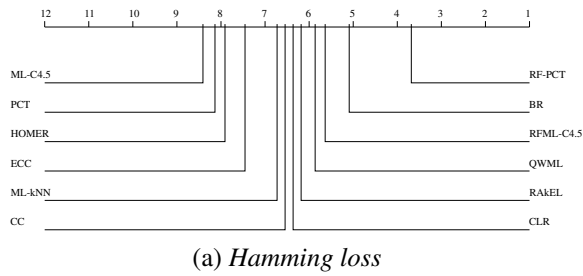


Figure B.8: The critical diagrams for the example-based evaluation measures: The results from the Nemenyi post-hoc test at 0.05 significance level on all the datasets. For *Hamming loss*, *precision*, *accuracy* and *subset accuracy* the differences are not statistically significant according to the Friedman test (see Table B.8), thus we show only the average ranks of the algorithms.

Appendix B.2. Results on the label-based evaluation measures

Table B.9: The performance of the multi-label learning approaches in terms of the *micro precision* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.684	0.698	0.685	0.680	0.471	0.607	0.607	0.584	0.586	0.579	0.783	0.783
scene	0.843	0.814	0.835	0.832	0.804	0.619	0.512	0.691	0.831	0.773	0.960	0.930
yeast	0.733	0.726	0.729	0.727	0.647	0.618	0.698	0.736	0.720	0.662	0.747	0.755
medical	0.225	0.229	0.669	0.667	0.807	0.796	0.826	0.807	0.881	0.834	0.884	0.885
enron	0.721	0.492	0.652	0.687	0.597	0.613	0.601	0.684	0.743	0.642	0.768	0.738
corel5k	0.061	0.061	0.338	0.339	0.308	0.160	0.000	0.730	0.000	0.333	0.750	0.696
tmc2007	0.947	0.948	0.940	0.941	0.922	0.940	0.689	0.757	0.938	0.869	0.963	0.992
mediamill	0.742	0.753	0.582	0.580	0.569	0.597	0.743	0.739	0.725	0.708	0.788	0.798
bibtex	0.753	0.744	0.734	0.736	0.547	0.359	1.000	0.819	DNF	0.948	0.940	0.957
delicious	0.658	0.660	DNF	DNF	0.396	0.000	0.000	0.651	DNF	DNF	0.589	0.695
bookmarks	DNF	DNF	DNF	DNF	DNF	0.632	0.947	0.850	DNF	DNF	0.878	0.895

Table B.10: The performance of the multi-label learning approaches in terms of the *micro recall* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.406	0.393	0.409	0.431	0.782	0.712	0.539	0.376	0.489	0.531	0.551	0.589
scene	0.694	0.708	0.695	0.692	0.727	0.570	0.521	0.634	0.721	0.751	0.572	0.523
yeast	0.587	0.588	0.595	0.595	0.702	0.603	0.492	0.543	0.602	0.655	0.491	0.521
medical	0.725	0.739	0.782	0.787	0.742	0.720	0.227	0.522	0.600	0.624	0.237	0.569
enron	0.464	0.472	0.532	0.438	0.585	0.440	0.246	0.353	0.435	0.532	0.366	0.422
corel5k	0.057	0.057	0.258	0.258	0.248	0.002	0.000	0.015	0.000	0.001	0.005	0.009
tmc2007	0.917	0.924	0.920	0.920	0.932	0.073	0.454	0.621	0.847	0.869	0.651	0.902
mediamill	0.415	0.385	0.066	0.066	0.537	0.004	0.351	0.432	0.315	0.333	0.418	0.435
bibtex	0.328	0.335	0.322	0.328	0.353	0.053	0.057	0.118	DNF	0.142	0.066	0.131
delicious	0.143	0.144	DNF	DNF	0.297	0.000	0.000	0.101	DNF	DNF	0.174	0.151
bookmarks	DNF	DNF	DNF	DNF	DNF	0.170	0.135	0.135	DNF	DNF	0.112	0.136

Table B.11: The performance of the multi-label learning approaches in terms of the *micro* F_1 measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.509	0.503	0.512	0.528	0.588	0.655	0.571	0.457	0.533	0.554	0.647	0.672
scene	0.761	0.757	0.758	0.756	0.764	0.593	0.516	0.661	0.772	0.762	0.717	0.669
yeast	0.652	0.650	0.655	0.654	0.673	0.610	0.577	0.625	0.656	0.658	0.593	0.617
medical	0.343	0.350	0.721	0.722	0.773	0.756	0.356	0.634	0.714	0.714	0.374	0.693
enron	0.564	0.482	0.585	0.535	0.591	0.512	0.349	0.466	0.548	0.582	0.496	0.537
corel5k	0.059	0.059	0.293	0.293	0.275	0.004	0.000	0.030	0.000	0.002	0.010	0.018
tmc2007	0.932	0.936	0.930	0.930	0.927	0.135	0.547	0.682	0.890	0.869	0.777	0.945
mediamill	0.533	0.509	0.118	0.119	0.553	0.007	0.477	0.545	0.440	0.453	0.546	0.563
bibtex	0.457	0.462	0.448	0.454	0.429	0.093	0.108	0.206	DNF	0.247	0.123	0.230
delicious	0.234	0.236	DNF	DNF	0.339	0.000	0.000	0.175	DNF	DNF	0.269	0.248
bookmarks	DNF	DNF	DNF	DNF	DNF	0.268	0.236	0.232	DNF	DNF	0.199	0.236

Table B.12: The performance of the multi-label learning approaches in terms of the *macro precision* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.721	0.581	0.677	0.660	0.464	0.602	0.628	0.518	0.547	0.531	0.828	0.802
scene	0.844	0.817	0.835	0.832	0.807	0.635	0.682	0.784	0.835	0.785	0.963	0.919
yeast	0.628	0.602	0.614	0.614	0.471	0.377	0.479	0.600	0.480	0.391	0.533	0.674
medical	0.399	0.391	0.288	0.285	0.287	0.263	0.018	0.267	0.269	0.266	0.190	0.269
enron	0.258	0.260	0.205	0.242	0.241	0.142	0.023	0.170	0.222	0.249	0.245	0.233
corel5k	0.052	0.053	0.059	0.059	0.044	0.004	0.000	0.031	0.000	0.001	0.007	0.015
tmc2007	0.972	0.972	0.964	0.965	0.954	0.925	0.386	0.780	0.973	0.938	0.994	0.997
mediamill	0.112	0.144	0.140	0.133	0.107	0.046	0.401	0.308	0.025	0.037	0.397	0.441
bibtex	0.528	0.539	0.503	0.490	0.391	0.128	0.006	0.192	DNF	0.121	0.080	0.127
delicious	0.299	0.303	DNF	DNF	0.154	0.000	0.000	0.134	DNF	DNF	0.422	0.293
bookmarks	DNF	DNF	DNF	DNF	DNF	0.292	0.018	0.414	DNF	DNF	0.388	0.522

Table B.13: The performance of the multi-label learning approaches in terms of the *macro recall* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.378	0.364	0.381	0.398	0.775	0.702	0.533	0.334	0.462	0.508	0.532	0.569
scene	0.703	0.716	0.704	0.701	0.734	0.573	0.529	0.647	0.727	0.757	0.381	0.533
yeast	0.355	0.357	0.361	0.361	0.466	0.375	0.269	0.308	0.352	0.388	0.257	0.286
medical	0.423	0.428	0.307	0.324	0.282	0.249	0.022	0.163	0.183	0.179	0.040	0.176
enron	0.120	0.146	0.139	0.120	0.163	0.107	0.030	0.075	0.097	0.129	0.082	0.100
corel5k	0.023	0.023	0.039	0.039	0.041	0.005	0.000	0.006	0.000	0.001	0.001	0.002
tmc2007	0.915	0.924	0.914	0.914	0.897	0.085	0.235	0.418	0.739	0.772	0.297	0.769
mediamill	0.049	0.044	0.028	0.028	0.074	0.002	0.029	0.088	0.020	0.023	0.065	0.080
bibtex	0.250	0.257	0.236	0.238	0.247	0.034	0.006	0.049	DNF	0.044	0.013	0.043
delicious	0.072	0.075	DNF	DNF	0.103	0.000	0.000	0.039	DNF	DNF	0.092	0.060
bookmarks	DNF	DNF	DNF	DNF	DNF	0.098	0.016	0.070	DNF	DNF	0.048	0.072

Table B.14: The performance of the multi-label learning approaches in terms of the *macro F₁* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.440	0.420	0.443	0.458	0.570	0.630	0.568	0.385	0.488	0.500	0.620	0.650
scene	0.765	0.762	0.762	0.759	0.768	0.596	0.593	0.692	0.777	0.770	0.514	0.658
yeast	0.392	0.390	0.392	0.394	0.447	0.370	0.293	0.336	0.359	0.350	0.283	0.322
medical	0.361	0.371	0.281	0.286	0.282	0.250	0.020	0.192	0.210	0.203	0.058	0.207
enron	0.143	0.153	0.149	0.143	0.167	0.115	0.026	0.087	0.115	0.140	0.102	0.122
corel5k	0.021	0.021	0.042	0.042	0.036	0.008	0.000	0.010	0.000	0.001	0.001	0.004
tmc2007	0.942	0.947	0.938	0.938	0.924	0.124	0.263	0.493	0.826	0.834	0.371	0.857
mediamill	0.056	0.052	0.037	0.037	0.073	0.003	0.031	0.113	0.019	0.022	0.088	0.112
bibtex	0.307	0.316	0.291	0.292	0.266	0.045	0.006	0.065	DNF	0.052	0.016	0.055
delicious	0.096	0.100	DNF	DNF	0.103	0.000	0.000	0.051	DNF	DNF	0.142	0.083
bookmarks	DNF	DNF	DNF	DNF	DNF	0.119	0.017	0.096	DNF	DNF	0.065	0.101

Table B.15: The p-values of the assessment of performance of the multi-label learning approaches by the Friedman test using the label-based evaluation measures. *Subset* shows the calculated p-values for the datasets on which all algorithms finished. *All* shows the calculated p-values for all datasets including those which did not finished.

	All	Subset
Macro precision	$3.5 \cdot 10^{-7}$	$4.8 \cdot 10^{-7}$
Macro recall	$2.8 \cdot 10^{-4}$	$1.1 \cdot 10^{-4}$
Macro F_1	$3.1 \cdot 10^{-4}$	$9.8 \cdot 10^{-5}$
Micro precision	$3.7 \cdot 10^{-9}$	$3.4 \cdot 10^{-8}$
Micro recall	$3.6 \cdot 10^{-4}$	$7.3 \cdot 10^{-5}$
Micro F_1	0.011	0.0022

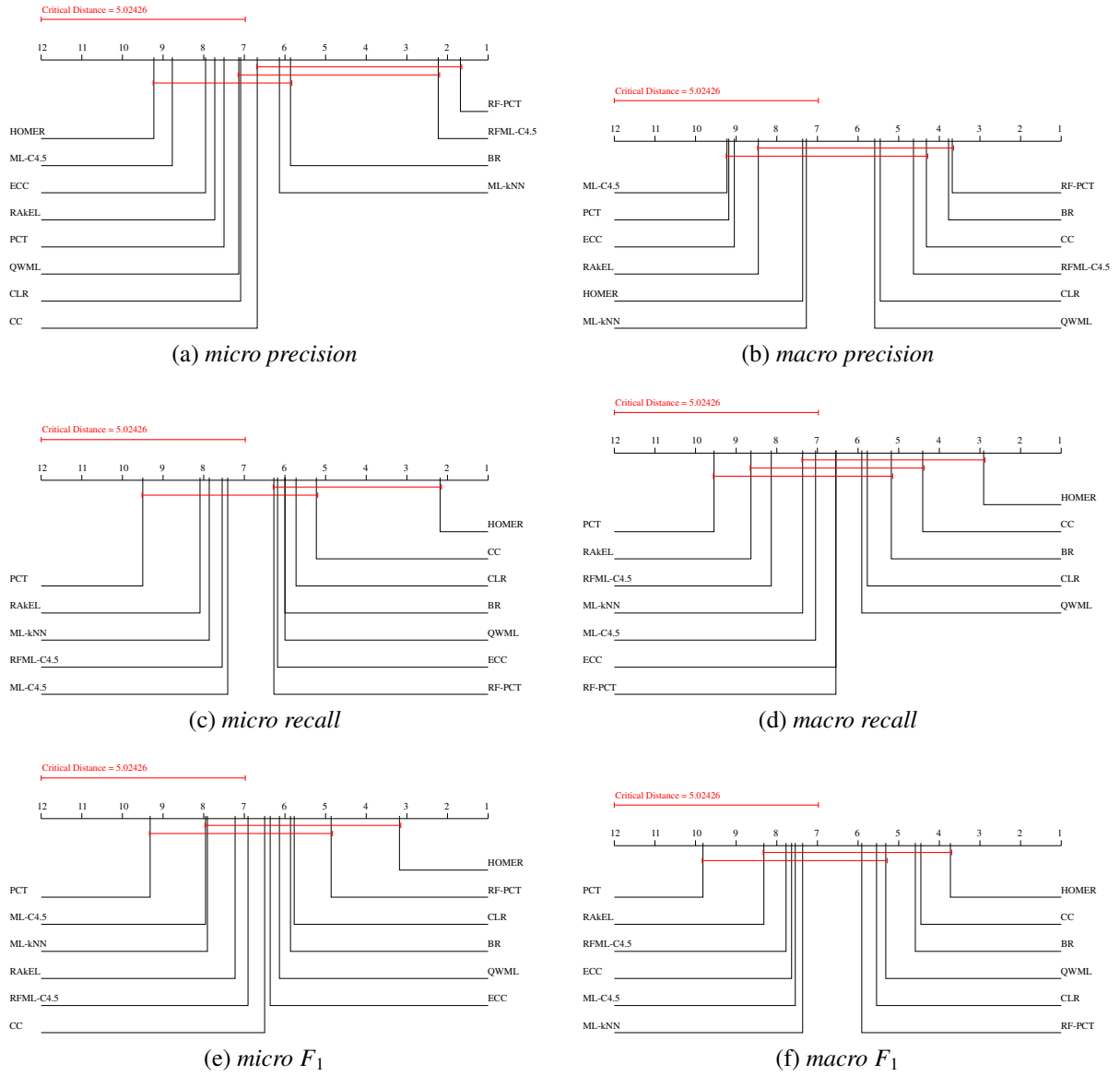


Figure B.9: The critical diagrams for the label-based evaluation measures: The results from the Nemenyi post-hoc test at 0.05 significance level on all the datasets.

Appendix B.3. Results on the ranking-based evaluation measures

Table B.16: The performance of the multi-label learning approaches in terms of the *ranking loss* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.246	0.245	0.264	0.331	0.297	0.210	0.270	0.283	0.281	0.310	0.153	0.151
scene	0.060	0.064	0.065	0.103	0.119	0.169	0.174	0.093	0.104	0.103	0.079	0.072
yeast	0.164	0.170	0.163	0.296	0.205	0.225	0.199	0.172	0.259	0.224	0.173	0.167
medical	0.021	0.019	0.028	0.027	0.090	0.048	0.104	0.045	0.159	0.152	0.028	0.024
enron	0.084	0.083	0.078	0.177	0.183	0.120	0.114	0.093	0.283	0.238	0.083	0.079
corel5k	0.117	0.118	0.100	0.245	0.352	0.479	0.140	0.130	0.673	0.749	0.122	0.117
tmc2007	0.003	0.003	0.005	0.039	0.028	0.043	0.100	0.031	0.031	0.032	0.007	0.006
mediamill	0.061	0.062	0.092	0.101	0.177	0.073	0.063	0.055	0.236	0.258	0.047	0.047
bibtex	0.068	0.067	0.065	0.207	0.255	0.260	0.255	0.217	DNF	0.394	0.126	0.093
delicious	0.114	0.117	DNF	DNF	0.379	0.174	0.172	0.129	DNF	DNF	0.140	0.106
bookmarks	DNF	DNF	DNF	DNF	DNF	0.194	0.258	0.181	DNF	DNF	0.129	0.104

Table B.17: The performance of the multi-label learning approaches in terms of the *one-error* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.386	0.376	0.391	0.391	0.411	0.347	0.386	0.406	0.396	0.426	0.277	0.262
scene	0.180	0.204	0.190	0.193	0.216	0.394	0.389	0.242	0.197	0.213	0.232	0.210
yeast	0.236	0.268	0.229	0.233	0.248	0.312	0.264	0.234	0.254	0.249	0.250	0.248
medical	0.135	0.123	0.168	0.165	0.216	0.198	0.612	0.279	0.312	0.315	0.243	0.174
enron	0.237	0.238	0.231	0.269	0.314	0.309	0.392	0.280	0.290	0.247	0.219	0.221
corel5k	0.660	0.674	0.588	0.592	0.652	0.762	0.776	0.706	0.758	0.992	0.644	0.608
tmc2007	0.029	0.026	0.033	0.033	0.050	0.145	0.306	0.190	0.047	0.052	0.071	0.006
mediamill	0.188	0.193	0.586	0.560	0.219	0.194	0.220	0.182	0.234	0.242	0.171	0.159
bibtex	0.346	0.342	0.388	0.380	0.466	0.529	0.783	0.576	DNF	0.666	0.544	0.433
delicious	0.354	0.367	DNF	DNF	0.509	0.411	0.592	0.416	DNF	DNF	0.368	0.332
bookmarks	DNF	DNF	DNF	DNF	DNF	0.643	0.817	0.639	DNF	DNF	0.607	0.541

Table B.18: The performance of the multi-label learning approaches in terms of the *coverage* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	2.307	2.317	2.386	2.807	2.634	2.069	2.356	2.490	2.465	2.619	1.801	1.827
scene	0.399	0.417	0.423	0.631	0.739	0.945	0.964	0.569	0.635	0.625	0.495	0.461
yeast	6.330	6.439	6.286	8.659	7.285	7.105	6.705	6.414	7.983	7.153	6.276	6.179
medical	1.610	1.471	2.036	1.832	5.324	3.033	5.813	2.844	8.520	7.994	1.889	1.619
enron	12.530	12.437	11.763	22.746	24.190	17.010	14.920	13.181	30.509	27.760	12.485	12.074
corel5k	104.800	105.428	91.506	206.880	250.800	279.900	115.676	113.046	340.398	348.160	110.356	107.412
tmc2007	1.311	1.302	1.363	2.796	2.369	2.671	4.572	2.155	2.498	2.494	1.416	1.219
mediamill	20.481	20.333	24.247	28.982	47.046	22.096	20.456	18.719	56.617	58.865	16.868	16.926
bibtex	20.926	21.078	18.540	57.343	65.626	58.016	58.599	56.266	DNF	87.841	32.580	25.854
delicious	530.126	537.388	DNF	DNF	933.956	620.155	691.622	589.898	DNF	DNF	624.572	504.999
bookmarks	DNF	DNF	DNF	DNF	DNF	58.353	73.780	54.528	DNF	DNF	40.903	34.185

Table B.19: The performance of the multi-label learning approaches in terms of the *average precision* measure. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	0.721	0.724	0.718	0.679	0.698	0.759	0.713	0.694	0.713	0.687	0.812	0.812
scene	0.893	0.881	0.886	0.864	0.848	0.751	0.745	0.851	0.862	0.856	0.862	0.874
yeast	0.768	0.755	0.768	0.698	0.740	0.706	0.724	0.758	0.715	0.734	0.749	0.757
medical	0.896	0.901	0.864	0.862	0.786	0.823	0.522	0.784	0.676	0.684	0.817	0.868
enron	0.693	0.695	0.699	0.604	0.604	0.629	0.546	0.635	0.522	0.576	0.680	0.698
corel5k	0.303	0.293	0.352	0.311	0.222	0.196	0.208	0.266	0.088	0.014	0.314	0.334
tmc2007	0.978	0.981	0.972	0.938	0.945	0.842	0.700	0.844	0.939	0.935	0.945	0.996
mediamill	0.686	0.672	0.450	0.492	0.583	0.669	0.654	0.703	0.492	0.453	0.728	0.737
bibtex	0.597	0.599	0.579	0.498	0.407	0.392	0.212	0.349	DNF	0.228	0.418	0.525
delicious	0.351	0.343	DNF	DNF	0.231	0.321	0.206	0.326	DNF	DNF	0.359	0.395
bookmarks	DNF	DNF	DNF	DNF	DNF	0.378	0.213	0.381	DNF	DNF	0.423	0.480

Table B.20: The p-values of the assessment of performance of the multi-label learning approaches by the Friedman test using the ranking-based evaluation measures. *Subset* shows the calculated p-values for the datasets on which all algorithms finished. *All* shows the calculated p-values for all datasets including those which did not finished.

	All	Subset
One error	$2.2 \cdot 10^{-7}$	$5.3 \cdot 10^{-6}$
Coverage	$1 \cdot 10^{-18}$	$2.3 \cdot 10^{-16}$
Ranking Loss	$1 \cdot 10^{-18}$	$1.2 \cdot 10^{-16}$
Average precision	$6.5 \cdot 10^{-14}$	$2 \cdot 10^{-11}$

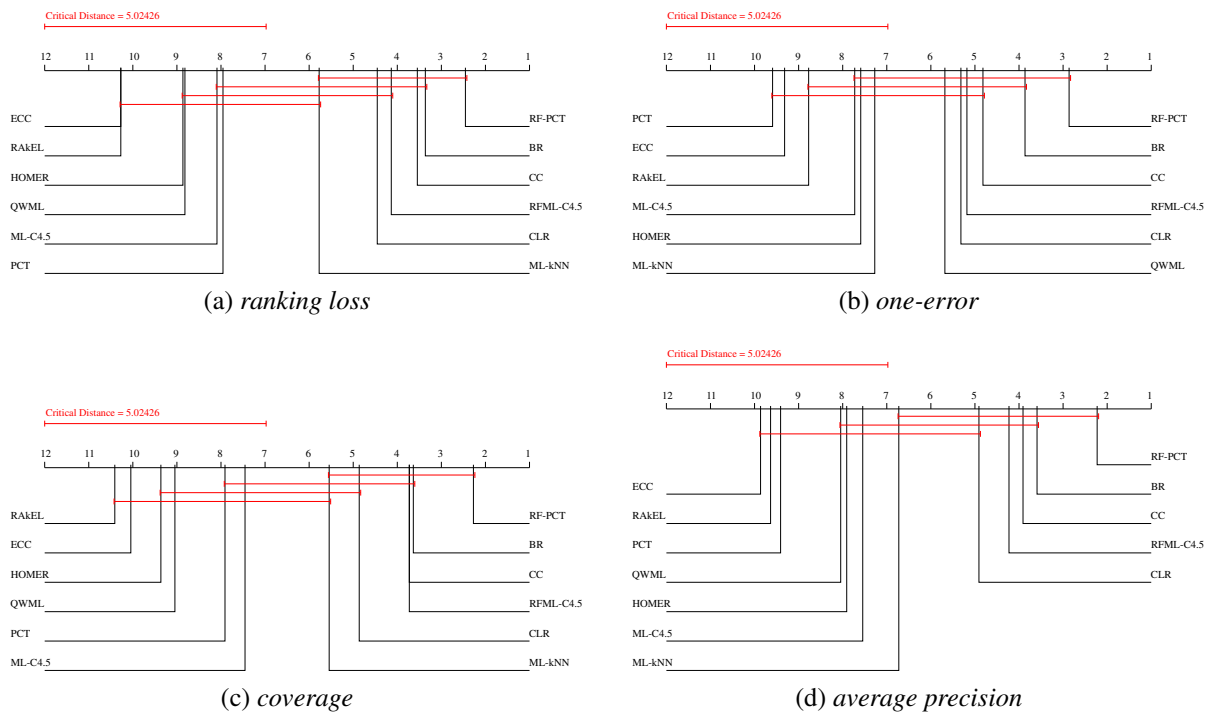


Figure B.10: The critical diagrams for the ranking-based evaluation measures: The results from the Nemenyi post-hoc test at 0.05 significance level on all the datasets.

Appendix B.4. Results on the efficiency of the methods

Table B.21: The performance of the multi-label learning approaches in terms of the *training time* measures in seconds. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	4.0	6.0	10.0	10.0	4.0	0.3	0.1	0.4	5.0	4.9	1.2	2.9
scene	71.0	99.0	195.0	195.0	68.0	8.0	2.0	14.0	79.0	319.0	10.0	23.0
yeast	145.0	206.0	672.0	672.0	101.0	14.0	1.5	8.2	157.0	497.0	19.0	25.0
medical	18.0	28.0	40.0	40.0	16.0	3.0	0.6	1.0	82.0	103.0	7.0	27.0
enron	318.0	440.0	971.0	971.0	158.0	15.0	1.1	6.0	493.0	1467.0	25.0	47.0
corel5k	926.0	1225.0	2388.0	2388.0	771.0	369.0	30.0	389.0	3380.0	20073.0	385.0	902.0
tmc2007	42645.0	46704.0	52427.0	52427.0	31300.0	469.0	11.5	737.0	102394.0	92169.0	460.0	557.0
mediamill	85468.0	100435.0	260156.0	260156.0	78195.0	2030.0	440.0	1094.0	33554.0	188957.0	4056.0	8360.0
bibtex	11013.0	12434.0	13424.0	13424.0	2896.0	566.0	16.4	124.0	DNF	29578.0	645.0	1550.0
delicious	57053.0	84903.0	DNF	DNF	21218.0	2738.0	70.0	236.0	DNF	DNF	21776.0	5376.0
bookmarks	DNF	DNF	DNF	DNF	DNF	4039.0	965.0	15990.0	DNF	DNF	5602.0	28900.0

Table B.22: The performance of the multi-label learning approaches in terms of the *testing time* measures in seconds. DNF (*Did Not Finish*) stands for the algorithms that did not construct a predictive model within one week under the available resources).

	BR	CC	CLR	QWML	HOMER	ML-C4.5	PCT	ML-kNN	RAKEL	ECC	RFML-C4.5	RF-PCT
emotions	1.0	1.0	3.0	2.0	1.0	0.0	0.0	0.4	2.0	6.6	0.1	0.3
scene	25.0	25.0	87.0	40.0	21.0	1.0	0.0	14.0	72.0	168.0	2.0	1.0
yeast	23.0	25.0	153.0	64.0	17.0	0.1	0.0	5.0	70.0	158.0	0.5	0.2
medical	4.0	6.0	90.0	25.0	1.5	0.1	0.0	0.2	24.0	46.0	0.5	0.5
enron	50.0	53.0	634.0	174.0	22.0	0.2	0.0	3.0	153.0	696.0	1.0	1.0
corel5k	25.0	31.0	2161.0	119.0	14.0	1.0	1.0	45.0	3613.0	2077.0	1.8	2.5
tmc2007	927.0	891.0	3282.0	1543.0	730.0	1.7	0.0	230.0	10985.0	10865.0	3.4	2.8
mediamill	6152.0	6125.0	76385.0	20317.0	6079.0	1.0	1.0	477.0	39001.0	50183.0	8.0	4.0
bibtex	654.0	661.0	16733.0	4710.0	155.0	6.5	0.0	64.0	DNF	10756.0	12.0	18.0
delicious	2045.0	1872.0	DNF	DNF	816.0	19.0	10.0	55.0	DNF	DNF	32.0	48.0
bookmarks	DNF	DNF	DNF	DNF	DNF	21.0	15.0	4084.0	DNF	DNF	28.0	58.0

Table B.23: The p-values of the assessment of performance of the multi-label learning approaches by the Friedman test using the efficiency measures. *Subset* shows the calculated p-values for the datasets on which all algorithms finished. *All* shows the calculated p-values for all datasets including those which did not finished.

	All	Subset
Training time	$1 \cdot 10^{-18}$	$1 \cdot 10^{-18}$
Testing time	$1 \cdot 10^{-18}$	$1 \cdot 10^{-18}$

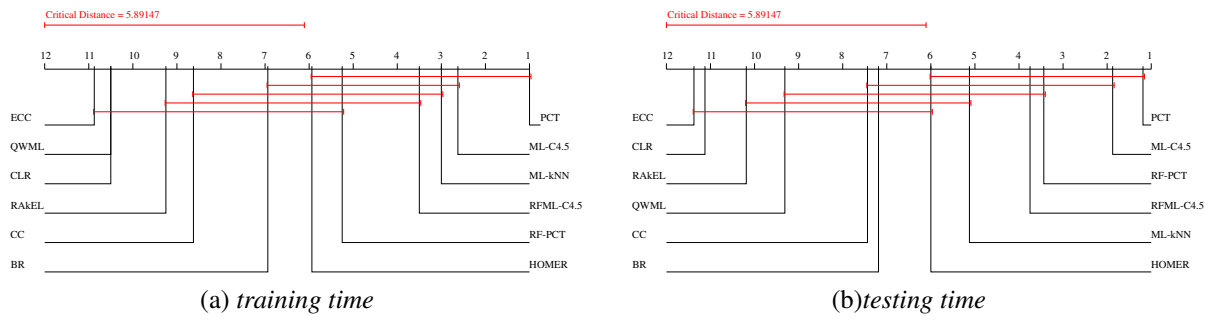


Figure B.11: The critical diagrams for the efficiency measures: The results from the Nemenyi post-hoc test at 0.05 significance level on all the datasets.