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Learning to Rank for Information Retrieval

By Tie-Yan Liu

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Abstract

Learning to rank for Information Retrieval (IR) is a task to automatically construct a ranking model using training data, such that the model can sort new objects according to their degrees of relevance, preference, or importance. Many IR problems are by nature ranking problems, and many IR technologies can be potentially enhanced by using learning-to-rank techniques. The objective of this tutorial is to give an introduction to this research direction. Specifically, the existing learning-to-rank algorithms are reviewed and categorized into three approaches: the pointwise, pairwise, and listwise approaches. The advantages and disadvantages with each approach are analyzed, and the relationships between the loss functions used in these approaches and IR evaluation measures are discussed. Then the empirical evaluations on typical learning-to-rank methods are shown, with the LETOR collection as a benchmark dataset, which seems to suggest that the listwise approach be the most effective one among all the approaches. After that, a statistical ranking theory is introduced, which can describe different learning-to-rank algorithms, and be used to analyze their querylevel generalization abilities. At the end of the tutorial, we provide a summary and discuss potential future work on learning to rank.

Introduction

With the fast development of the Web, every one of us is experiencing a flood of information. It was estimated that there are about 25 billion pages on the Web as of October 2008,¹ which makes it generally impossible for common users to locate desired information by browsing the Web. As a consequence, efficient and effective Information Retrieval (IR) has become more important than ever, and search engines (or IR systems) have become an essential tool for many people.

Ranking is a central problem in IR. Many IR problems are by nature ranking problems, such as document retrieval, collaborative filtering [58], key term extraction [30], definition finding [130], important email routing [23], sentiment analysis [94], product rating [36], and anti Web spam [56]. In this tutorial, we will mainly take document retrieval as an example. Note that document retrieval is not a narrow task. Web pages, emails, academic papers, books, and news articles are just a few of the many examples of documents. There are also many different ranking scenarios for document retrieval of our interest.

¹ http://www.worldwidewebsize.com/

Scenario 1: Rank the documents purely according to their relevance with regards to the query.

Scenario 2: Consider the relationships of similarity [117], website structure [35], and diversity [139] between documents in the ranking process. This is also referred to as relational ranking [103].

Scenario 3: Aggregate several candidate ranked lists to get a better ranked list. This scenario is also referred to as meta search [7]. The candidate ranked lists may come from different index servers or different vertical search engines, and the target ranked list is the final result presented to users.

Scenario 4: Find whether and to what degree a property of a webpage influences the ranking result. This is referred to as "reverse engineering" in search engine optimization (SEO).²

To tackle the problem of document retrieval, many heuristic ranking models have been proposed and used in IR literature. Recently, given the amount of potential training data available, it has become possible to leverage Machine Learning (ML) technologies to build effective ranking models. Specifically, we call those methods that learn how to combine predefined features for ranking by means of discriminative learning "learning-to-rank" methods.

In recent years, learning to rank has become a very hot research direction in IR, and a large number of learning-to-rank algorithms have been proposed, such as [9, 13, 14, 16, 17, 26, 29, 33, 34, 47, 49, 59, 63, 73, 78, 90, 97, 99, 102, 114, 119, 122, 129, 134, 136]. We foresee that learning to rank will have an even bigger impact on IR in the future.

When a research area comes to this stage, several questions naturally arise.

- To what respect are these learning-to-rank algorithms similar and in which aspects do they differ? What are the strengths and weaknesses of each algorithm?
- Empirically, which of those many learning-to-rank algorithms perform the best? What kind of datasets can be used to make fair comparison among different learning-to-rank algorithms?



² http://www.search-marketing.info/newsletter/reverse-engineering.htm

- Theoretically, is ranking a new ML problem, or can it be simply reduced to existing ML problems? What are the unique theoretical issues for ranking that should be investigated?
- Are there many remaining issues regarding learning to rank to study in the future? What are they?

The above questions have been brought to the attention of the IR and ML communities in a variety of contexts, especially during recent years. The aim of this tutorial is to review the recent work that attempts to answer these questions. Needless to say, the comprehensive understanding of the task of ranking in IR is the key to finding the right answers. Therefore, we will first give a brief introduction of ranking in IR, and then formalize the problem of learning to rank so as to set the stage for the upcoming detailed reviews.

1.1 Ranking in IR

In this subsection, we briefly review representative ranking models in IR literature, and introduce how these models are evaluated.

1.1.1 Conventional Ranking Models for IR

In IR literature, many ranking models have been proposed [8]; they can be roughly categorized as query-dependent models and query-independent models.

Query-dependent models

The early models retrieve documents based on the occurrences of the query terms in the documents. Examples include the *Boolean model* [8]. Basically these models can only predict whether a document is relevant to the query or not, but cannot predict the degree of relevance.

To further model the relevance degree, the *Vector Space model* (VSM) was proposed [8]. Both documents and queries are represented as vectors in a Euclidean space, in which the inner product of two vectors can be used to measure their similarities. To get an effective vector representation of queries and documents, TF–IDF weighting has been

widely used.³ The TF of term t in a vector is defined as the normalized number of its occurrences in the document, and the IDF of it is defined as follows:

$$\overline{\text{IDF}(t)} = \log \frac{N}{n(t)},\tag{1.1}$$

where N is the total number of documents in the collection, and n(t) is the number of documents containing term t.

While VSM implies the assumption on the independence between terms, Latent Semantic Indexing (LSI) [37] tries to avoid this assumption. In particular, Singular Value Decomposition (SVD) is used to linearly transform the feature space and thus a "latent semantic space" is generated. Similarity in this new space is then used to define the relevance between queries and documents.

As compared with the above, models based on the probabilistic ranking principle [83] garnered more attention and achieved more success in past decades. The famous ranking models like BM25 [111]⁴ and language model for IR can both be categorized as probabilistic ranking models.

The basic idea of BM25 is to rank documents by the log-odds of their relevance. Actually BM25 is not a single model, but it defines a whole family of ranking models, with slightly different components and parameters. One of the popular instantiations of the model is as follows.

Given query q, containing terms t_1, \ldots, t_M , the BM25 score of document d is computed as below:

$$BM25(d,q) = \sum_{i=1}^{M} \frac{IDF(t_i) \cdot TF(t_i, d) \cdot (k_1 + 1)}{TF(t_i, d) + k_1 \cdot \left(1 - b + b \cdot \frac{LEN(d)}{avdl}\right)},$$
(1.2)

where TF(t,d) is the term frequency of t in document d; LEN(d) is the length (number of words) of document d; avdl is the average document length in the text collection from which documents are drawn; k_1 and

³ Note that there are many different definitions of TF and IDF in IR literature. Some are purely based on the frequency and the others include smoothing or normalization [116]. Here we just give some simple examples to illustrate the main idea.

⁴The name of the actual model is BM25. However, it is usually referred to as "OKapi BM25", since the OKapi system was the first system to implement this model.

b are free parameters; IDF(t) is the IDF weight of term t, computed by using Equation (1.1), for example.

Language model for IR [96] is an application of the statistical language model on IR. A statistical language model assigns a probability to a sequence of terms. When used in IR, a language model is associated with a document. With query q as input, documents are ranked based on the query likelihood, or the probability that the document's language model would generate the terms in the query (i.e., P(q|d)). By further assuming the independence among terms, one has $P(q|d) = \prod_{i=1}^{M} P(t_i|d)$, if query q contains terms t_1, \ldots, t_M .

To learn the document's language model, a maximum likelihood method is used. As in many maximum likelihood methods, the issue of smoothing the estimate is critical. Usually a background language model estimated using the entire collection is used for this purpose. Then, the document's language model can be constructed as follows:

$$p(t_i|d) = (1-\lambda)\frac{\mathrm{TF}(t_i,d)}{\mathrm{LEN}(d)} + \lambda p(t_i|C), \tag{1.3}$$

where $p(t_i|C)$ is the background language model for term t_i , and $\lambda \in [0,1]$ is a smoothing factor.

There are many variants of language model for IR, some of them even go beyond the query likelihood retrieval model (e.g., the models based on K-L divergence [140]). We will not introduce more about them, and readers are encouraged to read the tutorial authored by Zhai [138].

In addition to the above examples, many other models have also been proposed to compute the relevance between a query and a document. Some of them [118] take the proximity of the query terms into consideration, and some others consider the relationship between documents in terms of content similarity [117], hyperlink structure [113], website structure [101], and topic diversity [139].

Query-independent models

In IR literature, there are also many models that rank documents based on their own importance. We will take PageRank [92] as an example for illustration. This model is particularly applicable to Web search because it makes use of the hyperlink structure of the Web for ranking. PageRank uses the probability that a surfer randomly clicking on links will arrive at a particular webpage to rank the web pages. In the general case, the PageRank value for any page d_u can be expressed as:

$$PR(d_u) = \sum_{d_v \in B_u} \frac{PR(d_v)}{U(d_v)}.$$
(1.4)

That is, the PageRank value for page d_u is dependent on the PageRank values for each page d_v out of the set B_u (containing all pages linking to page d_u), divided by $U(d_v)$, the number of outlinks from page d_v .

To get a meaningful solution to Equation (1.4), a smoothing term is introduced. When the random surfer walks on the link graph, she/he does not necessarily always follow the existing hyperlinks. There is a small probability that she/he will jump to any other page uniformly. This small probability can be represented by $(1 - \alpha)$, where α is called the damping factor. Accordingly, PageRank is refined as follows:

$$(d_u) = \alpha \sum_{d_v \in B_u} \frac{PR(d_v)}{U(d_v)} + \frac{(1-\alpha)}{N}, \qquad (1.5)$$

where N is the total number of pages on the Web.

There is much work discussing the theoretical properties, variations, and efficient implementations of PageRank. Furthermore, there are also many other link analysis algorithms, such as Hyperlink Induced Topic Search (HITS) [72] and TrustRank [57]. Some of these algorithms even leverage the content or topic information in the process of link analysis [91].

1.1.2 Query-level Position-based Evaluations in IR

Given the large number of ranking models as introduced in the previous subsection, a standard evaluation mechanism is needed to select the most effective model. For this purpose, one usually proceeds as follows:

 Collect a large number of (randomly sampled) queries to form a test set.

- For each query q,
 - Collect documents $\{d_j\}_{j=1}^m$ associated with the query.
 - Get the relevance judgment for each document by human assessment.
 - Use a given ranking model to rank the documents.
 - Measure the difference between the ranking results and the relevance judgment using an evaluation measure.
- Use the average measure on all the queries in the test set to evaluate the performance of the ranking model.

As for collecting the documents associated with a query, a number of strategies can be used. For example, one can simply collect all the documents containing the query word. One can also choose to use some predefined rankers to get documents that are more likely to be relevant. A popular strategy is the pooling method used in TREC.⁵ In this method a pool of possibly relevant documents is created by taking a sample of documents selected by various participating systems. In particular, the top 100 documents retrieved in each submitted run for a given query are selected and merged into the pool for human assessment. On average, an assessor judges the relevance of approximately 1500 documents per query.

As for the relevance judgment, three strategies were used in the literature.

(1) Specifying whether a document is relevant or not to the query (i.e., binary judgment 1 or 0), or further specifying the degree of relevance (i.e., multiple ordered categories, e.g., Perfect, Excellent, Good, Fair, or Bad). Suppose for document d_j associated with query q, we get its relevance judgment as l_j . Then for two documents d_u and d_v , if $l_u \succ l_v$, we say that document d_u is more relevant than document d_v , with regards to query q, according to the relevance judgment.



⁵ http://trec.nist.gov/

- (2) Specifying whether a document is more relevant than another with regards to a query. For example, if document d_u is judged to be more relevant than document d_v with regards to query q, we give the judgment $l_{u,v} = 1$; otherwise, $l_{u,v} = -1$. That is, this kind of judgment captures the relative preference between documents.⁶
- (3) Specifying the partial order or even total order of the documents with respect to a query. For the group of documents $\{d_i\}_{i=1}^m$ associated with query q, this kind of judgment is usually represented as a certain permutation of these documents, denoted as π_l , or a set of such permutations.

Given the vital role that relevance judgments play in a test collection, it is important to assess the quality of the judgments. In previous practices like TREC, both the completeness and the consistency of relevance judgments are of interest. Completeness measures the degree to which all the relevant documents for a topic have been found; consistency measures the degree to which the assessor has marked all the "truly" relevant documents relevant and the "truly" irrelevant documents irrelevant.

Since manual judgment is time consuming, it is almost impossible to judge all the documents with regards to a query. Consequently, there are always unjudged documents returned by the ranking model. As a common practice, one regards the unjudged documents as irrelevant in the evaluation process.⁷

With the relevance judgment, several evaluation measures have been proposed and used in IR literature. It is clear that understanding these measures will be very important for learning to rank, since to some extent they define the "true" objective function of ranking. Below we list some popularly used measures.

Mean reciprocal rank (MRR): For query q, the rank position of its first relevant document is denoted as r(q). Then $\frac{1}{r(q)}$ is defined as MRR for

 $^{^6}$ This kind of judgment can also be mined from click-through logs of search engines

⁷ In recent years, several new evaluation mechanisms [18] that consider the relevance probability of an unjudged document have also been proposed.

query q. It is clear that documents ranked below r(q) are not considered in MRR.

Mean average precision (MAP): To define MAP [8], one needs to define Precision at position k (P@k) first,

$$P@k(q) = \frac{\#\{\text{relevant documents in the top } k \text{ positions}\}}{k}.$$
 (1.6)

Then, the Average Precision (AP) is defined below:

$$AP(q) = \frac{\sum_{k=1}^{m} P@k(q) \cdot l_k}{\#\{\text{relevant documents}\}},$$
(1.7)

where m is the total number of documents associated with query q, and l_k is the binary judgment on the relevance of the document at the k-th position. The mean value of AP over all the test queries is named MAP.

Discounted cumulative gain (DCG): While the aforementioned measures are mainly designed for binary judgments, DCG [65, 66] can leverage the relevance judgment in terms of multiple ordered categories, and has an explicit position discount factor in its definition. More formally, suppose the ranked list for query q is π , then DCG at position k is defined as follows:

$$G@k(q) = \sum_{r=1}^{k} G(\pi^{-1}(r))\eta(r), \qquad (1.8)$$

where $\pi^{-1}(r)$ denotes the document ranked at position r of the list π , $G(\cdot)$ is the rating of a document (one usually sets $G(\pi^{-1}(r)) = (2^{l_{\pi^{-1}(r)}} - 1)$), and $\eta(r)$ is a position discount factor (one usually sets $\eta(r) = 1/\log_2(r+1)$).

By normalizing DCG@k with the maximum value of it (denoted as Z_k), we will get another measure named Normalized DCG (NDCG). That is:

NDCG@
$$k(q) = \frac{1}{Z_k} \sum_{r=1}^k G(\pi^{-1}(r)) \eta(r).$$
 (1.9)

It is clear that NDCG takes values from 0 to 1.

Rank correlation (RC): The correlation between the ranked list given by the model (denoted as π) and the relevance judgment

(denoted as π_l) can be used to define a measure. For example, when the weighted Kendall's τ is used, the RC measures the weighted pairwise inconsistency between two lists. Its definition is given below:

$$\frac{\tau_K(q) = \frac{\sum_{u < v} w_{u,v} (1 + \operatorname{sgn}((\pi(u) - \pi(v))(\pi_l(u) - \pi_l(v))))}{2\sum_{u < v} w_{u,v}}, \quad (1.10)$$

where $w_{u,v}$ is the weight, and $\pi(u)$ means the rank position of document d_u in list π .

To summarize, there are some common properties in these evaluation measures.

- (1) All these evaluation measures are calculated at the query level. That is, first the measure is computed for each query, and then averaged over all queries in the test set. No matter how poorly the documents associated with a particular query are ranked, it will not dominate the evaluation process since each query contributes similarly to the average measure.
- (2) All these measures are position based. That is, rank position is explicitly used. Considering that with small changes in the scores given by a ranking model the rank positions will not change until one document's score passes another, the position-based measures are usually non-continuous and non-differentiable with regards to the scores. This makes the optimization of these measures quite difficult. We will conduct more discussions on this in Section 4.1.

1.2 Learning to Rank

Many ranking models have been introduced in the previous subsection, most of which contain parameters. For example, there are parameters k_1 and b in BM25 (see Equation (1.2)), parameter λ in language model for IR (see Equation (1.3)), and parameter α in PageRank (see Equation (1.5)). In order to get a reasonably good ranking performance (in terms of IR evaluation measures), one needs to tune these parameters using a validation set. Nevertheless, the parameter tuning is far from trivial, especially considering that IR evaluation measures are non-continuous and non-differentiable with respect to the parameters.



In addition, a model perfectly tuned on the validation set sometimes performs poorly on unseen test queries. This is usually called overfitting. Another issue is about the combination of these ranking models. Given that many models have been proposed in the literature, it is natural to investigate how to combine these models and create an even more effective new model. This is, however, not straightforward either.

While IR researchers were facing these problems, machine learning has been demonstrating its effectiveness in automatically tuning parameters, combining multiple evidences, and avoiding over-fitting. Therefore, it seems quite promising to adopt ML technologies to solve the aforementioned problems.

1.2.1 ML Framework

In much ML research (especially discriminative learning), attention has been paid to the following key components.⁸

- (1) The *input space*, which contains the objects under investigation: Usually objects are represented by feature vectors, extracted according to different applications.
- (2) The output space, which contains the learning target with respect to the input objects: There are two related but different definitions of the output space in ML.⁹ The first is the output space of the task, which is highly dependent on the application. For example, in the regression problem the output space is the space of real numbers \mathbb{R} ; in classification, it is the set of discrete categories $\{0,1,\ldots,K-1\}$. The second is the output space to facilitate the learning process. This may differ from the output space of the task. For example, one can use regression algorithms to solve the problem of classification. In this case, the output space that facilitates learning is the space of real numbers but not discrete categories.
- (3) The *hypothesis* space, which defines the class of functions mapping the input space to the output space: The functions

⁸ For a more comprehensive introduction to the ML literature, please refer to [89].

⁹ In this tutorial, when we mention the output space, we mainly refer to the second type.

- operate on the feature vectors of the input objects, and make predictions according to the format of the output space.
- (4) In order to learn the optimal hypothesis, a training set is usually used, which contains a number of independent and identically distributed (i.i.d.) objects and their ground truth labels, sampled from the product of the input and output spaces. The loss function measures to what degree the prediction generated by the hypothesis is in accordance with the ground truth label. For example, widely used loss functions for classification include the exponential loss, the hinge loss, and the logistic loss. It is clear that the loss function plays a central role in ML, since it encodes the understanding of the target application (i.e., what prediction is correct and what is not). With the loss function, an empirical risk can be defined on the training set, and the optimal hypothesis is usually (but not always) learned by means of empirical risk minimization.

1.2.2Learning-to-Rank Framework

In recent years, more and more ML technologies have been used to train the ranking model, and a new research area named "learning to rank" has gradually emerged. Especially in the past several years, learning to rank has become one of the most active research areas in IR.

In general, we can call all those methods that use ML technologies to solve the problem of ranking "learning-to-rank" methods, 10 such as the work on relevance feedback¹¹ [39, 112] and automatically tuning the parameters of existing IR models [60, 120]. However, most of the state-of-the-art learning-to-rank algorithms learn the optimal way of combining features extracted from query-document pairs through discriminative training. Therefore, in this tutorial we define learning to rank in a more specific way to better summarize these algorithms.

 $^{^{10}}$ In ML literature, there is a topic named label ranking. It is to predict the ranking of multiple class labels for an individual document, but not to predict the ranking of documents. In this regard, it is largely different from the task of ranking for IR.

¹¹We will make further discussions on the relationship between relevance feedback and learning to rank in Section 2.

We call those ranking methods that have the following two properties learning-to-rank methods.

Feature based: All the documents under investigation are represented by feature vectors, 12 reflecting the relevance of the documents to the query. That is, for a given query q, its associated document d can be represented by a vector $x = \Phi(d,q)$, where Φ is a feature extractor. Typical features used in learning to rank include the frequencies of the query terms in the document, the BM25 and PageRank scores, and the relationship between this document and other documents. If one wants to know more about widely used features, please refer to Tables 6.2 and 6.3 in Section 6.

Even if a feature is the output of an existing retrieval model, in the context of learning to rank, one assumes that the parameter in the model is fixed, and only the optimal way of combining these features is learned. In this sense, the previous work on automatically tuning the parameters of existing models [60, 120] is not categorized as "learningto-rank" methods.

The capability of combining a large number of features is a very important advantage of learning to rank. It is easy to incorporate any new progress on the retrieval model by including the output of the model as one dimension of the features. Such a capability is highly demanding for real search engines, since it is almost impossible to use only a few factors to satisfy complex information needs of Web users.

Discriminative training: The learning process can be well described by the four components of discriminative learning as mentioned in the previous subsection. That is, a learning-to-rank method has its specific input space, output space, hypothesis space, and loss function.

In ML literature, discriminative methods have been widely used to combine different kinds of features, without the necessity of defining a probabilistic framework to represent the objects and the correctness of prediction. In this sense, previous works that train generative ranking



 $^{^{12}}$ Note that, hereafter in this tutorial, when we refer to a document, we will not use d any longer. Instead, we will directly use its feature representation x. Furthermore, since our discussions will focus more on the learning process, we will always assume the features are pre-specified, and will not purposely discuss how to extract them.

models are not categorized as "learning-to-rank" methods in this tutorial. If one has interest in such work, please refer to [74, 85, 141], etc.

Discriminative training is an automatic learning process based on the training data. This is also highly demanding for real search engines, because everyday these search engines will receive a lot of user feedback and usage logs indicating poor ranking for some queries or documents. It is very important to automatically learn from feedback and constantly improve the ranking mechanism.

Due to the aforementioned two characteristics, learning to rank has been widely used in commercial search engines, ¹³ and has also attracted great attention from the academic research community.

Figure 1.1 shows the typical "learning-to-rank" flow. From the figure we can see that since learning to rank is a kind of supervised learning, a training set is needed. The creation of a training set is very similar to

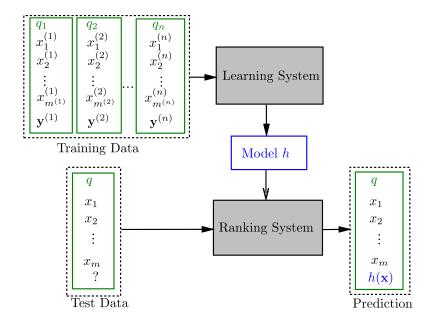


Fig. 1.1 Learning-to-rank framework.

 $^{^{13}}$ See http://blog.searchenginewatch.com/050622-082709, http://blogs.msdn.com/msnsearch/archive/2005/06/21/431288.aspx, and http://glinden.blogspot.com/2005/06/msn-search-and-learning-to-rank.html.

the creation of the test set for evaluation. For example, a typical training set consists of n training queries $q_i(i=1,\ldots,n)$, their associated documents represented by feature vectors $\mathbf{x}^{(i)} = \{x_j^{(i)}\}_{j=1}^{m^{(i)}}$ (where $m^{(i)}$ is the number of documents associated with query q_i), and the corresponding relevance judgments. Then a specific learning algorithm is employed to learn the ranking model (i.e., the way of combining the features), such that the output of the ranking model can predict the ground truth label in the training set a accurately as possible, in terms of a loss function. In the test phase, when a new query comes in, the model learned in the training phase is applied to sort the documents according to their relevance to the query, and return the corresponding ranked list to the user as the response to her/his query.

1.2.3 Approaches to Learning to Rank

Many learning-to-rank algorithms can fit into the above framework. In order to better understand them, we perform a categorization on these algorithms. In particular, we group the algorithms, according to the four pillars of ML, into three approaches: the pointwise approach, the pairwise approach, and the listwise approach. Note that different approaches model the process of learning to rank in different ways. That is, they define different input and output spaces, use different hypotheses, and employ different loss functions. Note that the output space is used to facilitate the learning process, which can be different from the relevance judgments on the documents. That is, even if provided with the same format of judgments, one can derive different ground truth labels from it, and use them for different approaches.

The pointwise approach

The *input space* of the pointwise approach contains the feature vector of each single document.



¹⁴ Please distinguish between the judgment for evaluation and the judgment for constructing the training set, although the processes of obtaining them may be very similar.

¹⁵ Hereafter, when we mention the ground truth labels in the remainder of the tutorial, we will mainly refer to the ground truth labels in the training set, although we assume every document has its intrinsic label no matter whether it is judged or not.

The *output space* contains the relevance degree of each single document. The ground truth label in the output space is usually defined in the following way. If the judgment is directly given as relevance degree l_i , the ground truth label for document x_i is defined as $y_i = l_i$. If the judgment is given as total order π_l , one can get the ground truth label by using a mapping function. 16 However, if the judgment is given as pairwise preference $l_{u,v}$, it is not straightforward to make use of it to generate the ground truth label.

The hypothesis space contains functions that take the feature vector of a document as the input and predict the relevance degree of the document. We usually call such a function f the scoring function. Note that, based on the scoring function, one can sort all the documents and produce the final ranked list.

The loss function examines the accurate prediction of the ground truth label for each single document. In different pointwise ranking algorithms, ranking is modeled as regression, classification, and ordinal regression (see Section 2). Therefore the corresponding regression loss, classification loss, and ordinal regression loss are used as the loss function. Note that the pointwise approach does not consider the interdependency among documents, and thus the position of a document in the final ranked list is invisible to its loss function. Furthermore, the approach does not make use of the fact that some documents are actually associated with the same query. Considering that most IR evaluation measures are query-level and position-based, intuitively speaking, the pointwise approach has its limitations.

Example algorithms belonging to the pointwise approach include [24, 25, 26, 31, 33, 34, 49, 53, 73, 78, 90, 114]. We will introduce some of them in Section 2.

The pairwise approach

The *input space* of the pairwise approach contains a pair of documents, both represented as feature vectors.

The *output space* contains the pairwise preference (which takes values from $\{1,-1\}$) between each pair of documents. The ground truth

 $[\]overline{}^{16}$ For example, the position of the document in π_l can be used to define the relevance degree.

label in the output space is usually defined in the following way. If the judgment is given as relevance degree l_j , then the order for document pair (x_u, x_v) can be defined as $y_{u,v} = 2 \cdot I_{\{l_u \succ l_v\}} - 1$. Here $I_{\{A\}}$ is an indicator function, which is defined to be 1 if predicate A holds and 0 otherwise. If the judgment is given directly as pairwise preference $l_{u,v}$, then it is straightforward to set $y_{u,v} = l_{u,v}$. If the judgment is given as total order π_l , one can define $y_{u,v} = 2 \cdot I_{\{\pi_l(u) < \pi_l(v)\}} - 1$.

The hypothesis space contains bi-variate functions h that take a pair of documents as the input and output the relative order between them. Some pairwise ranking algorithms directly define their hypotheses as such [29], however, in more algorithms, the hypothesis is still defined with a scoring function f for simplicity, i.e., $h(x_u, x_v) = 2 \cdot I_{\{f(x_u) > f(x_v)\}} - 1$.

The loss function measures the inconsistency between $h(x_u, x_v)$ and the ground truth label $y_{u,v}$. For example, in some algorithms, ranking is modeled as a pairwise classification, and the corresponding classification loss on a pair of documents is used as the loss function. Note that the loss function used in the pairwise approach only considers the relative order between two documents. When one looks at only a pair of documents, however, the position of the documents in the final ranked list can hardly be derived. Furthermore, the approach ignores the fact that some pairs are generated from the documents associated with the same query. Considering that most IR evaluation measures are query-level and position-based, intuitively speaking, there is still a gap between this approach and ranking for IR.

Example algorithms belonging to the pairwise approach include [9, 14, 16, 29, 47, 63, 97, 122]. We will introduce some of them in Section 3.

The listwise approach

The *input space* of the listwise approach contains the entire group of documents associated with query q, e.g., $\mathbf{x} = \{x_j\}_{j=1}^m$.

There are two types of *output spaces* used in the listwise approach. For some listwise ranking algorithms, the *output space* contains the relevance degrees of all the documents associated with a query. In this case, the ground truth label $\mathbf{y} = \{y_j\}_{j=1}^m$ can be derived from the judgment

in terms of the relevance degree or total order, in a similar manner to that of the pointwise approach. For some other listwise ranking algorithms, the *output space* contains the ranked list (or permutation) of the documents. In this case, the ground truth label, denoted as π_y , can be generated in the following way. When the judgment is given as total order π_l , we can define $\pi_y = \pi_l$. Otherwise, we can derive π_y by using the concept of the *equivalent permutation set* (see Section 4). When π_y is given as the ground truth label, the output space that facilitates the learning process is exactly the output space of the ranking task. Therefore, the theoretical analysis on the listwise approach has a more direct value where understanding the real ranking problem than the other approaches where there are mismatches between the output space that facilitates learning and the real output space of the task.

The hypothesis space contains multivariate functions h that operate on a group of documents, and predict their relevance degrees or their permutation. For practical reasons, the hypothesis h is also usually implemented with scoring function f. When the relevance degree comprises the output space, $h(\mathbf{x}) = f(\mathbf{x})$. When the ranked list (permutation) comprises the output space, h is defined as a compound function $h(\mathbf{x}) = \text{sort} \circ f(\mathbf{x})$. That is, first scoring function f is used to give a score to each document, and then these documents are sorted in the descending order of the scores to produce the desired ranked list.

There are also two types of loss functions, corresponding to the two types of output spaces. When the ground truth label is given as \mathbf{y} , the loss function is usually defined on the basis of the approximation or bound of widely used IR evaluation measures. When the ground truth label is given as π_y , the loss function measures the difference between the ranked list given by the hypothesis and the ground truth list. As compared to the pointwise and pairwise approaches, the advantage of the listwise approach lies in that its loss function can naturally consider the positions of documents in the ranked list of all the documents associated with the same query.

Example algorithms that belong to the listwise approach include [13, 17, 99, 102, 119, 129, 134, 136]. We will introduce some of them in Section 4.

It is noted that different loss functions are used in different approaches, while the same IR evaluation measures are used for testing their performances. A natural question that arises concerns the relationship between these loss functions and IR evaluation measures. The investigation on this issue can help us explain the empirical results of learning-to-rank algorithms. We will introduce some such investigations in Section 5. In addition, in Section 6, we will introduce a benchmark dataset for the research on learning to rank, named LETOR, and report some empirical results of representative learning-to-rank algorithms on the dataset.

Furthermore, one may have noticed that the scoring function, which is widely used in defining the hypotheses of different approaches, is a kind of "pointwise" function. However, it is not to say that all the approaches are in nature pointwise approaches. The categorization of the aforementioned three approaches is based on the four pillars of ML. That is, different approaches regard the same training data as in different input and output spaces, and define different loss functions and hypotheses accordingly. From the ML point of view, they have different assumptions on the i.i.d. distribution of the data and therefore the theoretical properties (e.g., generalization ability) of their corresponding algorithms will be largely different. We will further discuss this in Section 7, with the introduction of a new theory, which we call the statistical ranking theory.

1.3 About this Tutorial

As for the writing of the tutorial, we do not aim to be fully rigorous. Instead, we try to provide insights into the basic ideas. However, it is still unavoidable that we will use mathematics for better illustration of the problem, especially when we jump into the theoretical discussions on learning to rank. We will have to assume familiarity with basic concepts of probability theory and statistical learning in the corresponding discussions.

Furthermore, we will use the notation rules as listed in Table 1.1 throughout the tutorial. Here we would like to add one more note. Since in practice the hypothesis h is usually defined with scoring function f,

Table 1.1 Notation rules.

Meaning	Notation
Query	q , or q_i
A quantity z for query q_i	$z^{(i)}$
Number of training queries	n
Number of documents associated with query q	m
Number of document pairs associated with query q	$ ilde{m}$
Feature vector of a document associated with query q	x
Feature vectors of documents associated with query q	$\mathbf{x} = \{x_j\}_{j=1}^m$
Term frequency of query q in document d	$\mathrm{TF}(q,d)$
Inverse document frequency of query q	$\mathrm{IDF}(q)$
Length of document d	LEN(d)
Hypothesis	$h(\cdot)$
Scoring function	$f(\cdot)$
Loss function	$L(\cdot)$
Expected risk	$R(\cdot)$
Empirical risk	$\hat{R}(\cdot)$
Relevance degree for document x_i	l_j
Document x_u is more relevant than document x_v	$l_u \succ l_v$
Pairwise preference between documents x_u and x_v	$l_{u,v}$
Total order of document associated with the same query	π_l
Ground truth label for document x_j	y_j
Ground truth label for document pair (x_u, x_v)	$y_{u,v}$
Ground truth list for documents associated with query q	π_y
Ground truth permutation set for documents associated with query q	Ω_y
Original document index of the j-th element in permutation π	$\pi^{-1}(j)$
Rank position of document j in permutation π	$\pi(j)$
Number of classes	K
Index of class	k
VC dimension of a function class	V
Indicator function	$I_{\{\cdot\}}$
Gain function	$G(\cdot)$
Position discount function	$\eta(\cdot)$

we sometimes use L(h) and L(f) interchangeably to represent the loss function. When we need to emphasize the parameter in the scoring function, we will use f(w,x) instead of f(x) in the discussion, although they actually mean the same thing.

The Pointwise Approach

When using the technologies of ML to solve the problem of ranking, probably the most straightforward way is to check whether existing learning methods can be directly applied. This is exactly what the pointwise approach does. When doing so, one assumes that the exact relevance degree of each document is what we are going to predict, although this may not be necessary when the target is to produce a ranked list of the documents.

According to different ML technologies used, the pointwise approach can be further divided into three subcategories: regression based algorithms, classification based algorithms and ordinal regression based algorithms. For regression based algorithms, the output space contains real-valued relevance scores; for classification based algorithms, the output space contains non-ordered categories; and for ordinal regression based algorithms, the output space contains ordered categories. Documents together with their ground truth labels in the training set are regarded as i.i.d. random variables sampled from the product of the input and output spaces.

In the following, we will first introduce representative algorithms in the three subcategories of the pointwise approach, and then make discussions on their advantages and problems.

2.1 Regression based Algorithms

In this subcategory, the problem of ranking is reduced to a regression problem, by regarding the relevance degree as real numbers. Here we introduce two representative algorithms as examples.

2.1.1 Polynomial Regression Function

This is an early work on learning to rank [49], which uses least square regression to learn the scoring function.

Given a group of documents $\mathbf{x} = \{x_j\}_{j=1}^m$ associated with training query q, the ground truth label for x_j is defined as a vector. For binary judgments, $\vec{y}_j = (1,0)$ if the document is judged as relevant, and $\vec{y}_j = (0,1)$ otherwise. For multiple ordered categories, the k-th element of the vector \vec{y}_j is set to 1 and the other elements are set to 0, if the document is judged as belonging to the k-th category.

Then, the scoring function is defined as $\vec{f} = (f_1, f_2, ...)$, with each element f_k as a predictor of the k-th element in $\vec{y_j}$. Here, f_k is supposed to be selected from the polynomial function class, i.e.,

$$f_k(x_j) = w_{k,0} + w_{k,1} \cdot x_{j,1} + \dots + w_{k,T} \cdot x_{j,T} + w_{k,T+1} \cdot x_{j,1}^2 + w_{k,T+2} \cdot x_{j,1} \cdot x_{j,2} + \dots,$$
 (2.1)

where $w_{k,l}$ is the combination coefficient, $x_{j,l}$ is the l-th feature in the feature vector x_j , and T is the number of features in the representation of a document.

Next, the loss function is defined as the following square loss:

$$L(\vec{f}; x_j, \vec{y}_j) = ||\vec{y}_j - \vec{f}(x_j)||^2.$$
 (2.2)

Suppose we are given the binary judgment, then the loss function indicates that for a relevant document, only if the scoring function can exactly output (1,0), there will be zero loss. Otherwise, even if the output is (2,0), which seems to be an even stronger prediction of

relevance for this document, there will be some loss. This is, in some sense, not very reasonable.

2.1.2 Subset Ranking with Regression

Cossock and Zhang [33] have also solved the problem of ranking by reducing it to a regression problem.

Given $\mathbf{x} = \{x_j\}_{j=1}^m$, a group of documents associated with training query q, and the ground truth labels $\mathbf{y} = \{y_j\}_{j=1}^m$ of these documents in terms of multiple ordered categories, suppose scoring function f is used to rank these documents. Cossock and Zhang define the loss function as the following regression loss when performing learning to rank:

$$L(f;x_j,y_j) = (y_j - f(x_j))^2.$$
(2.3)

Furthermore, they have conducted some theoretical analysis on the use of such a loss function. The basic conclusion is that the regression loss can bound the ranking error (1–NDCG) (See Section 5 for more discussions).

However, it is clear that this work also suffers from the same problem as the polynomial regression function [49]. In many cases, it is not right to accurately predict the value of y_j since y_j is actually a qualitative judgment but not a quantitative value at all.

2.2 Classification based Algorithms

Analogously to reducing ranking to regression, one can also consider reducing ranking to a classification problem. Since the classification based algorithms do not regard the ground truth label as a quantitative value, it is more reasonable than the regression based algorithms. Here we introduce two representative algorithms in this subcategory.

2.2.1 Discriminative Model for IR

While most conventional ranking models for IR can be regarded as generative models (e.g., the language model for IR), ML literature has shown that discriminative models are preferred over generative models in many recent situations. Therefore, it is worth trying to find whether discriminative classifiers can lead to retrieval performances similar to or even better than retrieval performances of those generative IR ranking models. Actually, there has been some work that studied the use of a discriminative classification model for relevance ranking in IR, such as [31, 53, 90]. Here we take [90] as an example to illustrate the basic idea.

Given documents $\mathbf{x} = \{x_j\}_{j=1}^m$, and their binary relevance judgments $\mathbf{y} = \{y_j\}_{j=1}^m$ associated with query q, one regards all the relevant documents (i.e., $y_j = 1$) as positive examples while all the irrelevant documents (i.e., $y_j = 0$) as negative examples, and adopts the classification technology to learn the ranking model.

Two representative classification models, Maximum Entropy (ME) [54] and Support Vector Machines (SVM) [125, 126], were investigated in [90]. The principle of ME is to model all that is known and to assume nothing about the rest. So one can put all the information contained in the training data into a constraint set, and then maximize the entropy of the conditional probability with these constraints. SVM maximizes the margin on the constraint set of the training data. SVM has been proven to be one of the best classifiers in many classification tasks. It is also associated with a nice generalization theory based on the VC dimension, and therefore is theoretically guaranteed to have good performance even if the number of training samples is small.

Experiments on ad-hoc retrieval indicate that the ME-based algorithm is significantly worse than language models, but the SVM-based algorithm is comparable with and sometimes slightly better than language models. Based on this, the author argued that SVM is still preferred because of its ability to learn arbitrary features automatically, to make fewer assumptions, and to be more expressive [90].

2.2.2 Multi-class Classification for Ranking (McRank)

Li et al. [78] proposed using multi-class classification to solve the problem of ranking.

Given documents $\mathbf{x} = \{x_j\}_{j=1}^m$ associated with query q, and their relevance judgment $\mathbf{y} = \{y_j\}_{j=1}^m$, suppose we have a multi-class classifier, which makes prediction \hat{y}_j on x_j . Then the loss function used to

learn the classifier is defined as an upper bound of the following 0–1 classification error:

$$L(\hat{y}_j, y_j) = I_{\{y_j \neq \hat{y}_j\}}.$$
 (2.4)

In practice, different upper bounds yield different loss functions, such as the exponential loss, the hinge loss, and the logistic loss. All of them can be used to learn the classifier.

As for the testing process, the authors discussed how to convert classification results into ranking scores. In particular, the output of the classifier is converted to a probability using a logistic function, indicating the probability of a document belonging to a specific category. Suppose this probability is $P(\hat{y}_j = k), k = 0, ..., K-1$ (where K is the number of the categories given in the judgment). Then the following weighted combination is used to determine the final ranking scores of a document:

$$f(x_j) = \sum_{k=0}^{K-1} k \cdot P(\hat{y}_j = k). \tag{2.5}$$

2.3 Ordinal Regression based Algorithms

Ordinal regression¹ takes the ordinal relationship among the ground truth labels into consideration when learning the ranking model.

Suppose there are K ordered categories. The goal of ordinal regression is to find a scoring function, such that one can easily use thresholds $b_0 \leq b_2 \leq \cdots \leq b_{K-1}$ to distinguish the outputs of the scoring function into different ordered categories,² although this may not be necessary from the ranking point of view.

In the literature, there are several methods in this subcategory, such as [24, 25, 26, 34, 114]. We will introduce some of them as follows.

 $[\]overline{}$ Ordinal regression sometimes was also directly referred to as "ranking" in previous works [115].

² Note that there are some algorithms, such as [68], which were also referred to as ordinal regression based algorithms in the literature. According to our categorization, however, they belong to the pairwise approach since they do not really care about the accurate assignment of a document to one of the ordered categories. Instead, they focus more on the relative order between two documents.

2.3.1 Perceptron based Ranking (PRanking)

PRanking is a famous algorithm on ordinal regression [34]. The goal of PRanking is to find a direction defined by a parameter vector w, after projecting the documents onto which one can easily use thresholds to distinguish the documents into different ordered categories.

This goal is achieved by means of an iterative learning process. On iteration t, the learning algorithm gets an instance x_j associated with query q. Given x_j , the algorithm predicts $\hat{y}_j = \arg\min_k \{w^T x_j - b_k < 0\}$. It then receives the ground truth label y_j . If the algorithm makes a mistake by predicting the category of x_j as \hat{y}_j instead of y_j then there is at least one threshold, indexed by k, for which the value of $w^T x_j$ is on the wrong side of b_k . To correct the mistake, we need to move the values of $w^T x_j$ and b_k toward each other.

Let us see an example, shown in Figure 2.1. Suppose now we have model parameter w and document x_j . According to the output of the scoring function, this document seems to belong to the second category. However, its ground truth label indicates that it should belong to the fourth category. Then, the algorithm will lower down thresholds b_2 and b_3 . After that, the model parameter w is adjusted as $w = w + x_j$,

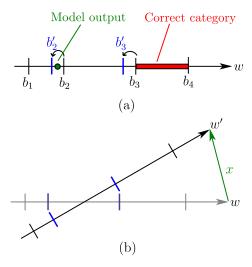


Fig. 2.1 Learning process of PRanking.

just as in many perceptron based algorithms. This process is repeated until the training process converges.

Harrington [59] later proposed using random sub-sampling to further improve the performance of PRanking. They first sub-sample the training data, and learn a PRanking model using each sample. After that, the weights and thresholds associated with the models are averaged to produce the final model. It has been proven that in this way a better generalization ability can be achieved [62].

2.3.2 Ranking with Large Margin Principles

Shashua and Levin [114] tried to use SVM to learn model parameter w and thresholds $b_k(k = 0, ..., K - 1)$, for ordinal regression.

Specifically, two strategies were proposed. The first one is referred to as the *fixed margin strategy*.

Given n training queries $\{q_i\}_{i=1}^n$, their associated documents $\mathbf{x}^{(i)} = \{x_j^{(i)}\}_{j=1}^{m^{(i)}}$, and the corresponding relevance judgments $\mathbf{y}^{(i)} = \{y_j^{(i)}\}_{j=1}^{m^{(i)}}$, the learning process is defined below, where the adoption of a linear scoring function is assumed. The constraints basically require every document to be correctly classified into its target ordered category, i.e., for documents in category k, $w^T x_j^{(i)}$ should exceed threshold b_{k-1} but be smaller than threshold b_k , with certain soft margins (i.e., $1 - \xi_{j,k-1}^{(i)*}$ and $1 - \xi_{j,k}^{(i)}$, respectively). The margin term $\frac{1}{2} \|w\|^2$ controls the complexity of model w.

$$\min \frac{1}{2} \|w\|^2 + C \sum_{i=1}^n \sum_{j=1}^{m^{(i)}} \sum_{k=0}^{K-2} \left(\xi_{j,k}^{(i)} + \xi_{j,k+1}^{(i)*} \right)$$
s.t. $w^T x_j^{(i)} - b_k \le -1 + \xi_{j,k}^{(i)}$, if $y_j^{(i)} = k$,
$$w^T x_j^{(i)} - b_k \ge 1 - \xi_{j,k+1}^{(i)*}$$
, if $y_j^{(i)} = k+1$,
$$\xi_{j,k}^{(i)} \ge 0$$
, $\xi_{j,k+1}^{(i)*} \ge 0$,
$$j = 1, \dots, m^{(i)}$$
, $i = 1, \dots, n$, $k = 0, \dots, K-2$. (2.6)

The second strategy is called the *sum of margins strategy*. In this strategy, some additional thresholds a_k are introduced, such that for

category k, b_{k-1} is its lower-bound threshold and a_k is its upper-bound threshold. Accordingly, the constraints become that for documents in category k, $w^T x_j^{(i)}$ should exceed threshold b_{k-1} but be smaller than threshold a_k , with certain soft margins (i.e., $1 - \xi_{j,k-1}^{(i)*}$ and $1 - \xi_{j,k}^{(i)}$, respectively). The corresponding learning process can be expressed as follows, from which we can see that the margin term $\sum_{k=0}^{K-1} (a_k - b_k)$ really has the meaning of "margin" (in Figure 2.2, $(b_k - a_k)$ is exactly the margin between category k+1 and category k):

$$\min \sum_{k=0}^{K-1} (a_k - b_k) + C \sum_{i=1}^{n} \sum_{j=1}^{m^{(i)}} \sum_{k=0}^{K-2} \left(\xi_{j,k}^{(i)} + \xi_{j,k+1}^{(i)*} \right)$$
s.t. $a_k \le b_k \le a_{k+1}$,
$$w^T x_j^{(i)} \le a_k + \xi_{j,k}^{(i)}, \quad \text{if } y_j^{(i)} = k,$$

$$w^T x_j^{(i)} \ge b_k - \xi_{j,k+1}^{(i)*}, \quad \text{if } y_j^{(i)} = k+1,$$

$$\|w\|^2 \le 1, \quad \xi_{j,k}^{(i)} \ge 0, \quad \xi_{j,k+1}^{(i)*} \ge 0,$$

$$j = 1, \dots, m^{(i)}, \quad i = 1, \dots, n, \quad k = 0, \dots, K-2. \tag{2.7}$$

Ideally in the above methods, one requires b_k $(k=0,\ldots,K-1)$ to be in an increasing order, i.e., $b_{k-1} \leq b_k$. However, in practice, since there are no clear constraints on the thresholds in the optimization problem, the learning process cannot always guarantee this. To tackle the problem, Chu and Keerthi [26] proposed adding explicit or implicit constraints on the thresholds to the optimization problem. The explicit constraint simply takes the form of $b_{k-1} \leq b_k$, while the implicit constraint uses redundant training examples to guarantee the ordinal relationship among thresholds.

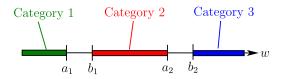


Fig. 2.2 Sum of margin strategy.

2.4 Discussions

In this subsection, we first discuss the relationship between the pointwise approach and some early learning methods in IR, such as relevance feedback. Then, we discuss the problems with the pointwise approach by considering the distinct properties of IR.

2.4.1 Relationship with Relevance Feedback

The pointwise approach of learning to rank, especially the classification based algorithms, has strong correlation with the relevance feedback algorithms [39, 112]. The relevance feedback algorithms, which have played an important role in IR literature, also leverage supervised learning technologies to improve the retrieval accuracy. The basic idea is to learn from explicit, implicit, or blind feedback to update the original query. Then the new query is used to retrieve a new list of documents. By doing this in an iterative manner, we can bring the original query closer to the optimal query so as to improve the retrieval performance.

Here we take the famous Rocchio algorithm [112] as an example to make discussions on the relationship between relevance feedback and learning to rank. The specific way that the Rocchio algorithm updates the query is as follows. First, both query q and its associated documents are represented in a vector space. Second, through relevance feedback, $\{x_j\}_{j=1}^{m^+}$ are identified as relevant documents (i.e., $y_j = 1$), and $\{x_j\}_{j=m^++1}^{m^+}$ are identified as irrelevant documents (i.e., $y_j = 0$). Third, the query vector is updated according to the following heuristic:

$$\tilde{q} = \alpha q + \beta \frac{1}{m^{+}} \sum_{j=1}^{m^{+}} x_{j} - \gamma \frac{1}{m^{-}} \sum_{j=m^{+}+1}^{m^{+}+m^{-}} x_{j}.$$
(2.8)

If we use the VSM model for retrieval, the documents will then be ranked according to their inner products with the new query vector \tilde{q} . Mathematically, we can define the corresponding ranking function as:

$$f(x_j) = \tilde{q}^T x_j. \tag{2.9}$$

In this sense, we can regard the query vector as the model parameter. For ease of discussion, we use w to represent this vector, i.e., $w = \tilde{q}$.

Then, as shown in [77], there is actually a hidden loss function behind the above query update process, which is a function of w and x. That is:

$$L(f, x_j, y_j) = \begin{cases} \frac{1}{m^+} \left(\frac{1-\alpha}{4} ||w||^2 - \beta w^T x_j \right), & y_j = 1, \\ \frac{1}{m^-} \left(\frac{1-\alpha}{4} ||w||^2 + \gamma w^T x_j \right), & y_j = 0. \end{cases}$$
(2.10)

In other words, the Rocchio algorithm also minimizes a certain pointwise loss function. In this sense, it looks quite similar to the pointwise approach of learning to rank. However, we would like to point out its significant differences from what we call learning to rank, as shown below.

- The feature space in the Rocchio algorithm is the standard vector space as used in VSM. In this space, both queries and documents are represented as vectors, and their inner product defines the relevance. In contrast, in learning to rank, the feature space contains features extracted from each querydocument pair. Only documents have feature representations, and the query is not a vector in the same feature space.
- The Rocchio algorithm learns the model parameter from the feedback on a given query, and then uses the model to rank the documents associated with the same query. It does not consider the generalization of the model across queries. However, in learning to rank, we learn the ranking model from a training set, and mainly use it to rank the documents associated with unseen test queries.
- The model parameter w in the Rocchio algorithm actually
 has its physical meaning, i.e., it is the updated query vector.
 However, in learning to rank, the model parameter does not
 have such a meaning and only corresponds to the importance
 of each feature to the ranking task.
- The goal of the Rocchio algorithm is to update the query formulation for a better retrieval but not to learn an optimal ranking function. In other words, after the query is updated, the fixed ranking function (e.g., the VSM model) is used to return a new list of related documents.

2.4.2 Problems with the Pointwise Approach

The algorithms belonging to the pointwise approach are highly correlated with each other. For example, when the number of categories K=2, the ordinal regression problem will naturally reduce to a binary classification problem. Therefore, these algorithms have similar problems when dealing with the task of learning to rank.

Since the input object in the pointwise approach is a single document, the relative order between documents cannot be naturally considered in the learning process, although ranking is more about predicting relative order than accurate relevance degree. Furthermore, the two intrinsic properties of the evaluation measures for ranking (i.e., query-level and position-based) cannot be well considered by the pointwise approach. First, the fact is ignored in the approach that some documents are associated with the same query and some others are not. As a result, when the number of associated documents varies largely for different queries,³ the overall loss function will be dominated by those queries with a large number of documents. Second, the position of each document in the ranked list is invisible to the pointwise loss functions. Therefore, the pointwise loss function may unconsciously overemphasize those unimportant documents (which are ranked low in the final ranked list and thus do not affect user experiences).

Given the above problems, the pointwise approach can only be a sub-optimal solution to ranking. To tackle the problem, people have made attempts at regarding a document pair, or the entire group of documents associated with the same query, as the input object. This results in the pairwise and listwise approaches of learning to rank. With the pairwise approach, the relative order among documents can be better modeled. With the listwise approach, the position information can be visible to the learning-to-rank process.

³ For the re-ranking scenario, the number of documents to rank for each query may be very similar, e.g., top 1000 documents per query. However, if we consider all the documents containing the query word, the difference between the number of documents for popular queries and that for tail queries may be very large.

The Pairwise Approach

The pairwise approach¹ does not focus on accurately predicting the relevance degree of each document; instead, it cares about the relative order between two documents. In this sense, it is closer to the concept of "ranking" than the pointwise approach.

In the pairwise approach, the ranking problem is reduced to a classification problem on document pairs. That is, the goal of learning is to minimize the number of miss-classified document pairs (i.e., the goal of the classification is to make positive predictions on those pairs whose first document is more relevant than the second document, and make negative predictions on other pairs). In the extreme case, if all the document pairs are correctly classified, all the documents will be correctly ranked. Note that this classification differs from the classification in the pointwise approach, since it operates on every two documents under investigation. Note that document pairs are not independent, which violates the basic assumption of classification. In this case, although one can still use classification algorithms to learn the ranking model, a different theoretical framework is needed to analyze the generalization of the learning process. We will make discussions on this in Section 7.

¹ Also referred to as preference learning in the literature.

In the rest of this section, we will introduce several representative algorithms that belong to the pairwise approach.

3.1 Example Algorithms

3.1.1 Ordering with Preference Function

In [29], a hypothesis $h(x_u, x_v)$ directly defined on a pair of documents is studied (i.e., without use of the scoring function f). In particular, given two documents x_u and x_v associated with a training query q, the loss function is defined below:

$$L(h; x_u, x_v, y_{u,v}) = \frac{|y_{u,v} - h(x_u, x_v)|}{2},$$
(3.1)

where the hypothesis is defined as $h(x_u, x_v) = \sum_t w_t h_t(x_u, x_v)$ and $h_t(x_u, x_v)$ is called the base preference function.

Suppose $h_t(x_u, x_v)$ only takes a value from $\{1, -1\}$, where a value of 1 indicates that document x_u is ranked before x_v , and a value of -1 indicates the opposite. Then, we can easily find if the ground truth label indicates that document x_u should be ranked before document x_v (i.e., $y_{u,v} = 1$) but if $h(x_u, x_v) = -1$, there will be a loss of one for this pair of documents. When all the pairs are incorrectly ranked, the average loss on the training set will reach its maximum value of one. On the other hand, when all the pairs are correctly ranked, we can get the minimum loss of zero.

With the above loss function, the weighted majority algorithm, e.g., the Hedge algorithm, is used to learn the parameters in hypothesis h. Note that h is actually a preference function, which cannot directly output the ranked list of the documents. In this case, an additional step is needed to convert the pairwise preference between any two documents to the total order of all the documents. To this end, one needs to find the ranked list π , which has the largest agreement with the pairwise preferences. This process is described below:

$$\max_{\pi} \sum_{u < v} h(x_{\pi^{-1}(u)}, x_{\pi^{-1}(v)}). \tag{3.2}$$

As we know, this is a typical problem called rank aggregation. It has been proven NP-hard to find the optimal solution to the above optimization problem. To tackle the challenge, a greedy ordering algorithm was proposed in [29], which can be much more efficient, and its agreement with the pairwise preferences is at least half the agreement of the optimal algorithm.

3.1.2 RankNet and FRank

RankNet [14] is probably the first learning-to-rank algorithm used by commercial search engines.²

In RankNet, the loss function is also defined on a pair of documents, but the hypothesis is defined with the use of a scoring function f. Given two documents x_u and x_v associated with training query q, a target probability $\bar{P}_{u,v}$ is constructed based on their ground truth labels. For example, we can define $\bar{P}_{u,v} = 1$, if $y_{u,v} = 1$; $\bar{P}_{u,v} = 0$, otherwise. Then, the modeled probability $P_{u,v}$ is defined based on the difference between the scores of these two documents given by the scoring function, i.e.,

$$P_{u,v}(f) = \frac{\exp(f(x_u) - f(x_v))}{1 + \exp(f(x_u) - f(x_v))}.$$
 (3.3)

Then the cross entropy between the target probability and the modeled probability is used as the loss function, which we refer to as the cross entropy loss for short.

$$L(f; x_u, x_v, y_{u,v}) = -\bar{P}_{u,v} \log P_{u,v}(f) - (1 - \bar{P}_{u,v}) \log (1 - P_{u,v}(f)).$$
(3.4)

A neural network is then used as the model and gradient descent as the optimization algorithm to learn scoring function f. In [84], a nested ranker is built on top of RankNet to further improve the retrieval performance.

Tsai et al. [122] pointed out some problems with the loss function used in RankNet. The curve of the cross entropy loss as a function of $f(x_u) - f(x_v)$ is plotted in Figure 3.1. From this figure, one can see that in some cases, the cross entropy loss has a non-zero minimum, indicating that there will always be some loss no matter what kind

 $^{^{2}}$ As far as we know, Microsoft Live Search (http://www.live.com/) is using the model trained with a variation of RankNet.

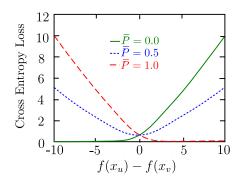


Fig. 3.1 Cross entropy loss as a function of $f(x_u) - f(x_v)$.

of model is used. This may not be in accordance with our intuition of a loss function. Furthermore, the loss is not bounded, which may lead to the dominance of some difficult document pairs in the training process.

To tackle these problems, a new loss function named the *fidelity loss* was proposed [122], which has the following form:

$$L(f; x_u, x_v, y_{u,v}) = 1 - \sqrt{\bar{P}_{u,v} P_{u,v}(f)} - \sqrt{(1 - \bar{P}_{u,v})(1 - P_{u,v}(f))}.$$
 (3.5)

The fidelity was originally used in quantum physics to measure the difference between two probabilistic states of a quantum. When being used to measure the difference between the target probability and the modeled probability, the fidelity loss has the shape as shown in Figure 3.2 as a function of $f(x_u) - f(x_v)$. By comparing the fidelity loss with the cross entropy loss, we can see that the fidelity loss is bounded between 0 and 1, and always has a zero minimum. These properties are nicer than those of the cross-entropy loss. On the other hand, however, while the cross-entropy loss is convex, the fidelity loss becomes non-convex. In theory, such a non-convex objective is more difficult to optimize. Overall, according to the experimental results reported in [122], FRank outperforms RankNet on several datasets.

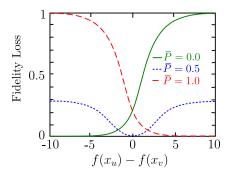


Fig. 3.2 Fidelity loss as a function of $f(x_u) - f(x_v)$.

3.1.3 RankBoost

The method of RankBoost [47] adopts AdaBoost [48] for the classification over document pairs. The only difference between RankBoost and AdaBoost is that the distribution in RankBoost is defined on document pairs while that in AdaBoost is defined on individual documents.

The algorithm flow of RankBoost is given in Algorithm 1, where \mathcal{D}_t is the distribution on document pairs, f_t is the weak ranker selected at the t-th iteration, and α_t is the weight for linearly combining the weak rankers. RankBoost actually minimizes the exponential loss defined below:

$$L(f; x_u, x_v, y_{u,v}) = \exp(-y_{u,v}(f(x_u) - f(x_v))).$$
(3.6)

Algorithm 1 Learning Algorithm for RankBoost

Input: document pairs

Given: initial distribution \mathcal{D}_1 on input document pairs.

For t = 1, ..., T

Train weak ranker f_t based on distribution \mathcal{D}_t .

Choose α_t

Update $\mathcal{D}_{t+1}(x_u^{(i)}, x_v^{(i)}) = \frac{1}{Z_t} \mathcal{D}_t(x_u^{(i)}, x_v^{(i)}) \exp(\alpha_t(f_t(x_u^{(i)}) - f_t(x_v^{(i)})))$ where $Z_t = \sum_{i=1}^n \sum_{u, v: y_{u,v}^{(i)} = 1} \mathcal{D}_t(x_u^{(i)}, x_v^{(i)}) \exp(\alpha_t(f_t(x_u^{(i)}) - f_t(x_v^{(i)}))).$

Output: $f(x) = \sum_t \alpha_t f_t(x)$.

From Algorithm 1, one can see that RankBoost learns the optimal weak ranker f_t and its coefficient α_t based on the current distribution of the document pairs (\mathcal{D}_t) . Three ways of choosing α_t are discussed in [47].

- First, most generally, for any given weak ranker f_t , it can be shown that Z_t , viewed as a function of α_t , has a unique minimum, which can be found numerically via a simple binary search.
- The second method is applicable in the special case that f_t takes a value from $\{0,1\}$. In this case, one can minimize Z_t analytically as follows. For $b \in \{-1,0,1\}$, let

$$W_{t,b} = \sum_{i=1}^{n} \sum_{u,v:y_{u,v}^{(i)}=1} \mathcal{D}_t \left(x_u^{(i)}, x_v^{(i)} \right) I_{\{f_t(x_u^{(i)}) - f_t(x_v^{(i)}) = b\}}. \tag{3.7}$$

Then

$$\alpha_t = \frac{1}{2} \log \left(\frac{W_{t,-1}}{W_{t,1}} \right). \tag{3.8}$$

• The third way is based on the approximation of Z_t , which is applicable when f_t takes a real value from [0, 1]. In this case, if we define:

$$r_{t} = \sum_{i=1}^{n} \sum_{u,v:u_{v,v}^{(i)} = 1} \mathcal{D}_{t}\left(x_{u}^{(i)}, x_{v}^{(i)}\right) \left(f_{t}\left(x_{u}^{(i)}\right) - f_{t}\left(x_{v}^{(i)}\right)\right), \quad (3.9)$$

then

$$\alpha_t = \frac{1}{2} \log \left(\frac{1 + r_t}{1 - r_t} \right). \tag{3.10}$$

Because of the analogy to AdaBoost, RankBoost inherits many nice properties from AdaBoost, such as the ability in feature selection, convergence in training, and certain generalization abilities.

3.1.4 Ranking SVM

Ranking SVM [63, 68] uses SVM for the task of pairwise classification. Given n training queries $\{q_i\}_{i=1}^n$, their associated document pairs

 $(x_u^{(i)}, x_v^{(i)})$, and the corresponding ground truth label $y_{u,v}^{(i)}$, the mathematical formulation of Ranking SVM is as shown below, where a linear scoring function is used, i.e., $f(x) = w^T x$,

$$\min \frac{1}{2} \|w\|^2 + C \sum_{i=1}^n \sum_{u,v:y_{u,v}^{(i)} = 1} \xi_{u,v}^{(i)}$$
s.t. $w^T (x_u^{(i)} - x_v^{(i)}) \ge 1 - \xi_{u,v}^{(i)}$, if $y_{u,v}^{(i)} = 1$,
$$\xi_{u,v}^{(i)} \ge 0, \quad i = 1, \dots, n.$$
(3.11)

As we can see, the objective function in Ranking SVM is exactly the same as in SVM, where the margin term $\frac{1}{2}||w||^2$ controls the complexity of model w. The difference between Ranking SVM and SVM lies in the constraints, which are constructed from document pairs. The loss function in Ranking SVM is a hinge loss defined on document pairs. For example, for a training query q, if document x_u is labeled as being more relevant than document x_v (mathematically, $y_{u,v} = +1$), then if $w^T x_u$ is larger than $w^T x_v$ by a margin of 1, there is no loss. Otherwise, the loss will be $\xi_{u,v}$.

Since Ranking SVM is well rooted in the framework of SVM, it inherits nice properties of SVM. For example, with the help of margin maximization, Ranking SVM can have good generalization ability. Kernel tricks can also be applied to Ranking SVM, so as to handle complex non-linear problems.

3.2 Discussions

3.2.1 Extension of the Pairwise Approach

Note that in the above algorithms, pairwise preference is used as the ground truth label. When we are given the relevance judgment in terms of multiple ordered categories, however, converting it to pairwise preference will lead to the absence of the information about the finer granularity in the relevance judgment.

To tackle the problem, Qin et al. [97] proposed a new algorithm named the multiple hyperplane ranker (MHR). The basic idea is "divide-and-conquer". Suppose there are K different categories of

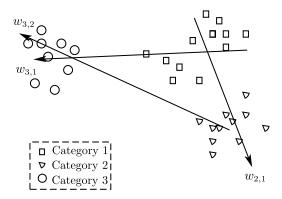


Fig. 3.3 Training multiple rankers.

judgments, then one can train K(K-1)/2 Ranking SVM models in total, with each model trained from the document pairs with two specific categories of judgments (see Figure 3.3). At the test phase, rank aggregation is used to merge the ranking results given by each model to produce the final ranking result. For instance, suppose that the model trained from categories k and l is denoted by $f_{k,l}$, then the final ranking results can be attained by using the weighted Borda Count aggregation:

$$f(x) = \sum_{k,l} \alpha_{k,l} f_{k,l}(x).$$
 (3.12)

Here the combination coefficient $\alpha_{k,l}$ can be pre-specified or learned from a separate validation set. The experimental results in [97] show that by considering more information about the judgment, the ranking performance can be significantly improved over Ranking SVM. Note that the technology used in MHR can actually be extended to any other pairwise ranking algorithm.

3.2.2 Improvement of the Pairwise Approach

It seems that the pairwise approach has its advantages as compared to the pointwise approach, since it can model the relative order between documents. However, in some cases, it faces even larger challenges than the pointwise approach. In Section 2.4, we have mentioned the problem of the pointwise approach when documents are distributed in an imbalanced manner across queries. Here this issue becomes even more serious in the pairwise approach. Considering that every two documents associated with the same query can create a document pair if their relevance degrees are different, in the worse case, the pair number can be quadratic to the document number. As a result, the difference in the numbers of document pairs is usually significantly larger than the difference in the numbers of documents. This phenomenon has been observed in some previous studies. For example, as reported in [99, 102], the distributions of pair numbers per query can be very skewed even if the document numbers of different queries are similar to each other (see Figure 3.4 for the distribution of a dataset from a commercial search engine), indicating that the above problem is really very serious in practice.

In this case, the pairwise loss function will be seriously dominated by the queries with large numbers of document pairs, and as a result the pairwise loss function will become inconsistent with the query-level IR evaluation measures. To tackle the problem, Cao et al. [16] and Qin et al. [99, 102] proposed introducing query-level normalization to the pairwise loss function. That is, the pairwise loss for a query will be normalized by the total number of document pairs associated with that query. In this way, the normalized pairwise losses with regards to different queries will become comparable with each other in their magnitude, no matter how many document pairs they are originally associated

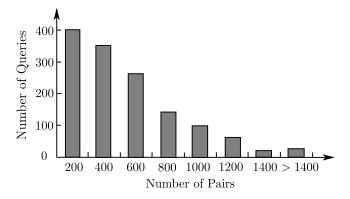


Fig. 3.4 Distribution of pair numbers per query.

with. With this kind of query-level normalization, Ranking SVM will become a new algorithm, which we call IR-SVM [16]. Specifically, given n training queries $\{q_i\}_{i=1}^n$, their associated document pairs $(x_u^{(i)}, x_v^{(i)})$, and the corresponding relevance judgment $y_{u,v}^{(i)}$, IR-SVM solves the following optimization problem:

$$\min \frac{1}{2} \|w\|^2 + C \sum_{i=1}^n \frac{\sum_{u,v:y_{u,v}^{(i)}=1} \xi_{u,v}^{(i)}}{\tilde{m}^{(i)}}$$
s.t. $w^T \left(x_u^{(i)} - x_v^{(i)} \right) \ge 1 - \xi_{u,v}^{(i)}, \text{ if } y_{u,v}^{(i)} = 1$

$$\xi_{u,v}^{(i)} \ge 0; \quad i = 1,\dots, n,$$
 (3.13)

where $\tilde{m}^{(i)}$ is the number of document pairs associated with query q_i . According to the experimental results in [16, 99, 102], a significant performance improvement has been observed after the query-level normalization is introduced.

The Listwise Approach

The listwise approach can be divided into two sub-categories. For the first sub-category, the output space contains the relevance degrees of all the documents associated with a query (i.e., \mathbf{y}), and the loss function is defined based on the approximation or bound of widely used IR evaluation measures. For the second sub-category, the output space contains the permutation of the documents associated with the same query (i.e., π_y), and the loss function measures the difference between the permutation given by the hypothesis and the ground truth permutation.

In the following, we will introduce both sub-categories and their representative algorithms.

4.1 Direct Optimization of IR Evaluation Measures

It might be the most straightforward choice to learn the ranking model by directly optimizing what is used to evaluate the ranking performance. This is exactly the motivation of the first sub-category of the listwise approach, which we call the direct optimization methods. However, the task is not as trivial as it seems. As we mentioned before, IR evaluation measures, such as NDCG and MAP, are position based, and thus non-continuous and non-differentiable [110, 132]. The difficulty in optimizing such objective functions stems from the fact that most existing optimization techniques were developed to handle continuous and differentiable cases.

To tackle the challenges, several attempts have been made. First, one can choose to optimize a continuous and differentiable approximation of the IR evaluation measure. By doing so, many existing optimization technologies can be leveraged. Example algorithms include SoftRank [119] and AppRank [98]. Second, one can alternatively optimize a continuous and differentiable (and sometimes even convex) bound of the IR evaluation measure. Example algorithms include SVM^{map} [136], SVM^{ndcg} [20], and PermuRank [132]. Actually, this trick has also been used in classification.¹ Third, one can choose to use optimization technologies that are able to optimize complex objectives. For example, one can leverage the Boosting framework for this purpose (the corresponding algorithm is called AdaRank [131]), or adopt the genetic algorithm for the optimization (the corresponding algorithm is called RankGP [134]).

In the rest of this subsection, we will take SoftRank, SVM^{map}, AdaRank, and RankGP as examples to introduce the direct optimization methods.

4.1.1 SoftRank

SoftRank [119] introduces randomness to the ranking scores of the documents, and then uses the expectation of NDCG as an approximation of the original IR evaluation measure NDCG.

First, SoftRank defines the score distribution. Given $\mathbf{x} = \{x_j\}_{j=1}^m$ associated with training query q, score s_j of document x_j is treated as no longer a deterministic value but a random variable. The random variable is governed by a Gaussian distribution whose variance is σ_s and mean is $f(x_j)$, the original score outputted by the scoring function. That is:

$$p(s_j) = N(s_j | f(x_j), \sigma_s^2).$$
 (4.1)

¹ Since the 0–1 classification loss is non-differentiable, convex upper bounds like the exponential loss have been used instead.

Second, SoftRank defines the rank distribution. Due to the randomness in the score, every document has the probability of being ranked at any position. Specifically, based on the score distribution, the probability of a document being ranked before another can be deduced as follows:

$$p_{u,v} = \int_0^\infty N(s | f(x_u) - f(x_v), 2\sigma_s^2) ds.$$
 (4.2)

On this basis, the rank distribution can be derived in an iterative manner. Let us consider adding the documents into the ranked list one after another. Suppose we already have document x_j in the ranked list, when adding document x_u , if document x_u can beat x_j the rank of x_j will be increased by one. Otherwise the rank of x_j will remain unchanged. Mathematically, the probability of x_j being ranked at position r (denoted as $p_j(r)$) can be computed as follows:

$$p_j^u(r) = p_j^{(u-1)}(r-1)p_{u,j} + p_j^{(u-1)}(r)(1-p_{u,j}).$$
(4.3)

Third, with the rank distribution, SoftRank computes the expectation of NDCG@m (where m is the total number of documents associated with the query) as the objective function for learning to rank (which we call SoftNDCG²). In other words, (1–SoftNDCG) corresponds to the loss function in SoftRank.

SoftNDCG
$$\triangleq \frac{1}{Z_m} \sum_{j=1}^m (2^{y_j} - 1) \sum_{r=0}^{m-1} \eta(r) p_j(r).$$
 (4.4)

In order to learn ranking model f by maximizing SoftNDCG, one can use a neural network as the model, and gradient descent as the optimization algorithm. In [55], the Gaussian process is used to further enhance SoftRank, where σ_s is no longer a pre-specified constant but a learned parameter.

$4.1.2 \quad \text{SVM}^{map}$

 ${\rm SVM}^{map}$ [136] uses the framework of structured SVM [70, 123] to optimize the IR evaluation measure AP.

 $^{^2\,\}mathrm{For}$ ease of reference, we also refer to the objective functions like SoftNDCG as the surrogate measure.

Suppose $\mathbf{x} = \{x_j\}_{j=1}^m$ represents all the documents associated with training query q, its corresponding ground truth label is $\mathbf{y} = \{y_j\}_{j=1}^m$ $(y_j = 1, \text{ if document } x_j \text{ is labeled as relevant; } y_j = 0, \text{ otherwise}), \text{ and any incorrect label of } \mathbf{x} \text{ is represented as } \mathbf{y}^c.$ Then SVM^{map} is formulated as follows, where AP is used in the constraints of structured SVM. It has been proven that the sum of slacks in SVM^{map} can bound (1 - AP) from above.

$$\min \frac{1}{2} ||w||^2 + \frac{C}{n} \sum_{i=1}^n \xi^{(i)}$$
s.t. $\forall \mathbf{y}^{c(i)} \neq \mathbf{y}^{(i)}$,
$$w^T \Psi(\mathbf{y}^{(i)}, \mathbf{x}^{(i)}) \ge w^T \Psi(\mathbf{y}^{c(i)}, \mathbf{x}^{(i)}) + 1 - \text{AP}(\mathbf{y}^{c(i)}) - \xi^{(i)}. \quad (4.5)$$

Here Ψ is called the joint feature map, whose definition is given as below:

$$\Psi(\mathbf{y}, \mathbf{x}) = \sum_{u, v: y_u = 1, y_v = 0} (x_u - x_v). \tag{4.6}$$

$$\Psi(\mathbf{y}^c, \mathbf{x}) = \sum_{u, v: y_u = 1, y_v = 0} (y_u^c - y_v^c)(x_u - x_v).$$
 (4.7)

As we know, there are an exponential number of incorrect labels for the documents, and thus the optimization problem has an exponential number of constraints for each query. Therefore, it is a big challenge to directly solve such an optimization problem. To tackle the challenge, a working set is maintained, which only contains those constraints with the largest violation (defined below), and the optimization is performed only with respect to this working set.

Violation
$$\triangleq 1 - AP(\mathbf{y}^c) + w^T \Psi(\mathbf{y}^c, \mathbf{x}).$$
 (4.8)

Then the problem is to efficiently find the most violated constraints for a given scoring function $f(x) = w^T x$. To this end, the property of AP is considered. That is, if the relevance at each position is fixed, AP will be the same no matter which document appears at that position. Furthermore, with the same AP, if the documents are sorted according to the descending order of their scores, $w^T \Psi(\mathbf{y}^c, \mathbf{x})$ will be maximized.

Therefore, an efficient strategy to find the most violated constraint can be designed [136], whose time complexity is $O(m \log m)$, where m is the number of documents associated with query q.

In [20, 21], the idea of SVM^{map} is further extended to optimize other IR evaluation measures, and the corresponding algorithms are named as SVM^{ndcg} and SVM^{mrr} . Basically, different feature maps or different strategies of searching the most violated constraints are used in these extensions, but the idea remains the same as that of SVM^{map} .

4.1.3 AdaRank

Xu and Li [131] found that IR evaluation measures can be plugged into the framework of Boosting and get effectively optimized. This process does not require IR evaluation measures to be continuous and differentiable. The resultant algorithm is called AdaRank.

As we know, in conventional AdaBoost the exponential loss is used to update the distribution of input objects and to determine the combination coefficient α_t at each round of iteration. Analogously, in AdaRank, IR evaluation measures are used to update the distribution of queries and to compute the combination coefficient. The algorithm flow is shown below, where $M(f, \mathbf{x}, \mathbf{y})$ represents the IR evaluation measure.

Due to the analogy to AdaBoost, AdaRank can focus on hard queries. Furthermore, a condition for the convergence of the training

Algorithm 2 Learning Algorithms for AdaRank

Input: document group for each query

Given: initial distribution \mathcal{D}_1 on input queries

For t = 1, ..., T

Train weak ranker $f_t(\cdot)$ based on distribution \mathcal{D}_t .

Choose
$$\alpha_t = \frac{1}{2} \log \frac{\sum_{i=1}^n \mathcal{D}_t(i)(1+M(f_t,\mathbf{x}^{(i)},\mathbf{y}^{(i)}))}{\sum_{i=1}^n \mathcal{D}_t(i)(1-M(f_t,\mathbf{x}^{(i)},\mathbf{y}^{(i)}))}$$

Train weak ranker
$$f_t(\cdot)$$
 based on distribution \mathcal{D}_t .
Choose $\alpha_t = \frac{1}{2} \log \frac{\sum_{i=1}^n \mathcal{D}_t(i)(1+M(f_t,\mathbf{x}^{(i)},\mathbf{y}^{(i)}))}{\sum_{i=1}^n \mathcal{D}_t(i)(1-M(f_t,\mathbf{x}^{(i)},\mathbf{y}^{(i)}))}$
Update $\mathcal{D}_{t+1}(i) = \frac{\exp(-M(\sum_{s=1}^t \alpha_s f_s,\mathbf{x}^{(i)},\mathbf{y}^{(i)}))}{\sum_{j=1}^n \exp(-M(\sum_{s=1}^t \alpha_s f_s,\mathbf{x}^{(j)},\mathbf{y}^{(j)}))}$,

Output: $\sum_t \alpha_t f_t(\cdot)$.

process was given in [131], with a similar derivation technique to that for AdaBoost. The condition requires $|M(\sum_{s=1}^t \alpha_s f_s, \mathbf{x}, \mathbf{y}) - M(\sum_{s=1}^{t-1} \alpha_s f_s, \mathbf{x}, \mathbf{y}) - \alpha_t M(f_t, \mathbf{x}, \mathbf{y})|$ to be very small. This implies the assumption on the linearity of IR evaluation measure M, as a function of f_t . However, this assumption may not be well satisfied in practice. Therefore, it is possible that the training process of AdaRank does not naturally converge and some additional stopping criteria are needed.

4.1.4 Genetic Programming based Algorithms

There are some methods originally designed for optimizing complex objectives. Genetic programming is just one such method. In the literature of learning to rank, there have been several attempts on using genetic programming to optimize IR evaluation measures. Representative algorithms include [4, 41, 42, 43, 44, 45, 46, 121, 134].

Here we take the algorithm named RankGP [134] as an example to illustrate how genetic programming can be used to learn the ranking model. In this algorithm, the ranking model is defined as a tree, whose leaf nodes are features or constants, and non-leaf nodes are operators such as $+,-,\times,\div$ (see Figure 4.1). Then single population genetic programming is used to perform learning on the tree. Crossover, mutation, reproduction, and tournament selection are used as evolution mechanisms, and the IR evaluation measure is used as the fitness function.

In addition to the examples introduced above, there are also some other works [13, 86, 110] that directly optimize IR evaluation measures. Due to space restrictions, we will not introduce them in detail.

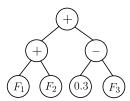


Fig. 4.1 Ranking function used in RankGP.

4.2 Minimization of Listwise Ranking Losses

In the second sub-category of the listwise approach, the loss function measures the inconsistency between the output of the ranking model and the ground truth permutation π_y . Although IR evaluation measures are not directly optimized here, if one can consider the distinct properties of ranking in IR in the design of the loss function, it is also possible that the model learned can have good performance in terms of IR evaluation measures. We refer to these algorithms as "algorithms that minimize listwise ranking losses". In this subsection, we will introduce two representative algorithms of them, i.e., ListNet [17] and ListMLE [129].

4.2.1 ListNet

In [17], a listwise ranking loss is proposed, which is based on the probability distribution on permutations.

Actually the distributions on permutations have been well studied in the field of probability theory. Many famous models have been proposed to represent permutation probability distributions, such as the Luce model [81, 95] and the Mallows model [82]. Since a permutation has a natural one-to-one correspondence with a ranked list, these researches can be naturally applied to ranking. ListNet [17] is just such an example, demonstrating how to apply the Luce model to learning to rank.

Given the relevance scores of the documents outputted by scoring function f (i.e., $\mathbf{s} = \{s_j\}_{j=1}^m$, where $s_j = f(x_j)$), the Luce model defines a probability for each possible permutation π of the documents, based on the chain rule, as follows:

$$P(\pi \mid \mathbf{s}) = \prod_{j=1}^{m} \frac{\varphi(s_{\pi^{-1}(j)})}{\sum_{u=j}^{m} \varphi(s_{-1}_{\pi(u)})},$$
(4.9)

where $\pi^{-1}(j)$ denotes the document ranked at the j-th position of permutation π , φ is a transformation function, which can be linear, exponential, or sigmoid. Each item $\frac{\varphi(s_{\pi^{-1}(j)})}{\sum_{u=j}^{m} \varphi(s_{\pi^{-1}(u)})}$ is a conditional probability as shown in the following example.

Suppose we have in total three documents A, B, and C associated with query q. The probability of permutation $\pi = (A, B, C)$ is equal to the product of the following three probabilities (i.e., $P_{\pi} = P_1 P_2 P_3$).

• P_1 : the probability of document A being ranked at the top position in π . This probability is computed by comparing the score of A with those of all the other documents.

$$P_1 = \frac{\varphi(s_A)}{\varphi(s_A) + \varphi(s_B) + \varphi(s_C)}. (4.10)$$

• P_2 : the conditional probability of document B being ranked at the second position given that A has already been ranked at the first position. This probability is computed by comparing the score of B with those of the other documents except A. In this simple example, there is only document C to be compared with.

$$P_2 = \frac{\varphi(s_B)}{\varphi(s_B) + \varphi(s_C)}. (4.11)$$

• P_3 : the conditional probability of document C being ranked on the third position given that documents A and B have already been ranked in the top two positions respectively. In our simple case, it is easy to see $P_3 = 1$.

With the Luce model, for a given query q, ListNet first defines the permutation probability distribution based on the scores given by scoring function f. Then it defines another permutation probability distribution $P_y(\pi)$ based on the ground truth label.³ For the next step, ListNet uses the K-L divergence between these two distributions to define its listwise ranking loss (which we call the K-L divergence loss for short).

$$L(f; \mathbf{x}, \pi_y) = D(P(\pi \mid \varphi(f(w, \mathbf{x}))) \parallel P_y(\pi)). \tag{4.12}$$

³ For example, if the ground truth is given in terms of relevance degree, it can be directly substituted into the Luce model to define a probability distribution. If the ground truth label is given as a ranked list, one can simply define the probability distribution as a delta function, or use a mapping function to map it to real-valued scores of the documents and then apply the Luce model. One can also use other ways such as the Mallows model to define the permutation probability distribution of the ground truth label.

A neural network model is employed in ListNet, and the gradient descent approach is used to minimize the K-L divergence loss. As shown in [17], the training curve of ListNet well demonstrates the correlation between the K-L divergence loss and 1 - NDCG@5.

As one may have noticed, there is a computational issue with ListNet. Due to the use of the scoring function, the testing complexity of ListNet can be the same as those of the pointwise and pairwise approaches, however the training complexity of ListNet is much higher. The training complexity of ListNet is in the exponential order of m (and thus intractable in practice), because the evaluation of the K-L divergence loss for each query q requires the addition of m-factorial terms. Comparatively speaking, the training complexities of the pointwise and pairwise approaches are roughly proportional to the number of documents (i.e., O(m)) and the number of document pairs (i.e., $O(\tilde{m})$). To tackle the problem, a top-k version of the K-L divergence loss is further introduced in [17], which is based on the top-k Luce model and can reduce the training complexity from m-factorial to the polynomial order of m.

4.2.2 ListMLE

Even if the top-k K-L divergence loss is used in ListNet, one still cannot avoid its following limitation. When k is set to be large, the time complexity of evaluating the K-L divergence loss is still very high. However, when k is set to be small, information about the permutation will be significantly lost and the effectiveness of the ListNet algorithm is questionable [17].

To tackle these problems, a new algorithm named ListMLE was proposed [129]. ListMLE is also based on the Luce model. For each query q, with the permutation probability distribution defined with the output of the scoring function, it uses the negative log likelihood of the ground truth permutation as the listwise ranking loss. We denote this new listwise ranking loss as the *likelihood loss* for short.

$$L(f; \mathbf{x}, \pi_y) = -\log P(\pi_y | \varphi(f(w, \mathbf{x}))). \tag{4.13}$$

It is clear that in this way the training complexity can be greatly reduced as compared to ListNet, since one only needs to compute the probability of a single permutation π_y but not all the permutations. Once again, one can use a neural network model to optimize the likelihood loss.

It should be noted that ListMLE has assumed the ground truth labels to be given as the total order of the documents. This is an advantage when it is really the case, however, if the judgments are given in other terms, ListMLE will not work. To tackle this challenge and further enhance the application scope of ListMLE, one needs to conduct the following preprocessing of the training data.

Given the judgment in terms of relevance degree, one can define an equivalent permutation set as follows:

$$\Omega_y = \left\{ \pi_y \, | \, u < v, \text{ if } l_{\pi_y^{-1}(u)} \succ l_{\pi_y^{-1}(v)} \right\}.$$

Similarly, given the judgment in terms of pairwise preferences, one can define Ω_y as below:

$$\Omega_y = \left\{ \pi_y \, | \, u < v, \text{ if } l_{\pi_y^{-1}(u), \pi_y^{-1}(v)} = 1 \right\}.$$

As compared to the judgment in terms of total order, we can regard the above judgments as incomplete. In other words, they are the necessary conditions of being ground truth permutations. However, permutations satisfying these constraints might not always be the ground truth permutations. This situation is very similar to that in multi-instance learning. Therefore, one can define the loss function, following a similar idea proposed for multi-instance learning [6], as below:

$$L(f; \mathbf{x}, \Omega_y) = \min_{\pi_y \in \Omega_y} \left(-\log P(\pi_y | \varphi(f(w, \mathbf{x}))) \right). \tag{4.14}$$

In [129], some analyses have been made on ListNet and ListMLE. Basically, it has been proven that both the K-L divergence loss and the likelihood loss are continuous, differentiable, and convex. In this way, they can be easily optimized, e.g., using the gradient descent algorithm.

4.3 Discussions

As shown in this subsection, different kinds of listwise ranking algorithms have been proposed. Intuitively speaking, they model the ranking problem in a more natural way than the pointwise and pairwise approaches, and thus can address some problems that these two

approaches have encountered. As we have discussed in the previous sections, for the pointwise and pairwise approaches, the position information is invisible to their loss functions, and they ignore the fact that some documents (or document pairs) are associated with the same query. Comparatively speaking, the listwise approach takes all the documents associated with the same query as the input and their ranked list (or their relevance degrees) as the output. In this way, it has the potential to distinguish documents from different queries, and to consider the position information in the output ranked list in its learning process. According to some previous studies, the performances of the listwise ranking algorithms are really better than previous approaches [129]. This is also verified by the discussions in Section 6, which is about the empirical ranking performances of different learning-to-rank algorithms, with the LETOR benchmark dataset as the experimental platform.

On the other hand, the listwise approach also has certain aspects that could be improved. For example, the training complexities of some listwise ranking algorithms (e.g., ListNet) are high since the evaluation of their loss functions are permutation based. A more efficient learning algorithm is needed to make the listwise approach more practical. Moreover, we would like to point out that the use of the position information in some listwise ranking algorithms is insufficient. For example, there is no explicit position discount considered in the loss functions of ListNet and ListMLE. As a result, even if the algorithms can see different positions in the output ranked list, they have not fully utilized them in the learning process. By introducing certain position discount factors, the performance improvement of these algorithms can be expected.

Analysis of the Approaches

In the previous three sections, we have introduced the pointwise, pairwise, and listwise approaches to learning to rank. The major differences between these approaches are the loss functions. Note that the loss functions are mainly used to guide the learning process, while the evaluation of the learned ranking model is based on IR evaluation measures. Therefore, an important issue to discuss is the relationship between the loss functions used in these approaches and the IR evaluation measure. This is exactly the motivation of this section. Without the loss of generality, we will take NDCG@m as an example in the discussions. Here m is the total number of documents associated with the query. For simplicity, we refer to NDCG@m as NDCG.

For ease of our discussion, we assume that the judgment is given in terms of multiple ordered categories, and the ground truth label is represented by a permutation set Ω_y as defined in Equation (4.14) (in other cases, one can obtain similar results). With this assumption, NDCG with respect to a given ranked list π can be defined

as follows¹:

$$NDCG(\pi, \Omega_y) = \frac{1}{Z_m} \sum_{t=1}^m G(\pi_y^{-1}(t)) \eta(\pi(\pi_y^{-1}(t))), \quad \forall \pi_y \in \Omega_y.$$
 (5.1)

It is easy to verify that for $\forall \pi_y \in \Omega_y$, the right-hand side of the equation takes the same value. Sometimes, we need to emphasize the ranking model, i.e., when $\pi = \operatorname{sort} \circ f$. In this case, we will denote the above-defined NDCG as NDCG $(f, \mathbf{x}, \Omega_y)$.

5.1 The Pointwise Approach

As mentioned in Section 2, Cossock and Zhang [33] have established the theoretical foundation for reducing ranking to regression.² Given $\mathbf{x} = \{x_j\}_{j=1}^m$, a group of documents associated with training query q, and the ground truth $\mathbf{y} = \{y_j\}_{j=1}^m$ of these documents in terms of multiple ordered categories, suppose a scoring function f is used to rank these documents. The authors proved a theory showing that the ranking error in terms of NDCG can be bounded by the following regression loss:

1 - NDCG
$$(f, \mathbf{x}, \Omega_y) \le \frac{1}{Z_m} \left(2 \sum_{j=1}^m \eta(j)^2 \right)^{\frac{1}{2}} \left(\sum_{j=1}^m (f(x_j) - y_j)^2 \right)^{\frac{1}{2}},$$
 (5.2)

where Z_m is the maximum DCG value and $\eta(j)$ is the discount factor used in NDCG.³

In other words, if one can really minimize the regression loss to zero, one can also minimize (1 - NDCG) to zero. This seems to be a very nice property of the regression based methods.

With similar proof techniques to those used in [33], Li et al. [78] showed that (1 - NDCG) can also be bounded by the multi-class

¹ Note that this definition of NDCG is equivalent to that given in Equation (1.9), although the index of the summation changes from the rank position in π (i.e., r) to the position in the ground truth list (i.e., t). Furthermore, we assume that the gain function $G(\cdot)$ can access the original relevance degree of the document.

² Note that the bounds given in the original papers are with respect to DCG, and here we give their equivalent form in terms of NDCG for ease of comparison.

³ Note that the original bound was given with regards to DCG. Here we use its equivalent form in terms of NDCG for ease of comparison.

classification loss as shown below (it is assumed K = 5 in the inequality).

 $1 - \text{NDCG}(f, \mathbf{x}, \Omega_y)$

$$\leq \frac{15}{Z_m} \sqrt{2 \left(\sum_{j=1}^m \eta(j)^2 - m \prod_{j=1}^m \eta(j)^{\frac{2}{m}} \right)} \cdot \sqrt{\sum_{j=1}^m I_{\{y_j \neq \hat{y}_j\}}}, \quad (5.3)$$

where \hat{y}_j is the prediction on the label of x_j by the multi-class classifier, and $f(x_j) = \sum_{k=0}^{K-1} k \cdot P(\hat{y}_j = k)$.

In other words, if one can really minimize the classification loss to zero, one can also minimize (1 - NDCG) to zero at the same time.

However, on the other hand, please note that when (1 - NDCG) is zero (i.e., the documents are perfectly ranked), the regression loss and the classification loss might not be zero (and can still be very large). In other words, the minimization of the regression loss and the classification loss is only a sufficient condition but not a necessary condition for optimal ranking in terms of NDCG.

Let us have a close look at the classification bound in inequality (5.3) with an example.⁴ Note that a similar example has been given in [1] to show the problem of reducing ranking to a binary classification problem.

Suppose for a particular query q, we have four documents (i.e., m=4) in total, and their ground truth labels are 4, 3, 2, and 1, respectively (i.e., $y_1=4$, $y_2=3$, $y_3=2$, $y_4=1$). We use the same discount factor and gain function as used in [78]. Then it is easy to compute that $Z_m = \sum_{j=1}^4 \frac{1}{\log(j+1)} (2^{y_j-1}) \approx 21.35$.

Then, suppose the outputs of the multi-class classifier are $\hat{y}_1=3$,

Then, suppose the outputs of the multi-class classifier are $\hat{y}_1 = 3$, $\hat{y}_2 = 2$, $\hat{y}_3 = 1$, and $\hat{y}_4 = 0$, with 100% confidence in the prediction for each document. It is easy to compute that $1 - \text{NDCG}(f, \mathbf{x}, \Omega_y)$ is 0 and we actually get a perfect ranking based on the classifier. However, in terms of multi-class classification, we made errors in all the four documents, i.e., $\sum_{j=1}^m I_{\{y_j \neq \hat{y}_j\}} = 4$. Furthermore, if we compute the bound

⁴One can get similar results for the regression bound given in inequality (5.2).

given by inequality (5.3), we obtain:

$$\frac{15}{Z_m} \sqrt{2 \left(\sum_{j=1}^m \left(\frac{1}{\log(j+1)} \right)^2 - m \prod_{j=1}^m \left(\frac{1}{\log(j+1)} \right)^{\frac{2}{m}} \right) \cdot \sum_{j=1}^m I_{\{y_j \neq \hat{y}_j\}}}$$

$$\approx \frac{24.49}{21.35} = 1.15.$$

It is clear the bound is meaningless since it is even larger than one. Actually the loose bound is not difficult to understand. The left-hand side of inequality (5.3) contains the position information, while the right-hand side does not. When the same amount of classification loss occurs in different positions, the ranking error will be quite different. In order to make the inequality always hold, the price one has to pay is that the bound must be very loose.

5.2 The Pairwise Approach

It has been shown in [22] that many of the pairwise loss functions are upper bounds of a quantity, named the essential loss for ranking. Furthermore, the essential loss is an upper bound of (1 - NDCG), and therefore these loss functions are also upper bounds of (1 - NDCG).

To better illustrate this result, we first introduce the concept of the essential loss, which is constructed by modeling ranking as a sequence of classification tasks.

Given a group of documents \mathbf{x} and their ground truth permutation $\pi_y \in \Omega_y$, the ranking problem can be decomposed into several sequential steps. For each step t, one tries to distinguish $\pi_y^{-1}(t)$, the document ranked at the t-th position in π_y , from all the documents ranked below the t-th position in π_y , using a scoring function f. Denote $\mathbf{x}_{(t)} = \{x_{\pi_y^{-1}(t)}, \dots, x_{\pi_y^{-1}(m)}\}$. One can define a classifier based on f, whose target output is $\pi_y^{-1}(t)$:

$$T \circ f(\mathbf{x}_{(t)}) = \arg \max_{j=\pi_y^{-1}(t),\dots,\pi_y^{-1}(m)} f(x_j).$$
 (5.4)

It is clear that there are m-t possible outputs of this classifier, i.e., $\{\pi_y^{-1}(t), \dots, \pi_y^{-1}(m)\}$. The 0–1 loss for this classification task can be

written as follows, where the second equation is based on the definition of $T \circ f$:

$$L_{t}(f; \mathbf{x}(t), \pi_{y}^{-1}(t)) = I_{\{T \circ f(\mathbf{x}_{(t)}) \neq \pi_{y}^{-1}(t)\}}$$

$$= 1 - \prod_{j=t+1}^{m} I_{\{f(\pi_{y}^{-1}(t)) > f(\pi_{y}^{-1}(j))\}}.$$
(5.5)

By summing up the losses at all the steps (t = 1, ..., m - 1), one can obtain,

$$\tilde{L}(f; \mathbf{x}, \pi_y) = \sum_{t=1}^{m-1} I_{\{T \circ f(\mathbf{x}_{(t)}) \neq \pi_y^{-1}(t)\}}.$$
(5.6)

By further taking a minimization over the permutation set Ω_y , we will get the so-called essential loss:

$$\tilde{L}(f; \mathbf{x}, \Omega_y) = \min_{\pi_y \in \Omega_y} \sum_{t=1}^{m-1} I_{\{T \circ f(\mathbf{x}_{(t)}) \neq \pi_y^{-1}(t)\}}.$$
 (5.7)

It has been proven in [22] that the essential loss is an upper bound of (1 - NDCG). As a result, the minimization of it will lead to the effective maximization of NDCG:

$$1 - \text{NDCG}(f, \mathbf{x}, \Omega_y) \le \frac{2^{K-1} - 1}{Z_m} \left(\sum_{t=1}^{m-1} \eta(t)^{\alpha} \right)^{\frac{1}{\alpha}} (\tilde{L}(f; \mathbf{x}, \Omega_y))^{\frac{1}{\beta}}, \quad (5.8)$$

where $\frac{1}{\alpha} + \frac{1}{\beta} = 1$.

As compared to the bounds given in the previous subsection, one can see that the essential loss has a nicer property. When (1 - NDCG) is zero, the essential loss is also zero. In other words, the zero value of the essential loss is not only a sufficient condition but also a necessary condition of the zero value of (1 - NDCG).

Furthermore, it has been proven in [22] that the essential loss has the following property:

$$\tilde{L}(f; \mathbf{x}, \Omega_y) \le \max_{\pi_y \in \Omega_y} a(T \circ f(\mathbf{x}_{(t)}), \pi_y^{-1}(t)), \tag{5.9}$$

where $a(\cdot,\cdot)$ is a cost sensitive function, i.e., a(i,j) = 0 if $l_i = l_j$ and a(i,j) = 1 otherwise.

Based on the above property, one can prove that the widely used pairwise loss functions are upper bounds of the essential loss.

$$\tilde{L}(f; \mathbf{x}, \Omega_y) \leq \sum_{t=1}^{m-1} \sum_{\substack{j=t+1, \\ l_{\pi_y^{-1}(t)} \neq l_{\pi_y^{-1}(j)}}}^{n} \phi\left(f(x_{\pi_y^{-1}(t)}) - f(x_{\pi_y^{-1}(j)})\right), \\
\forall \pi_y \in \Omega_y, \quad (5.10)$$

where when function ϕ is the hinge function, the exponential function, and the logistic function, the right hand side of the inequality represents exactly the loss functions of Ranking SVM, RankBoost, and RankNet.

Therefore, the minimization of the loss functions in the aforementioned pairwise ranking algorithms will all lead to the minimization of the essential loss. Further, considering the relationship between the essential loss and (1 - NDCG), these algorithms can also effectively minimize (1 - NDCG).

5.3 The Listwise Approach

5.3.1 Listwise Ranking Loss

One sub-category of the listwise approach minimizes a listwise ranking loss in the training process. Here, we take ListMLE as an example to perform the discussion on this sub-category. Actually, the technique used in the discussions on the pairwise loss functions can be used again.

Specifically, based on the essential loss, one can prove that the following inequality holds [22]:

$$\tilde{L}(f; \mathbf{x}, \Omega_y) \le \frac{1}{\log 2} \cdot \min_{\pi_y \in \Omega_y} \sum_{t=1}^{m-1} \left(f(x_{\pi_y^{-1}(t)}) + \log \left(\sum_{j=t}^m \exp\left(f(x_{\pi_y^{-1}(j)}) \right) \right) \right).$$

It is clear that the right hand side of the inequality contains the likelihood loss used in ListMLE (see Equation (4.14) when the exponential function is used as the transformation function φ). Recalling the connection between the essential loss and (1–NDCG) as discussed in the

previous section, the likelihood loss can also upper bound (1–NDCG). Therefore, the minimization of the likelihood loss in the training process will lead to the minimization of (1–NDCG) [22].

5.3.2 Loss Functions in Direct Optimization Methods

The other sub-category of the listwise approach optimizes a loss function derived from the IR evaluation measure. It seems that the discussion on the relationship between such a loss function and the corresponding IR evaluation measure is more straightforward, since they have natural connections. However, in order to make a formal discussion on the issue, a new quantity named "directness" needs to be introduced [61]. Basically, directness indicates whether a surrogate measure is a good approximation of the corresponding IR evaluation measure. Note that in the following definition, we assume the use of a linear scoring function, i.e., $f(x) = w^T \mathbf{x}$.

Definition 5.1. (Directness) For any query q, suppose its associated documents and the ground truth are \mathbf{x} and Ω_y , respectively. For a ranking model w, denote $\tilde{M}(w,\mathbf{x},\Omega_y)$ and $M(w,\mathbf{x},\Omega_y)$ as a surrogate measure and its corresponding IR evaluation measure, respectively. The directness of \tilde{M} with respect to M is defined as:

$$D(w, \tilde{M}, M) = \frac{1}{\sup_{\mathbf{x}, \Omega_y} |M(w, \mathbf{x}, \Omega_y) - \tilde{M}(w, \mathbf{x}, \Omega_y)|}.$$
 (5.11)

As can be seen, the directness is determined by the maximum difference between the surrogate measure and its corresponding IR evaluation measure with respect to a ranking model w, over the entire input and output spaces. In the extreme case, when the difference becomes zero, the directness will become infinite, and the surrogate measure will become exactly the IR evaluation measure.

As examples, the directness of SoftRank and SVM^{map} have been analyzed in [61]. The corresponding result for SoftRank is listed in the following theorem.

Theorem 5.1. For query q, suppose its associated documents and ground truth labels are \mathbf{x} and Ω_y , respectively. Assume $\forall i$ and j, $|f(x_i) - f(x_j)| \ge \delta > 0$ and $\forall q, m \le M$. If $\sigma_s < \frac{\delta}{2\mathrm{erf}^{-1}\left(\sqrt{\frac{5M-9}{5M-5}}\right)}$, then:

$$D(w, \text{SoftNDCG}, \text{NDCG}) \ge \frac{1}{M \cdot 2^{K-1} \cdot (\varepsilon_1 + \varepsilon_2)},$$
 (5.12)

where

$$\varepsilon_{1} = \frac{(M-1)\sigma_{s}}{2\delta\sqrt{\pi}}e^{-\frac{\delta^{2}}{4\sigma_{s}^{2}}}, \quad \varepsilon_{2} = \sqrt{\frac{\varepsilon_{3}(\sigma_{s})}{1-5\varepsilon_{3}(\sigma_{s})}} + 5\varepsilon_{3}(\sigma_{s}),$$

$$\varepsilon_{3} = \frac{M-1}{4}\left[1 - \operatorname{erf}^{2}\left(\frac{\delta}{2\sigma_{s}}\right)\right], \quad \operatorname{erf}(x) = \frac{2}{\sqrt{\pi}}\int_{0}^{x} e^{-t^{2}}dt,$$

and K is the number of relevance degrees in the judgment.

From the above theorem, we can see that the lower bound of the directness of SoftNDCG is a decreasing function of σ_s . Furthermore, when $\sigma_s \to 0$, the directness of SoftNDCG becomes extremely large. The intuitive explanation of this result is as follows. When the variance of the score distribution is small enough,⁵ it is highly probable that the introduction of the score distribution will not change the original order of any two documents. Therefore, the value of SoftNDCG can be very close to that of NDCG, even if it has become continuous and differentiable.

Similar analysis on SVM^{ndcg} [20], however, shows that for any ranking model, there always exists such inputs and outputs that will result in the large difference between its surrogate measure and the corresponding IR evaluation measure [61]. Consequently, it is not guaranteed that these algorithms can lead to the effective optimization of the IR evaluation measures.⁶

⁵ Since σ_s is a pre-defined parameter in the SoftRank algorithm, we are able to make SoftRank as direct as desired by setting σ_s as small as possible regardless of the joint probability distribution of the inputs and outputs.

⁶ Note that the same conclusion also applies to SVM^{map} and PermuRank [132].

5.4 Discussions

In this subsection, we have reviewed the relationships between different loss functions in learning to rank and the IR evaluation measure. The discussions well explain why different learning-to-rank algorithms perform reasonably well in practice (see the Experimental Results in Section 6).

While the analyses introduced in this section look quite nice, there are still several issues that have not been adequately solved. First, although the essential loss can be used to explain the relationship between (1 - NDCG) and the loss functions in the pairwise and listwise approaches, it is not clear whether it can also be used to explain the pointwise approach. If it is the case, the essential loss will become really "essential". Second, from the ML point of view, being an upper bound of the evaluation measure might not be sufficient for a good loss function. The reason is that what we really care about is the optimal solution with regards to the loss function. Even if a loss can be the upper bound of (1 - NDCG) everywhere, its optimum might not correspond to the optimum of (1 - NDCG).

The discussions on the "directness" is one step toward solving this problem. A more principled solution should be obtained by investigating the so-called "consistency" of the loss functions, which exactly describes whether the optima with regards to the loss function and the measure can be the same. The consistency of learning methods has been well studied in classification, but not yet for ranking (see Discussions in Section 7). This should be important future work on learning to rank, from the theoretical point of view.

Benchmarking Learning-to-Rank Algorithms

In this section, we introduce a benchmark dataset for learning to rank and investigate the empirical performances of some representative learning-to-rank algorithms on the dataset.

As we know, a standard dataset with standard features and evaluation measures is very helpful for the research on ML. For example, there are benchmark datasets such as Reuters¹ and RCV-1² for text classification, and UCI³ for general ML. However, there were no such benchmark datasets for ranking until the LETOR collection [79] was released in early 2007. In recent years, the LETOR collection has been widely used in the experiments of learning-to-rank papers, and has helped to greatly move forward the research on learning to rank.

6.1 The LETOR Collection

In this subsection, we introduce the LETOR collection, including five aspects: document corpora (together with query sets), document sampling, feature extraction, meta information, and cross validation.

 $^{^{1}\,\}mathrm{http://www.daviddlewis.com/resources/testcollections/reuters21578/}$

 $^{^2\,}http://jmlr.csail.mit.edu/papers/volume5/lewis04a/lyrl2004_rcv1v2_README.htm.$

³ http://archive.ics.uci.edu/ml/

6.1.1 Document Corpora

Two document corpora together with seven query sets were used in the LETOR collection.

6.1.1.1 The "Gov" corpus and six query sets

In TREC 2003 and 2004, a special track for Web IR, named the Web track,⁴ was organized. The track used the "Gov" corpus, which is based on a January, 2002 crawl of the "Gov" domain. There are in total 1,053,110 html documents in this corpus.

Three search tasks were designed in the Web track: topic distillation (TD), homepage finding (HP), and named page finding (NP). TD aims to find a list of entry points for good websites principally devoted to the topic. HP aims at returning the homepage of the query. NP aims to return the page whose name is exactly identical to the query. Generally speaking, there is only one answer for HP and NP. The numbers of queries in these three tasks are shown in Table 6.1.

Due to the large scale of the corpus, it is not feasible to check every document and judge whether it is relevant to a given query. Therefore, the pooling strategy as introduced in Section 1 was used [35].

Many research papers [97, 101, 131, 133] have been published using the three tasks on the "Gov" corpus as their experimental platform.

6.1.1.2 The OHSUMED corpus

The OHSUMED corpus [64] is a subset of MEDLINE, a database on medical publications. It consists of 348,566 records (out of over 7 million) from 270 medical journals during the years of 1987–1991.

Table 6.1 Number of queries in TREC web track.

Topic distillation 50 75 Homepage finding 150 75	Task	TREC2003	TREC2004
1 0 0	Topic distillation	50	75
	Homepage finding	150	75
Named page finding 150 75	Named page finding	150	75

 $[\]overline{^4}$ http://trec.nist.gov/tracks.html.

The fields of a record include title, abstract, MeSH indexing terms, author, source, and publication type.

A query set with 106 queries on the OHSUMED corpus was used in much previous work [97, 131], with each query describing a medical search need (associated with patient information and topic information). The relevance degrees of the documents with respect to the queries are judged by humans, on three levels: definitely relevant, partially relevant, and irrelevant. There are a total of 16,140 query—document pairs with relevance judgments.

6.1.2 Documents Sampling

It is not feasible to extract feature vectors of all the documents in the corpora, due to a reason similar to selecting documents for labeling. A reasonable strategy is to sample some "possibly" relevant documents, and then extract feature vectors for the corresponding query-document pairs.

For the "Gov" corpus, following the suggestions in [88] and [100], the documents were sampled in the following way. First, the BM25 model was used to rank all the documents with respect to each query, and then the top 1000 documents for each query were selected for feature extraction. Please note that this sampling strategy is to ease the experimental investigation, and it is by no means to say that learning to rank can only be applicable in such a re-ranking scenario.

Different from the "Gov" corpus where unjudged documents are regarded as irrelevant, in OHSUMED, the judgments explicitly contain the category of "irrelevant" and the unjudged documents are ignored in the evaluation [64]. Following this practice, in LETOR, only judged documents were used for feature extraction in OHSUMED and all the unjudged documents were ignored. On average, a query has about 152 documents sampled for feature extraction.

6.1.3 Features Extraction

In this subsection, we introduce the feature representation of each document in LETOR. For the "Gov" corpus, 64 features were extracted for each query–document pair, as shown in Table 6.2.

For the OHSUMED corpus, 40 features were extracted in total, as shown in Table 6.3.

Table 6.2 Learning features of TREC.

ID	Feature description
1	Term frequency (TF) of body
2	TF of anchor
3	TF of title
4	TF of URL
5	TF of whole document
6	Inverse document frequency (IDF) of body
7	IDF of anchor
8	IDF of title
9	IDF of URL
10	IDF of whole document
11	TF*IDF of body
12	TF*IDF of anchor
13	TF*IDF of title
14	TF*IDF of URL
15	TF*IDF of whole document
16	Document length (DL) of body
17	
18	DL of title
19	DL of URL
20	DL of whole document
21	BM25 of body
22	BM25 of anchor
23	BM25 of title
24	BM25 of URL
$\frac{25}{26}$	BM25 of whole document LMIR.ABS of body
$\frac{20}{27}$	v .
28	LMIR.ABS of title
29	LMIR.ABS of URL
30	LMIR.ABS of whole document
31	LMIR.DIR of body
32	LMIR.DIR of anchor
33	LMIR.DIR of title
34	LMIR.DIR of URL
35	LMIR.DIR of whole document
36	LMIR.JM of body
37	v
38	LMIR.JM of title
39	LMIR.JM of URL
	(Continued)

 $({\it Continued})$

Table 6.2 (Continued)

ID	Feature description
40	LMIR.JM of whole document
41	Sitemap based term propagation
42	Sitemap based score propagation
43	Hyperlink base score propagation: weighted in-link
44	Hyperlink base score propagation: weighted out-link
45	Hyperlink base score propagation: uniform out-link
46	Hyperlink base feature propagation: weighted in-link
47	Hyperlink base feature propagation: weighted out-link
48	Hyperlink base feature propagation: uniform out-link
49	HITS authority
50	HITS hub
51	PageRank
52	HostRank
53	Topical PageRank
54	Topical HITS authority
55	Topical HITS hub
56	Inlink number
57	Outlink number
58	Number of slash in URL
59	Length of URL
60	Number of child page
61	BM25 of extracted title
62	LMIR.ABS of extracted title
63	LMIR.DIR of extracted title
64	LMIR.JM of extracted title

Table 6.3 Learning features of OHSUMED.

ID	Feature description
1	$\sum_{q_i \in q \cap d} \mathrm{TF}(q_i, d)$ in title
2	$\sum_{q_i \in q \cap d} \log (\mathrm{TF}(q_i, d) + 1)$ in title
3	$\sum_{q_i \in q \cap d} \frac{\operatorname{TF}(q_i, d)}{\operatorname{LEN}(d)}$ in title
4	$\sum_{q_i \in q \cap d} \log \left(\frac{\operatorname{TF}(q_i, d)}{\operatorname{LEN}(d)} + 1 \right)$ in title
5	$\sum_{q_i \in q \cap d} \log(\text{IDF}(q_i))$ in title
6	$\sum_{q_i \in q \cap d} \log(\log(\mathrm{IDF}(q_i)))$ in title
7	$\sum_{q_i \in q \cap d} \log \left(\frac{N}{\mathrm{TF}(q_i, C)} + 1 \right)$ in title
8	$\sum_{q_i \in q \cap d} \log \left(\frac{\operatorname{TF}(q_i, d)}{\operatorname{LEN}(d)} \cdot \log \left(\operatorname{IDF}(q_i) \right) + 1 \right) \text{ in title}$
9	$\sum_{q_i \in q \cap d} \mathrm{TF}(q_i, d) \cdot \log(\mathrm{IDF}(q_i))$ in title
10	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} \cdot \frac{N}{TF(q_i, C)} + 1 \right)$ in title
11	BM25 of title
12	log(BM25) of title

(Continued)

Table 6.3 (Continued)

ID	Feature description
13	LMIR.DIR of title
14	LMIR.JM of title
15	LMIR.ABS of title
16	$\sum_{q_i \in q \cap d} \mathrm{TF}(q_i, d)$ in abstract
17	$\sum_{q_i \in q \cap d} \log (\mathrm{TF}(q_i, d) + 1)$ in abstract
18	$\sum_{q_i \in q \cap d} \frac{\text{TF}(q_i, d)}{\text{LEN}(d)}$ in abstract
19	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} + 1 \right)$ in abstract
20	$\sum_{q_i \in q \cap d} \log(\mathrm{IDF}(q_i))$ in abstract
21	$\sum_{q_i \in q \cap d} \log(\log(\mathrm{IDF}(q_i)))$ in abstract
22	$\sum_{q_i \in q \cap d} \log \left(\frac{N}{\mathrm{TF}(q_i, C)} + 1 \right)$ in abstract
23	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} \cdot \log (\text{IDF}(q_i)) + 1 \right)$ in abstract
24	$\sum_{q_i \in q \cap d} \mathrm{TF}(q_i, d) \cdot \log(\mathrm{IDF}(q_i))$ in abstract
25	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} \cdot \frac{N}{\text{TF}(q_i, C)} + 1 \right)$ in abstract
26	BM25 of abstract
27	log(BM25) of abstract
28	LMIR.DIR of abstract
29	LMIR.JM of abstract
30	LMIR.ABS of abstract
31	$\sum_{q_i \in q \cap d} \mathrm{TF}(q_i, d)$ in title + abstract
32	$\sum_{q_i \in q \cap d} \log (TF(q_i, d) + 1)$ in title + abstract
33	$\sum_{q_i \in q \cap d} \frac{\operatorname{TF}(q_i, d)}{\operatorname{LEN}(d)}$ in title + abstract
34	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} + 1 \right)$ in title + abstract
35	$\sum_{q_i \in q \cap d} \log(\text{IDF}(q_i))$ in title + abstract
36	$\sum_{q_i \in q \cap d} \log(\log(\text{IDF}(q_i)))$ in title + abstract
37	$\sum_{q_i \in q \cap d} \log \left(\frac{N}{\text{TF}(q_i, C)} + 1 \right)$ in title + abstract
38	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} \cdot \log (\text{IDF}(q_i)) + 1 \right)$ in title + abstract
39	$\sum_{q_i \in q \cap d} \mathrm{TF}(q_i, d) \cdot \log(\mathrm{IDF}(q_i))$ in title + abstract
40	$\sum_{q_i \in q \cap d} \log \left(\frac{\text{TF}(q_i, d)}{\text{LEN}(d)} \cdot \frac{N}{\text{TF}(q_i, C)} + 1 \right)$ in title + abstract
41	BM25 of title + abstract
42	log(BM25) of title + abstract
43	LMIR.DIR of title + abstract
44	LMIR.JM of title $+$ abstract
45	LMIR.ABS of title + abstract

6.1.4 Meta Information

In addition to the features, meta information that can be used to reproduce these features and even other new features has also been provided

in LETOR. There are three kinds of meta information:

- Statistical information about the corpus, such as the total number of documents, the number of streams, and the number of (unique) terms in each stream.
- Raw information on the documents associated with each query, such as the term frequency and the document length.
- Relational information, such as the hyperlink graph, the sitemap information, and the similarity relationship matrix of the corpus.

With the meta information, one can reproduce existing features, tune their parameters, investigate new features, and perform advanced research such as relational ranking [103, 104].

6.1.5 Cross Validation

In total, there are seven datasets in the LETOR collection, i.e., TD2003, TD2004, NP2003, NP2004, HP2003, HP2004, and OHSUMED. Each of these datasets was partitioned into five parts with about the same number of queries, denoted as S1, S2, S3, S4, and S5, in order to conduct five-fold cross validation. For each fold, three parts are used for training the ranking model, one part for tuning the hyper parameters of the ranking algorithm (e.g., the number of iterations in RankBoost, and the combination coefficient in the objective function of Ranking SVM), and the remaining part to evaluate the ranking performance of the learned model (see Table 6.4). The average performance over the five folds is used to measure the overall performance of a learning-to-rank algorithm.

Table 6.4 Data partitioning for five-fold cross validation.

Folds	Training set	Validation set	Test set
Fold1	{S1,S2,S3}	S4	S5
Fold2	${S2,S3,S4}$	S5	S1
Fold3	${S3,S4,S5}$	S1	S2
Fold4	${S4,S5,S1}$	S2	S3
Fold5	{S5,S1,S2}	S3	S4

The LETOR collection, containing the aforementioned feature representations of documents, their relevance judgments with respective queries, and the partitioned training, validation, and test sets can be downloaded from http://research.microsoft.com/~LETOR/.5

6.2 Experimental Results on LETOR

In this subsection, we introduce the experiments on LETOR to evaluate several representative learning-to-rank algorithms, and make discussions on the experimental results.⁶ Three widely used measures are adopted for the evaluation: precision at position k (P@k) [8], MAP [8], and NDCG at position k (N@k) [66]. The official evaluation tool provided with LETOR was used in the evaluation process.

We have evaluated seven representative learning-to-rank algorithms in total. For the pointwise approach, we tested the regression based method. For the pairwise approach, we tested Ranking SVM, RankBoost, and FRank. For the listwise approach, we tested ListNet, AdaRank, and SVM^{map}. To make fair comparisons, we tried to use the same setting for all the algorithms. Firstly, most algorithms use the linear ranking function, except RankBoost, which uses binary weak rankers. Secondly, all the algorithms used MAP on the validation set for model selection. Some detailed experimental settings are listed as follows.

As for the linear regression based algorithm, the validation set was used to select a good mapping from the ground truth labels to real numbers. For Ranking SVM, the public tool of SVMlight⁷ was employed and the validation set was used to tune the parameter C in its loss function. For RankBoost, the weak ranker was defined on the basis of a single feature with 255 possible thresholds. The validation set was used to determine the best number of iterations. For FRank, to efficiently minimize the fidelity loss, a generalized additive model was adopted. The validation set was used to determine the number of weak learners in the additive model. For ListNet, the validation set was used to deter-

 $[\]overline{^5}$ Note that the LETOR collection is being frequently updated. It is expected that more datasets will be added in the future.

 $^{^6}$ These results are also published at the website of LETOR.

⁷ http://svmlight.joachims.org/.

mine the best mapping from the ground truth label to real numbers in order to use the Luce model, and to determine the optimal number of iterations in the gradient descent process. For AdaRank, MAP was set as the IR evaluation measure to be optimized, and the validation set was used to determine the number of iterations. For SVM^{map} [136], the publicly available tool SVM^{map} was employed, and the validation set was used to determine the parameter C in its loss function.

The ranking performances of the aforementioned algorithms are listed in Tables 6.5–6.11. According to these experimental results, we find that listwise ranking algorithms perform very well on most datasets. Among the three listwise ranking algorithms, ListNet seems

Table 6.5 Results on TD2003.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.320	0.307	0.326	0.320	0.260	0.178	0.241
RankSVM	0.320	0.344	0.346	0.320	0.293	0.188	0.263
RankBoost	0.280	0.325	0.312	0.280	0.280	0.170	0.227
FRank	0.300	0.267	0.269	0.300	0.233	0.152	0.203
ListNet	0.400	0.337	0.348	0.400	0.293	0.200	0.275
AdaRank	0.260	0.307	0.306	0.260	0.260	0.158	0.228
SVM^{map}	0.320	0.320	0.328	0.320	0.253	0.170	0.245

Table 6.6 Results on TD2004.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.360	0.335	0.303	0.360	0.333	0.249	0.208
RankSVM	0.413	0.347	0.307	0.413	0.347	0.252	0.224
RankBoost	0.507	0.430	0.350	0.507	0.427	0.275	0.261
FRank	0.493	0.388	0.333	0.493	0.378	0.262	0.239
ListNet	0.360	0.357	0.317	0.360	0.360	0.256	0.223
AdaRank	0.413	0.376	0.328	0.413	0.369	0.249	0.219
SVM^{map}	0.293	0.304	0.291	0.293	0.302	0.247	0.205

Table 6.7 Results on NP2003.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.447	0.614	0.665	0.447	0.220	0.081	0.564
RankSVM	0.580	0.765	0.800	0.580	0.271	0.092	0.696
RankBoost	0.600	0.764	0.807	0.600	0.269	0.094	0.707
FRank	0.540	0.726	0.776	0.540	0.253	0.090	0.664
ListNet	0.567	0.758	0.801	0.567	0.267	0.092	0.690
AdaRank	0.580	0.729	0.764	0.580	0.251	0.086	0.678
SVM^{map}	0.560	0.767	0.798	0.560	0.269	0.089	0.687

⁸ http://svmrank.yisongyue.com/svmmap.php

Table 6.8 Results on NP2004.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.373	0.555	0.653	0.373	0.200	0.082	0.514
RankSVM	0.507	0.750	0.806	0.507	0.262	0.093	0.659
RankBoost	0.427	0.627	0.691	0.427	0.231	0.088	0.564
FRank	0.480	0.643	0.729	0.480	0.236	0.093	0.601
ListNet	0.533	0.759	0.812	0.533	0.267	0.094	0.672
AdaRank	0.480	0.698	0.749	0.480	0.244	0.088	0.622
SVM^{map}	0.520	0.749	0.808	0.520	0.267	0.096	0.662

Table 6.9 Results on HP2003.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.420	0.510	0.594	0.420	0.211	0.088	0.497
RankSVM	0.693	0.775	0.807	0.693	0.309	0.104	0.741
RankBoost	0.667	0.792	0.817	0.667	0.311	0.105	0.733
FRank	0.653	0.743	0.797	0.653	0.289	0.106	0.710
ListNet	0.720	0.813	0.837	0.720	0.320	0.106	0.766
AdaRank	0.733	0.805	0.838	0.733	0.309	0.106	0.771
SVM^{map}	0.713	0.779	0.799	0.713	0.309	0.100	0.742

Table 6.10 Results on HP2004.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.387	0.575	0.646	0.387	0.213	0.08	0.526
RankSVM	0.573	0.715	0.768	0.573	0.267	0.096	0.668
RankBoost	0.507	0.699	0.743	0.507	0.253	0.092	0.625
FRank	0.600	0.729	0.761	0.600	0.262	0.089	0.682
ListNet	0.600	0.721	0.784	0.600	0.271	0.098	0.690
AdaRank	0.613	0.816	0.832	0.613	0.298	0.094	0.722
SVM^{map}	0.627	0.754	0.806	0.627	0.280	0.096	0.718

Table 6.11 Results on OHSUMED.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	0.446	0.443	0.411	0.597	0.577	0.466	0.422
RankSVM	0.496	0.421	0.414	0.597	0.543	0.486	0.433
RankBoost	0.463	0.456	0.430	0.558	0.561	0.497	0.441
FRank	0.530	0.481	0.443	0.643	0.593	0.501	0.444
ListNet	0.533	0.473	0.441	0.652	0.602	0.497	0.446
AdaRank	0.539	0.468	0.442	0.634	0.590	0.497	0.449
SVM^{map}	0.523	0.466	0.432	0.643	0.580	0.491	0.445

to be better than the others. AdaRank and SVM^{map} obtain similar performances. Pairwise ranking algorithms obtain good ranking accuracy on some (although not all) datasets. For example, RankBoost offers the best performance on TD2004 and NP2003; Ranking SVM shows very

promising results on NP2003 and NP2004; and FRank achieves very good results on TD2004 and NP2004. Comparatively speaking, simple linear regression performs worse than the pairwise and listwise ranking algorithms. Its performance is not so good on most datasets.

We have also observed that most ranking algorithms perform differently on different datasets. They may perform very well on some datasets but not so well on other datasets. To evaluate the overall ranking performances of an algorithm, we use the number of other algorithms that it can beat over all the seven datasets as a measure. That is:

$$S_i(M) = \sum_{j=1}^{7} \sum_{k=1}^{7} I_{\{M_i(j) > M_k(j)\}},$$

where j is the index of a dataset, i and k are the indexes of algorithms, $M_i(j)$ is the performance of i-th algorithm on j-th dataset, and $I_{\{\cdot\}}$ is the indicator function.

It is clear that the larger the $S_i(M)$, the better the *i*-th algorithm performs. For ease of reference, we call this measure the *winning number*. Table 6.12 shows the winning number for all the algorithms under investigation. From this table, we make the following observations:

(1) In terms of NDCG@1, the listwise ranking algorithms perform the best, followed by the pairwise ranking algorithms, while the pointwise algorithm performs the worst. Among the three listwise ranking algorithms, ListNet is better than AdaRank, while SVM^{map} performs a little worse than the others. The three pairwise ranking algorithms achieve comparable results, among which Ranking SVM seems to be slightly better than the other two.

Table 6.12 Winning number of each algorithm.

Algorithm	N@1	N@3	N@10	P@1	P@3	P@10	MAP
Regression	4	4	4	5	5	5	4
RankSVM	21	22	22	21	22	22	24
RankBoost	18	22	22	17	22	23	19
FRank	18	19	18	18	17	23	15
ListNet	29	31	33	30	32	35	33
AdaRank	26	25	26	23	22	16	27
SVM^{map}	23	24	22	25	20	17	25

- (2) In terms of NDCG@3 and NDCG@10, ListNet and AdaRank perform much better than the pairwise and pointwise ranking algorithms, while the performance of SVM^{map} is very similar to the pairwise ranking algorithms.
- (3) In terms of P@1 and P@3, the listwise ranking algorithms perform the best, followed by the pairwise ranking algorithms, while the pointwise ranking algorithm performs the worst. Among the three listwise ranking algorithms, ListNet is better than AdaRank and SVM^{map}. The three pairwise ranking algorithms achieve comparable results, among which Ranking SVM seems to be slightly better than the other two algorithms.
- (4) In terms of P@10, ListNet performs much better than the pairwise and pointwise ranking algorithms, while the performances of AdaRank and SVM^{map} are not as good as those of the pairwise ranking algorithms.
- (5) In terms of MAP, the listwise ranking algorithms are in general better than the pairwise ranking algorithms. Furthermore, the variance among the three pairwise ranking algorithms in terms of MAP is much larger than the variance among the three algorithms in terms of other measures (e.g., P@1,3, and 10). The possible explanation is that since MAP involves all the documents associated with a query in the evaluation process, it can better differentiate algorithms.

To summarize, the experimental results show that the listwise algorithms have certain advantages over other algorithms, especially for top positions of the ranking result.

Here, we would like to point out that the above experimental results are still primal, since the result of almost every algorithm can be further improved. For example, for regression, we can add a regularization item to make it more robust; for Ranking SVM, we can run more steps of the iteration so as to guarantee a better convergence of the optimization; for ListNet, we can also add a regularization item to its loss function and make it more generalizable to the test set.

Statistical Ranking Theory

As a new ML problem, ranking is not only about effective algorithms but also about the theory behind these algorithms. In this section, we will introduce the so-called statistical ranking theory [75, 76], and will focus on the generalization analysis of learning-to-rank methods.

Actually, theoretical analysis on an algorithm always plays an important role in ML. This is because in practice, one can only observe experimental results on small-scale datasets (e.g., the experimental results on the LETOR collection as introduced in Section 6). To some extent, such empirical results might not be fully trustworthy, because a small training set sometimes cannot fully realize the potential of a learning algorithm, and a small test set sometimes cannot reflect the true performance of an algorithm (the input and output spaces are too large to be well represented by a small number of samples). In this regard, theories are solely needed in order to analyze the performance of a learning algorithm when the training data is infinite and the test data is perfectly sampled from the input and output spaces.

For example, the generalization analysis on an algorithm is concerned with whether and at what rate its empirical risk computed from the training data will converge to the expected risk on any test data in the input and output spaces, when the number of training samples approaches infinity. Sometimes, we alternatively represent the generalization ability of an algorithm using the bound of the difference between its expected risk and empirical risk, and see whether and at what rate the bound will converge to zero when the number of training samples approaches infinity. In the context of learning to rank, the expected risk measures the average error that a ranking model would make on a randomly sampled input instance (document, document pairs, or all documents associated with a query), while the empirical risk is an estimate of the expected risk based on the training data. In general, an algorithm is regarded as better than the other algorithm if its empirical risk can converge to the expected risk but that of the other cannot. Furthermore, an algorithm is regarded as better than the other if its corresponding convergence rate is faster than that of the other.

To facilitate the discussions on (but not limited to) the generalization ability, a theoretical framework is needed. We will first review some conventional generalization analyses on ranking and their corresponding theoretical frameworks. After that, a recently-proposed query-level ranking framework and its use in analyzing the generalization ability of ranking methods will be introduced.

Note that it is unavoidable that a lot of mathematics will be used in this section. It is safe, however, to skip this whole section, if one only wants to know the algorithmic development of learning to rank.

7.1 Conventional Generalization Analyses on Ranking

Most of the conventional generalization analyses on ranking are for the pairwise approach of learning to rank. The theoretical framework behind these analyses is given in the following subsection.

7.1.1 Theoretical Framework for Pairwise Ranking

The basic assumption in the theoretical framework is that the documents and their relevance degrees are i.i.d. random variables. In this case, the document pairs, as random variables, are not independent of each other. To handle the challenge, *U*-statistics is used as the tool to perform generalization analysis.

Assume that (x_u, y_u) and (x_v, y_v) are i.i.d. random variables according to distribution P_{XY} , where x stands for the document and y stands for the ground truth label of the document. (x_u, y_u) and (x_v, y_v) construct a pair. Given the scoring function f, loss comes up if the documents are not ranked according to their ground truth labels. Suppose the loss function is $L(f, x_u, x_v, y_{u,v})$, where $y_{u,v} = 2 \cdot I_{\{y_u \succ y_v\}} - 1$. Then the expected risk for ranking is defined as:

$$R(f) = \int_{(\mathcal{X} \times \mathcal{Y})^2} L(f, x_u, x_v, y_{u,v}) P_{XY}(dx_u, dy_u) P_{XY}(dx_v, dy_v).$$
 (7.1)

Intuitively, the expected risk means the average loss that a ranking model f would make for a random document pair. Since it is almost impossible to compute the expected risk, in practice, the empirical risk on the training set is used as an estimate of the expected risk. In particular, given the training data $\{(x_j, y_j)\}_{j=1}^m$, the empirical risk can be defined with the following U-statistics:

$$\hat{R}(f) = \frac{1}{C_m^2} \sum_{u=1}^m \sum_{v=u+1}^m L(f, x_u, x_v, y_{u,v}), \tag{7.2}$$

where $C_m^2 = m(m-1)/2$.

Specifically, when the ground truth is given as a binary relevance degree, the relevant (positive) document is denoted as x^+ according to P^+ and the irrelevant (negative) document is denoted as x^- according to P^- . Given the training data $\mathbf{x}^+ = \{x_j^+\}_{j=1}^{m^+}$ and $\mathbf{x}^- = \{x_j^-\}_{j=1}^{m^-}$ (where m^+ and m^- are the numbers of positive and negative documents in the training data, respectively), the expected risk and the empirical risk are refined as follows:

$$R(f) = \int_{\mathcal{X}^2} L(f, x^+, x^-) P^+(dx^+) P^-(dx^-), \tag{7.3}$$

$$\hat{R}(f) = \frac{1}{m^+ m^-} \sum_{u=1}^{m^+} \sum_{v=1}^{m^-} L(f, x^+, x^-).$$
 (7.4)

7.1.2 Generalization Analysis on the Pairwise Approach

With the above theoretical framework, several tools have been used to conduct generalization analyses for the pairwise approach of learning to rank. Here we give three examples. The VC dimension and rank shatter coefficient measure the complexity of the function class, while stability measures the robustness of a ranking algorithm. Basically, simpler ranking models generalize better, and more robust ranking algorithms generalize better.

Based on the VC dimension: In [47], the VC dimension [125, 126] is used to obtain the generalization bound for RankBoost. It is clear that the bound converges to zero at a rate of $O(\max\{\sqrt{\log(m^+)/m^+}, \sqrt{\log(m^-)/m^-}\})$.

Theorem 7.1. Assume that all the weak learners belong to the function class \mathcal{F}' , which has a finite VC dimension V, the scoring function belongs to function class \mathcal{F} . Let S^+ and S^- be positive and negative samples of size m^+ and m^- , respectively. Then with probability at least $1 - \delta$, the following inequality holds:

$$\forall f \in \mathcal{F}, |R(f) - \hat{R}(f)| \le 2\sqrt{\frac{V'(\log\frac{2m^{+}}{V'} + 1) + \log\frac{18}{\delta}}{m^{+}}} + 2\sqrt{\frac{V'(\log\frac{2m^{-}}{V'} + 1) + \log\frac{18}{\delta}}{m^{-}}},$$
(7.5)

where $V' = 2(V+1)(T+1)\log_2(e(T+1))$ and T is the number of weak rankers in RankBoost.

The above theorem is only applicable to the bipartite case. Clemencon et al. [27] proposed another theorem using the properties of U-Statistics, which can handle the general case.

Based on the rank shatter coefficient: The notion of bipartite rankshatter coefficient, denoted as $r(\mathcal{F}, m^+, m^-)$, was proposed by Agarwal et al. [1]. Here \mathcal{F} is the class of scoring functions, m^+ is the number of positive documents and m^- is the number of negative documents. This new notion has a similar meaning to the shattering coefficient (growth function) in the VC theory [125, 126].

Using the bipartite rank-shatter coefficient as a tool, the following generalization theorem has been proven. As shown in [1], for

the class of linear ranking functions in the one-dimensional feature space, $r(\mathcal{F}, m^+, m^-)$ is a constant, regardless of the values of m^+ and m^- . In this case, the bound converges to zero at a rate of $O(\max\{1/\sqrt{m^+}, 1/\sqrt{m^-}\})$, and is therefore tighter than the bound for RankBoost given in inequality (7.5). For the class of linear ranking functions in the d-dimensional feature space (d > 1), $r(\mathcal{F}, m^+, m^-)$ is of the order $O((m^+m^-)^d)$, and in this case the bound has a similar convergence rate to that based on the VC dimension, i.e., $O(\max\{\sqrt{\log(m^+)/m^+}, \sqrt{\log(m^-)/m^-}\})$.

Theorem 7.2. Let \mathcal{F} be the class of real-valued functions on \mathcal{X} , then $\forall 0 < \delta < 1$, with probability at least $1 - \delta$,

$$\forall f \in \mathcal{F}, |R(f) - \hat{R}(f)| \le \sqrt{\frac{8(m^{+} + m^{-})\left(\log\frac{4}{\delta} + \log r(\mathcal{F}, 2m^{+}, 2m^{-})\right)}{m^{+}m^{-}}}.$$
 (7.6)

Furthermore, Rajaram and Agarwal [108] generalized the above theory to the k-partite case using the k-partite rank-shatter coefficient.

Based on the stability theory: Agarwal and Niyogi [2] used stability as a tool and derived the generalization bound for some bipartite ranking algorithms.

Let \mathcal{A} be a ranking algorithm, and let L be the loss function that is minimized in \mathcal{A} . Suppose we have learned a ranking model f_1 from the training data using algorithm \mathcal{A} . Then we replace a positive training document with another document and learn a new model f_2 from the new training data. Similarly, we replace a negative training document with another document and learn a new model f_3 . We say that \mathcal{A} has uniform loss stability (α, β) with respect to L, if the difference between the losses with respect to f_1 and f_2 on any unseen document pair (x+,x-) is smaller than $\alpha(m^+,m^-)$, and that with respect to f_1 and f_3 is smaller than $\beta(m^+,m^-)$. In other words, if the model learned from the training data is robust to the small change in the training data, the algorithm is regarded as having certain stability.

The generalization bound they obtained is as shown in Theorem 7.3. As shown in [12], for many algorithms, $\alpha(m^+, m^-)$ and

 $\beta(m^+,m^-)$ are of the order $O(\frac{1}{m^+})$ and $O(\frac{1}{m^-})$, respectively. Therefore, the bound given in Theorem 7.3 will converge to zero at a rate of $O(\max\{1/\sqrt{m^+},1/\sqrt{m^-}\})$. In this regard, this bound is tighter than the bound based on the Rank Shatter Coefficient (see Inequality (7.6)) except for the case of using linear ranking functions in the one-dimensional feature space. This is understandable since the use of stability can lead to a data-dependent bound [12], which is usually tighter than data-independent bounds.

Theorem 7.3. Let L be the loss function with $L(f; x^+, x^-) \in [0, B]$, and \mathcal{A} have uniform loss stability (α, β) . Then $\forall 0 < \delta < 1$, with probability at least $1 - \delta$, the following inequality holds:

$$R_{L}(f_{S^{+},S^{-}})$$

$$\leq \hat{R}_{L}(f_{S^{+},S^{-}}) + \alpha(m^{+},m^{-}) + \beta(m^{+},m^{-})$$

$$+ \sqrt{\frac{\{m^{-}(2m^{+}\alpha(m^{+},m^{-}) + B)2 + m^{+}(2m^{-}\beta(m^{+},m^{-}) + B)^{2}\}\log\frac{1}{\delta}}{2m^{+}m^{-}}}.$$
(7.7)

Due to the basic assumption of the theoretical framework, the aforementioned theorems can only be used when the ground truth labels are given as relevance degrees of all the documents. However, as we mentioned before, the ground truth labels can also be given in terms of the pairwise preference and even total order of the documents. In this case, a new framework needs to be developed.

When the ground truth is given as a pairwise preference, it is more reasonable to assume document pairs and their ground truth labels are i.i.d. random variables. In this way, ranking is actually formalized as a standard classification problem on document pairs. By using the theoretical results of classification, one can also get a generalization bound. For ease of reference, we refer to this way as taking the "average view" on the generalization analysis, and refer to the work introduced in this subsection as taking the "U-statistics view" on the generalization analysis.

Note that the "average view" is also technically sound. The intuition is not right that two document pairs cannot be independent of each other when they share a common document. The reason is that the dependence (or independence) is actually defined with regards to random variables but not their values. Therefore, as long as two document pairs are sampled and labeled in an independent manner, they are i.i.d. random variables no matter whether their values (the specific documents in the pair) have overlap or not. More discussions about the average view and the U-statistics view will be given in the next subsection.

7.2 A Query-level Ranking Framework

In [76], Lan et al. pointed out the limitation of the aforementioned generalization analyses, from the IR perspective.

As we know, the generalization ability is concerned with the difference between the empirical risk and the expected risk of learning, and these risks are highly related to how one evaluates the performance of a learning algorithm. As mentioned earlier in the tutorial, all the IR evaluation measures are defined at the query level. Therefore, the definition of the risks in ranking and the generalization analysis for ranking should also be at the query level.

In contrast, the previous generalization analyses are either performed at the document level or at the document pair level. Therefore, Lan et al. argued that a novel theoretical framework needs to be developed for learning to rank, to facilitate (but not limited to) the query-level generalization analysis. They call such a framework the query-level ranking framework.

Let \mathcal{Q} be the query space, \mathcal{D} be the document space, and \mathcal{X} be the d-dimensional space containing a feature representation of each document. Let q be a random variable defined on the query space with an unknown probability distribution $P_{\mathcal{Q}}$. Denote f as the scoring function, and use a loss function L(f;q) to measure the loss for each query. Then the goal of ranking is to minimize the following expected query-level risk:

$$R(f) = \int_{\mathcal{Q}} L(f;q) P_{\mathcal{Q}}(dq). \tag{7.7}$$

Intuitively, the expected query-level risk means the average loss that the ranking model f would make on a randomly sampled query. Since the distribution $P_{\mathcal{Q}}$ is unknown, the following empirical query-level risk on the training set is used as the estimate of the expected query-level risk,

$$\hat{R}(f) = \frac{1}{n} \sum_{i=1}^{n} L(f; q_i), \tag{7.8}$$

where q_1, \ldots, q_n are i.i.d. observations with the same distribution as that of q.

The forms of L(f;q) corresponding to the pointwise, pairwise, and listwise approaches are given as follows.

7.2.1 The Pointwise Approach

In the pointwise approach, each document x is given a ground truth label $y \in \mathcal{Y}$, which stands for the relevance degree of x with regards to a specific query q.

For a fixed query q, (x,y) is assumed to be a random variable sampled according to probability distribution $\mathcal{D}_q^{(1)}$ (which is dependent on query q). Then, L(f;q) is defined as below:

$$L(f;q) = \int_{\mathcal{X} \times \mathcal{V}} L(f;x,y) \mathcal{D}_q^{(1)}(dx,dy), \tag{7.9}$$

where L(f;x,y) is the loss² of the scoring function f on sample (x,y).

As the distribution $\mathcal{D}_q^{(1)}$ is unknown, the average of the loss over a group of training samples is used to estimate L(f;q):

$$\hat{L}(f;q) = \frac{1}{m} \sum_{j=1}^{m} L(f;x_j, y_j), \tag{7.10}$$

where $\{x_j, y_j\}_{j=1}^m$ stands for m i.i.d. observations with distribution $\mathcal{D}_q^{(1)}$.

¹ For example, $\mathcal{Y} = \{0,1\}$ for binary classification based algorithms, and $\mathcal{Y} = \mathbb{R}$ (where \mathbb{R} stands for the space of real values) for regression based algorithms.

² For example, L(f;x,y) can be defined as $I_{\{\operatorname{sgn}(f(x))\neq y\}}$ (indicator function) in binary classification and $(f(x)-y)^2$ in regression.

7.2.2 The Pairwise Approach

As introduced in the previous subsection, there are two views on the generalization analysis of the pairwise approach: the *U-statistics* view and the *average* view. For both views, it is not difficult to extend them to be fitted into the query-level ranking framework.

(1) The U-statistics view

With the U-statistics view, one assumes i.i.d. distribution of the documents and their ground truth labels with respect to a query. Given two documents associated with query q and their ground truth labels, (x_1, y_1) and (x_2, y_2) , (x_1, x_2, y_1, y_2) is regarded as a random variable with probabilistic distribution $\mathcal{D}_q^{(2)}$. Then L(f;q) can be defined as follows:

$$L(f;q) = \int_{(\mathcal{X}\times\mathcal{Y})^2} L(f;x_1,x_2,y_1,y_2) \mathcal{D}_q^{(2)}(dx_1,dx_2,dy_1,dy_2), \quad (7.11)$$

where $L(f; x_1, x_2, y_1, y_2)$ is the loss of f on sample (x_1, x_2, y_1, y_2) .

As the distribution $\mathcal{D}_q^{(2)}$ is unknown, the following *U*-statistics equation is used to estimate L(f;q):

$$\hat{L}(f;q) = \frac{1}{C_m^2} \sum_{u=1}^m \sum_{v=u+1}^m L(f; x_u, x_v, y_u, y_v), \tag{7.12}$$

where $(x_u, x_v, y_u, y_v), u, v = 1, ..., m$ can be viewed as C_m^2 observations with distribution $\mathcal{D}_q^{(2)}$, which might not be independent of each other.

Since two document pairs are not independent under the aforementioned assumptions, to perform the generalization analysis, one needs to use some advanced statistical tools such as [27, 124].

(2) The average view

The average view assumes the i.i.d. distribution of document pairs. More specifically, with the average view, each document pair (x_1, x_2) is given a ground truth label $y \in \mathcal{Y} = \{-1, 1\}$, where y = 1 indicates that document x_1 is more relevant than x_2 and y = -1 otherwise. Then (x_1, x_2, y) is assumed to be a random variable with probabilistic

distribution $\mathcal{D}_q^{(3)}$, and L(f;q) can be defined as follows:

$$L(f;q) = \int_{\mathcal{X} \times \mathcal{X} \times \mathcal{Y}} L(f;x_1, x_2, y) \mathcal{D}_q^{(3)}(dx_1, dx_2, dy),$$
 (7.13)

where $L(f; x_1, x_2, y)$ is the loss of f on sample (x_1, x_2, y) .

As the distribution $\mathcal{D}_q^{(3)}$ is unknown, the average of the loss over training document pairs is used to estimate L(f;q):

$$\hat{L}(f;q) = \frac{1}{\tilde{m}} \sum_{j=1}^{\tilde{m}} L(f; x_{j_1}, x_{j_2}, y_j), \tag{7.14}$$

where $\{x_{j_1}, x_{j_2}, y_j\}_{j=1}^{\tilde{m}}$ stands for \tilde{m} i.i.d. observations with distribution $\mathcal{D}_q^{(3)}$, and \tilde{m} is the number of document pairs associated with query q in the training data.

7.2.3 The Listwise Approach

In the listwise approach, each query is represented by a group of documents (denoted by \mathbf{x}) and their ground truth labels, and queries are assumed to be i.i.d. random variables. As mentioned earlier, there are two types of output spaces used in the listwise approach. For simplicity and clarity, in this subsection, our discussion focuses on the case where π_y is used as the ground truth label. Furthermore, we will take re-ranking as our target application,³ where there are the same number of documents (i.e., m) for every query under investigation.

Let \mathcal{X}^m be the input space, whose elements are m feature vectors corresponding to m documents associated with a query.⁴ Let \mathcal{Y} be the output space, whose elements are permutations of m documents. Then (\mathbf{x}, π_y) can be regarded as a random variable sampled from the space $\mathcal{X}^m \times \mathcal{Y}$ according to an unknown probability distribution $P(\cdot, \cdot)$.

Define

$$L(f;q) = L(f; \mathbf{x}, \pi_y), \tag{7.15}$$

³ In the re-ranking application, "all the documents" may mean all the documents under investigation, e.g., the top 1000 documents with respect to a query returned by a conventional ranking model like BM25.

⁴ We call m the list length in this tutorial, and suppose $m \ge 3$. Otherwise the listwise approach will be reduced to the pointwise or pairwise approach.

where $L(f; \mathbf{x}, \pi_y)$ denotes a listwise ranking loss (e.g., the likelihood loss in [129] and the K-L divergence loss in [17]) defined on the random variable (\mathbf{x}, π_y) and a scoring function f.

7.3 Query-level Generalization Analysis

In this subsection, we review the work on query-level generalization analysis by using the aforementioned query-level ranking framework. Here, the query-level generalization analysis is concerned with whether and at what convergence rate, the empirical query-level risk converges to the expected query-level risk, when the number of training queries approaches infinity. Please note the differences between the query-level generalization analysis and the previous work on generalization analysis as reviewed in Section 7.1.1. As for the query-level generalization analysis, one only cares about the number of training queries, but not the number of training documents. In previous work, however, it is the number of training documents that really matters for the generalization bound.

7.3.1 On the Pairwise Approach

In [76], the stability theory [12] was extended to perform the query-level generalization analysis on the pairwise approach. The *average view* is taken in the analysis.

To assist the analysis, the definition of uniform leave-one-query-out pairwise loss stability (also referred to as query-level stability for short) was given in [76]. Suppose we have learned a ranking model f_1 from a training set with n queries, using an algorithm \mathcal{A} . Suppose L is the loss function that is minimized in algorithm \mathcal{A} . Then we randomly remove a query and all its associated document pairs from the training data, and learn a ranking model f_2 from the new training set. If the difference between the losses with respect to f_1 and f_2 on any unseen document pair (x_1, x_2) is smaller than $\tau(n)$, we say the algorithm \mathcal{A} has uniform leave-one-query-out pairwise loss stability with coefficient τ with respect to L.

Based on the concept of query-level stability, a query-level generalization bound has been derived in [76], as shown in Theorem 7.4.

The theorem states that if a pairwise ranking algorithm has querylevel stability, then with a large probability, the expected query-level risk can be bounded by the empirical query-level risk and a term that depends on the query number and the stability of the algorithm.

Theorem 7.4. Let \mathcal{A} be a learning-to-rank algorithm, $(q_1, S^{(1)}), \ldots,$ $(q_n, S^{(n)})$ be n training queries, and let L be the pairwise loss function. If

- (1) $\forall (q_1, S^{(1)}), \dots, (q_n, S^{(n)}), \forall q \in \mathcal{Q}, (x_1, x_2, y) \in \mathcal{X} \times \mathcal{X} \times \mathcal{Y},$ $|L(f_{(q_i, S^{(i)})_{i=1}^n}, x_1, x_2, y)| \leq B,$ (2) \mathcal{A} has query-level stability with coefficient τ ,

then $\forall \delta \in (0,1)$ with probability at least $1-\delta$ over the samples of $\{(q_i, S^{(i)})\}_{i=1}^n$ in the product space $\prod_{i=1}^n \{\mathcal{Q} \times (\mathcal{X} \times \mathcal{X} \times \mathcal{Y})^{\infty}\}$, the following inequality holds:

$$R_L(f_{\{(q_i,S^{(i)})\}_{i=1}^n}) \le \widehat{R}_L(f_{\{(q_i,S^{(i)})\}_{i=1}^n}) + 2\tau(n) + (4n\tau(n) + B)\sqrt{\frac{\log\frac{1}{\delta}}{2n}}.$$
 (7.16)

When using this theorem to perform the query-level generalization analysis on pairwise ranking algorithms, what one needs to do is to compute the query-level stability coefficient $\tau(n)$ of the algorithms.

Query-level generalization bound for ranking SVM: As proven in [76], Ranking SVM has query-level stability with coefficient $\tau(n) = \frac{4\kappa^2}{\lambda n} \times$ $\max \frac{\tilde{m}^{(i)}}{\frac{1}{n}\sum_{i=1}^{n}\tilde{m}^{(i)}}$, where $\tilde{m}^{(i)}$ is the number of document pairs associated with query q_i , and $\forall x \in \mathcal{X}, K(x,x) \leq \kappa^2 < \infty$.

On this basis, one can have the following discussions regarding the query-level generalization ability of Ranking SVM:

• When the number of training queries approaches infinity, with a large probability the empirical query-level risk of Ranking SVM will converge to its expected query-level risk, at a rate of $O(\frac{1}{\sqrt{n}})$.

• When the number of training queries is finite, the expected query-level risk and the empirical query-level risk are not necessarily close to each other.

Query-level Generalization Bound for IR-SVM: As proven in [76], IR-SVM [16] has a query-level stability with coefficient $\tau(n) = \frac{4\kappa^2}{\lambda n}$. On this basis, one can find that:

- When the number of training queries approaches infinity, with a large probability the empirical query-level risk of IR-SVM will converge to its expected query-level risk, at a convergence rate of $O(\frac{1}{\sqrt{n}})$.
- When the number of queries is finite, the query-level generalization bound is a decreasing function of the number of training queries.

By comparing the query-level generalization abilities of Ranking SVM and IR-SVM, we can find that the convergence rates of the empirical query-level risks to the expected query-level risks for these two algorithms are both $O(\frac{1}{\sqrt{n}})$. However, by comparing the case with a finite number of training queries, the bound for IR-SVM is much tighter than that for Ranking SVM.

7.3.2 On the Listwise Approach

In [75], the theory of the Rademacher average [10, 11] was extended to perform the query-level generalization analysis on the listwise approach. Specifically, the algorithms that minimize listwise ranking losses introduced in Section 4.2 were taken as examples in the analysis.

The Rademacher average measures how much the function class can fit random noise, which is defined below.

Definition 7.1. For a function class \mathcal{G} , the empirical Rademacher average is defined as:

$$\widehat{\mathcal{R}}(\mathcal{G}) = E_{\sigma} \sup_{g \in \mathcal{G}} \frac{1}{n} \sum_{i=1}^{n} \sigma_{i} g(z_{i}), \tag{7.17}$$

where $z_i, i = 1, ..., n$ are i.i.d. random variables, and $\sigma_i, i = 1, ..., n$ are i.i.d. random variables, with probability $\frac{1}{2}$ of taking a value of 1 or -1.

By fitting the theory of the Rademacher average [10, 11] to the query-level ranking framework, the query-level generalization bound for listwise ranking algorithms have been derived [75], as shown in the following theorem. Here it is assumed that $\forall x \in \mathcal{X}, ||x|| \leq M$, and the ranking function f is learned from the linear function class $\mathcal{F} = \{x \to w^T x : ||w|| \le B.\}, \text{ for simplicity. In this case, one has } \forall x \in$ $\mathcal{X}, |f(x)| \leq BM.$

Theorem 7.5. Let \mathcal{A} denote a listwise ranking algorithm, and let $L_{\mathcal{A}}(f;\mathbf{x},\pi_y)$ be its listwise loss, given the training data S= $\{(\mathbf{x}^{(i)}, \pi_y^{(i)}), i = 1, \dots, n\}, \text{ if } \forall f \in \mathcal{F}, (\mathbf{x}, \pi_y) \in \mathcal{X}^m \times \mathcal{Y}, L_{\mathcal{A}}(f; \mathbf{x}, \pi_y) \in \mathcal{X}^m \times \mathcal{X}$ [0,1], then with probability at least $1-\delta$, the following inequality holds:

$$\sup_{f \in \mathcal{F}} (R_{L_{\mathcal{A}}}(f) - \widehat{R}_{L_{\mathcal{A}}}(f;S)) \le 2C_{\mathcal{A}}(\varphi)N(\varphi)\widehat{\mathcal{R}}(\mathcal{F}) + \sqrt{\frac{2\log\frac{2}{\delta}}{n}}, \quad (7.18)$$

where $\widehat{\mathcal{R}}(\mathcal{F})$ is the Rademacher average of the scoring function class (for the linear scoring function, we have $\widehat{\mathcal{R}}(\mathcal{F}) \leq \frac{2BM}{\sqrt{n}}$); $N(\varphi) =$ $\sup_{x\in[-BM,BM]}\varphi'(x)$ measures the smoothness of the transformation function φ ; and $C_{\mathcal{A}}(\varphi)$ is an algorithm-dependent factor.

The expressions of $N(\varphi)$ and $C_{\mathcal{A}}(\varphi)$ for ListNet and ListMLE, with respect to three representative transformation functions,⁵ are listed in Table 7.1.

From Theorem 7.5, one can see that when the number of training queries n approaches infinity, the query-level generalization bound will converge to zero at a rate of $O(\frac{1}{\sqrt{n}})$. Furthermore, by comparing the query-level generalization bound for different listwise ranking

⁵ The three transformation functions are:

[•] Linear Functions: $\varphi_L(x) = ax + b, x \in [-BM, BM].$

[•] Exponential Functions: $\varphi_E(x) = e^{ax}, x \in [-BM, BM]$. • Sigmoid Functions: $\varphi_S(x) = \frac{1}{1+e^{-ax}}, x \in [-BM, BM]$.

Table 7.1 $N(\varphi)$ and $C_A(\varphi)$ for listwise ranking algorith	able 7.1 $N(\varphi)$ and	$d C_{\Delta}(\varphi)$ for	listwise ranking	algorithms.
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φ	$N(\varphi)$	$C_{ ext{ListMLE}}(arphi)$	$C_{ ext{ListNet}}(arphi)$
φ_L	a	$\frac{2}{(b-aBM)\left(\log m + \log \frac{b+aBM}{b-aBM}\right)}$	$\frac{2m!}{(b-aBM)\left(\log m + \log\frac{b+aBM}{b-aBM}\right)}$
φ_E	ae^{aBM}	$rac{2e^{aBM}}{\log m + 2aBM}$	$\frac{2m!e^{aBM}}{\log m + 2aBM}$
φ_S	$\frac{ae^{aBM}}{(1+e^{-aBM})^2}$	$rac{2(1+e^{aBM})}{\log m + aBM}$	$\frac{2m!(1+e^{aBM})}{\log m + aBM}$

algorithms, and with regards to different transformation functions, one can have the following observations:

- The query-level generalization bound for ListMLE is much tighter than that for ListNet, especially when m, the length of the list, is large.
- The query-level generalization bound for ListMLE decreases monotonously, while that of ListNet increases monotonously, with respect to m.
- The linear transformation function is the best choice in terms of the query-level generalization bound in most cases.

7.4 **Discussions**

Since learning to rank is still a new research area, many theoretical issues are still left open and significant efforts are still needed to make it a legitimate branch of ML.

Actually, the full story of the statistical ranking theory should consist of two parts: the statistical consistency and the generalization ability of learning-to-rank methods. We have briefly reviewed the recent advances on generalization analysis in terms of a given surrogate loss function L, i.e., when the number of training queries approaches infinity, whether the empirical risk defined with the surrogate loss L can converge to the expected risk, which is also defined with the same surrogate loss L. The statistical consistency further discusses whether the minimization of the expected risk defined with the surrogate loss L can lead to the minimization of the expected risk defined with the true loss (which is not clearly defined for ranking yet).

Although there has been some work discussing the consistency for ranking in the literature, the problem has yet been adequately solved. For example, the consistency for bipartite ranking was studied in [28], which is, however, more like a classification problem and much simpler than real ranking problems. For another example, in [27], the consistency for pairwise ranking was discussed, but no necessary discussions were provided on how to extend this result to listwise ranking. In [33], the consistency issue with respect to DCG was studied. However, the authors only discussed the regression loss and thus their work is insufficient to explain the majority of learning-to-rank algorithms that are based on pairwise or listwise loss functions. In [129], the consistency of listwise ranking was investigated. However, the true loss is defined as a 0-1 loss at the permutation level, which is clearly not in accordance with our understanding of the loss in ranking (e.g., errors at different positions should lead to different losses). Therefore, all the aforementioned work cannot be used to well explain the consistency issue in learning to rank for IR.

To move forward, it is very important to define a reasonable true loss for ranking. One choice is to directly use the IR evaluation measures to define the true loss. However, there are still several problems with this. For example, there are many different IR evaluation measures, and it is not clear which one should be regarded as the true loss. Since these measures are not necessarily consistent with each other, no matter which one we choose as the true loss, it looks not that "true" after all. Second, according to the practice of classification, the true loss is usually not what we use to evaluate the performance of a learning algorithm. For example, while the 0-1 loss is used as the true loss of classification, precision, recall and F-scores are widely used as evaluation measures. Based on these discussions, it is not clear yet how to define the true loss for ranking, but it is relatively easy to get some principles of defining such a true loss. For example, the true loss for ranking should be defined at the query level and should consider the position. This could be future work of the theoretical study on learning to rank.

Summary and Outlook

In this tutorial, we have mainly introduced three approaches to learning to rank. The first is called the pointwise approach, which reduces ranking to regression, classification, or ordinal regression on each single document. The second is called the pairwise approach, which basically formulates ranking as a pairwise classification problem. The third is called the listwise approach, which regards ranking as a new problem, and tries either to directly optimize the non-smooth IR evaluation measures, or to minimize listwise ranking losses. We have introduced the representative algorithms of these three approaches, discussed their advantages and problems, and validated their empirical effectiveness on the LETOR benchmark dataset. In addition, we have also introduced the statistical ranking theory, and analyzed the query-level generalization ability of several learning-to-rank methods.

As a summary, we plot the representative algorithms introduced in this tutorial in Figure 8.1. From the figure, one can find that learning to rank for IR has become hotter and hotter in recent years, and more and more attention has been paid to the listwise approach.

Note that this tutorial is by no means a complete review of the area of learning to rank, especially considering that this is still a

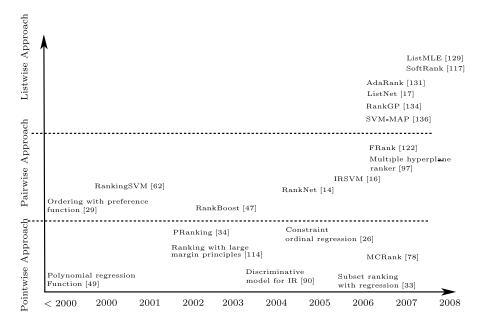


Fig. 8.1 Learning to rank algorithms.

developing research area. There is much other work that has not been covered, some of which cannot be easily categorized into the three approaches:

- Ground truth mining [3, 68, 105], which targets automatically mining ground truth labels for learning to rank, mainly from click-through logs of search engines.
- Feature engineering [51], which includes feature selection, dimensionality reduction, and effective feature learning.
- Query-dependent ranking [52, 71], which adopts different ranking models for different types of queries, based on either hard query type classification or the soft nearest neighbor based approach.
- Supervised rank aggregation [80], which learns the ranking model not to combine features, but to aggregate candidate ranked lists.

- Semi-supervised/active ranking [5, 40, 67, 106, 135], which leverages the large number of unlabeled queries and documents to improve the performance of ranking model learning.
- Relational/global ranking [38, 103, 104, 107, 137], which does not only make use of the scoring function for ranking, but also considers the inter-relationship between documents to define more complex ranking models.
- Other learning-to-rank algorithms [15, 19, 32, 93, 109, 127, 142, 143, 144] that are based on association rules, decision systems, and other technologies; other theoretical analysis on ranking [50]; and applications of learning-to-rank methods [87, 128].

At the end of this tutorial, let us come back to the questions we raised in the introduction, and give their possible answers.

(1) To what respect are these learning-to-rank algorithms similar and in which aspects do they differ? What are the strengths and weaknesses of each algorithm?

The answer can be found by the algorithm description and categorization. Basically all the algorithms belonging to the pointwise approach reduce ranking to either regression, classification, or ordinal regression. Almost all algorithms belonging to the pairwise approach reduce ranking to a pairwise classification. The advantage of these two approaches is that many existing theories and tools in ML can be directly applied. However, distinct properties of ranking have not been considered in such formulations. For example, most IR evaluation measures are query-level and position-based. However, neither the query information nor the position information is visible to the loss functions of these two approaches. The listwise approach instead treats ranking as a new problem, and defines specific algorithms for it. It can better leverage the concept of query, and consider the position information in the ranking results when training the ranking model. The problem with the listwise approach is that it is in general more complex than the pointwise and pairwise approaches. Furthermore, a new theoretical foundation is needed to explain the behaviors of listwise ranking algorithms.

According to the analysis in Section 5, the loss functions of most learning-to-rank methods, no matter pointwise, pairwise, or listwise, are upper bounds of (1 - NDCG). Therefore, the minimization of these loss functions can lead to the minimization of (1 - NDCG), or the maximization of NDCG.

(2) Empirically speaking, which of those many learning-to-rank algorithms perform the best? What kind of datasets can be used to make a fair comparison among different learning-to-rank algorithms?

According to the discussions in Section 6, the LETOR benchmark dataset has recently been widely used. Due to the standard data collection, feature representation, dataset partitioning, and evaluation tools in LETOR, it is possible to perform fair comparisons among different learning-to-rank methods. Empirical studies on LETOR have shown that the listwise ranking algorithms seem to have certain advantages over other algorithms, especially for top positions of the ranking result, and the pairwise ranking algorithms seem to outperform the pointwise algorithms. These results are in accordance with the discussions in this tutorial. However, as pointed out in Section 6, these experimental results are still primal and by carefully tuning the optimization process, the performance of every algorithm can be further improved.

(3) Theoretically speaking, is ranking a new ML problem, or can it be simply reduced to existing ML problems? What are the unique theoretical issues for ranking that should be investigated?

According to the discussions in Section 7, we can clearly see that it is better to regard ranking as a new ML problem, rather than reducing it to existing problems. Unique properties of ranking for IR as compared to classification and regression lie in that the evaluation of a ranking model is performed at the query level and is position based. Therefore, the risks should be defined at the query level as well, and a query-level theoretical framework is desired for conducting analyses on learning-to-rank methods. Furthermore, the "true loss" for ranking should consider the position information in the ranking result, but not as simple as the 0–1 loss in classification.

(4) Are there many remaining issues regarding learning to rank we should study in the future? What are they?

As mentioned previously, there are still many open problems. We list some of them as follows, as future work on learning to rank.

Learning from Logs

• The existing benchmark datasets are all of relatively small scales. From a ML point of view, hundreds of gueries cannot reasonably guarantee the effectiveness of a learning-torank algorithm. Developing realistically sized datasets is very important. Click-through log mining is one of the possible approaches to construct large-scale training data. Some work has been done along this direction [3, 68, 105], however, they also have certain limitations. Basically, the work has tried to map the click-through logs to pairwise preferences or multiple ordered categories. However, this process is not always necessary (and sometimes even not reasonable). As we know, the multiple ordered categories are designed for human labelers, which cannot cover all the rich information contained in the click-through logs, e.g., the user sessions, the frequency of clicking a certain document, the frequency of a certain click pattern, and the diversity in the intentions of different users. If converting the log data to multiple ordered categories or pairwise preferences, the information will be missing. Therefore, it is meaningful to reconsider the problem, and probably change the learning algorithms to adapt to the log data. For example, one can regard the click-through logs (without mining) as the ground truth, and define the loss function based on its likelihood.

Feature Engineering

• After one extracts a set of features for each document, it seems the learning-to-rank problem becomes a standard prediction task. However, one should notice that ranking is deeply rooted in IR, so the eventual goal of learning to rank

is not only to develop a set of new algorithms and theories, but also to substantially improve the ranking performance. For this purpose, feature engineering cannot be overlooked. It is a killer aspect whether we can encode the knowledge on IR accumulated in the past half a century in the extracted features. Currently, this kind of work has not been given enough attention. In the future, we should study the possibility of learning effective features.

Advanced Ranking Methods

- In most existing learning-to-rank algorithms, a scoring function is used for the sake of simplicity and efficiency. However, sometimes such a simplification cannot handle complex ranking problems. People have made some attempts on leveraging the inter-relationships between objects and some relational (global) ranking algorithms [103, 104, 107] have been proposed. However, this is not yet the most straightforward way of defining the hypothesis for ranking, especially for the listwise approach. Since the output space of the listwise approach is composed of permutations of documents, the ranking hypothesis should better directly output permutations of the documents, rather than output scores for each of the individual documents. In this regard, defining the ranking hypothesis as a multi-variate function directly on permutations could be a future research topic. Note that the task is challenging because permutation-based ranking functions can be very complex due to the extremely large number of permutations, but we think it is worthy and also possible to find efficient algorithms to deal with this situation.
- Motivated by the recent advances in ML, one should expect corresponding progress in learning to rank. For example, transfer ranking, multi-task ranking, semi-supervised ranking, and active ranking can all be promising future research topics. However, when performing such research, one should pay attention to the unique properties of ranking, and make

necessary adaptions when introducing these concepts and technologies. For example, we have mentioned some previous work on semi-supervised ranking [40, 67, 5]. Basically, the work simply borrows some concepts and algorithms from semi-supervised classification. However, the validity of doing so needs further checking. For instance, since similarity is essential to classification ("similar documents should have the same class label"), it looks very natural and reasonable to propagate labels cross similar documents. However, in ranking, similarity does not play the same central role. It seems that preference is more fundamental than similarity. Then it is questionable to still conduct similarity based label propagation for semi-supervised ranking. Furthermore, in classification, if we do not have class labels, we know nothing about the objects. However, in ranking, even if we do not have ground truth labels, we still have several very strong rankers, such as BM25 and language models for IR, which can give us a relatively reasonable guess on which document should be ranked higher. Therefore, we can assume that we have some knowledge about the unlabeled data. If we can incorporate such knowledge into the semi-supervised ranking process, we may have the chance to do a better job.

 As introduced in this tutorial, most efforts on learning to rank have been given to discriminative learning. However, as we notice, generative learning is also a very important branch of ML. There is no reason that generative learning cannot be used in ranking. This could be a promising research direction of learning to rank, from both algorithmic and theoretical points of view.

$Ranking\ Theory$

• Regarding existing algorithms directly optimizing IR evaluation measures, although a theory has been proposed to quantify the "directness" of some algorithms in the large sample limit, it is not clear how such algorithms will perform on the

- test data when the training data is limited. This could also be a future research direction.
- As compared to the efforts on algorithms, the theoretical work on ranking is not yet sufficient. For example, it is still unclear about the "true loss" in ranking, and in addition to the generalization analysis, statistical consistency and fast convergence rate have not been adequately studied. Furthermore, some fundamental questions have not been answered with regards to the ranking theory, such as the complexity of the function class in ranking. This could also be an important research direction for learning to rank.

Overall, this tutorial is just a stage-wise summary of this hot research field. Given the fast development of learning to rank, we can foresee that many new algorithms and theories will appear in the future. We hope that this tutorial can motivate more people to work on learning to rank, so as to make this research direction more impactful in both the IR and ML communities.

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