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Structure-Tags Improve Text Classification for Scholarly Document Quality Prediction

Gideon Maillette de Buy Wenniger†, Thomas van Dongen†, Eleri Aedmaa‡, Herbert Teun Kruitbosch†, Edwin A. Valentijn§, and Lambert Schomaker†

†Bernoulli Institute for Mathematics, Computer Science and Artificial Intelligence
University of Groningen, Groningen, The Netherlands
gmdbw AT gmail.com t.a.van.dongen AT student.rug.nl
‡Center for Information Technology, University of Groningen, Groningen, The Netherlands
§Kapteyn Astronomical Institute, University of Groningen, Groningen, The Netherlands

Abstract

Training recurrent neural networks on long texts, in particular scholarly documents, causes problems for learning. While hierarchical attention networks (HANs) are effective in solving these problems, they still lose important information about the structure of the text. To tackle these problems, we propose the use of HANs combined with structure-tags which mark the role of sentences in the document. Adding tags to sentences, marking them as corresponding to title, abstract or main body text, yields improvements over the state-of-the-art for scholarly document quality prediction. The proposed system is applied to the task of accept/reject prediction on the PeerRead dataset and compared against a recent BiLSTM-based model and joint textual+visual model as well as against plain HANs. Compared to plain HANs, accuracy increases on all three domains. On the computation and language domain our new model works best overall, and increases accuracy 4.7% over the best literature result. Applications include (short) text classification (Rao and Spasojevic, 2016), entailment (Rocktäschel et al., 2015) and neural machine translation (MT) (Bahdanau et al., 2014; Luong et al., 2015). Newer attention-based models, particularly the transformer model (Vaswani et al., 2017) are even more apt at using all of the possible context when encoding sentences, further improving performance. Transformers are also used to build general sentence embeddings with the BERT model (Devlin et al., 2018). In comparison, the accurate classification of full documents remains challenging. To be effective, a deep learning model for longer text should fulfill the following three criteria:

1. Trainability: being trainable on long texts.
2. Computational efficiency: efficiency as well as parallelizability, to effectively use GPUs.
3. Rich context: having access to rich context at sentence and document level. And avoiding therefore: 1) the assumption that sentences at different locations are independent, 2) the even more crippling assumption of statistical independence of document words.

Plain sequential models such as RNNs and LSTMs model text as unstructured word sequences. This causes problems on longer texts because of the vanishing gradient and exploding gradient problem (Pascanu et al., 2013), which hampers trainability. Gradient bounding methods including gradient clipping (Hochreiter, 1998), can help to reduce these problems, but provide no solution for docu-
ments with thousands of words. Transformers and BERT are not a good match for long texts either, as these models have a computational cost that grows quadratically with sentence length. Arguably, bag-of-word models, including models performing average pooling over word embeddings accomplish trainability and computational efficiency. However, their computational cheapness is achieved at the price of making very strong statistical independence assumptions that harm prediction quality.

A group of models does fulfill all three criteria: hierarchical versions of sequential models, in particular hierarchical attention networks (HANs) (Yang et al., 2016). HANs produce hierarchical text encodings using a hierarchical stacking of LSTMs with attention, for the sentence and text level. This massively increases parallelization while simultaneously reducing the number of steps the gradient signal needs to be back-propagated during training, increasing learnability. HAN text encodings can still take much context into account at every level in the representation, thanks to the use of LSTMs.

While HANs are highly effective in forming adequate representations of longer texts, they are still deficient in the use of structure information inherent in the text. The reason is simple: these models have only one ((Bi)LSTM) encoding sub-model per level in the hierarchy. This sub-model is used to encode all the inputs at that level, without access to relevant structure context. In this work we observe that this problem can be tackled by adding XML-like structure-tags at the beginning and end of each input sentence. The effectiveness of our approach is demonstrated on two tasks:

A Paper accept/reject prediction on the PeerRead dataset (Kang et al., 2018).

B Number of citations prediction for scholarly documents, on a new dataset with 88K articles compiled from the Allen AI S2ORC dataset.

The experiments for both tasks show that using just three tags to mark abstract, title and body text, already provides substantial improvements: A) outperforming all models on the computation and language domain and HAN without tags on all domains, B) outperforming all other models. Larger gains can likely be made by further enriching the tag-set. The proposed tagging approach is particularly useful in the domain of scholarly document understanding, since while these document are typically long, they are also highly structured.

The rest of the paper is structured as follows. In section 2 we discuss the various existing and alternative NLP models for the aforementioned tasks of quality prediction. Section 3 describes the proposed HAN model combined with structure-tags. Section 4 and 5 respectively discuss their use for accept/reject and number of citations prediction.

2 Related Work

Multiple methods have been proposed to estimate the quality of scientific papers. The most common approach is to use the citation counts as a measure of quality, to be predicted by models. Fu and Aliferis (2008) proposed one of the first models which used both the papers content in the form of the paper title, abstract and keywords as well as bibliometric information. Notably they used automated scripts to retrieve bibliometric information, even so their final corpus is still relatively small, containing 3788 papers.

Limited recent research is available on the subject of predicting the quality of papers with deep learning using the textual content. Shen et al. (2019) combine visual and textual content using a CNN and LSTM respectively. The authors make use of the Wikipedia and the arXiv datasets. The authors propose a joint model that classifies the quality of papers. To generate textual embeddings, the authors use a bi-directional LSTM model similar to the one proposed by the same authors in (Shen et al., 2017). The input to the model is the word embeddings of a paper, obtained using GloVe, and the output is a textual embedding.

Some recent work focuses on predicting the number of citations from the paper text augmented with review text. To do so, Li et al. (2019) created a dataset of abstracts and reviews from the ICLR and NIPS conferences: 1739 abstracts with a total of 7171 reviews for ICLR and 384 abstracts with 1119 reviews for NIPS. Plank and van Dale (2019) collect a dataset of 3427 papers with 12260 reviews. Both papers show improvement in the results from using the review information.

Hierarchical sequential models

Hierarchical versions of sequential models have already been pioneered in the literature for a long time in the form of hierarchical RNNs (Hihi and Bengio, 1996). More recently however, use of LSTMs instead of RNNs and use of attention resulted in the now popular HAN model (Yang et al., 2016), which was successfully applied to sentiment
Figure 1: Most important models compared in this work.

Adding structure through additional inputs
Our proposed structure-tag framework resembles the approach that been used for neural MT translating multiple source languages to multiple target languages using a unified model (Johnson et al., 2016), in which a special “command token” is used to indicate which kind of translation is desired. Related also is the idea of using multiple embeddings for different types of information, as introduced in neural MT by Sennrich and Haddow (2016), which was later also exploited in the transformer model (Vaswani et al., 2017). In contrast to the latter approaches which change the embedding layer, like (Johnson et al., 2016) we leave the (HAN) model exactly as is and only change the input.

3 Models
In this work we use and refine state-of-the-art text-based deep learning models for text classification and regression tasks: accept/reject prediction and number of citations prediction respectively. Our contributions focus on HANs, which we show for these tasks to be competitive with models that use a flat BiLSTM encoder at their core (Shen et al., 2019). Figure 1a shows a diagram of our HAN model with structure-tags added to the input, and

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1Our HAN implementation is adapted from https://github.com/cedias/Hierarchical-Sentiment

Figure 1b shows a diagram of the BiLSTM-based model, our baseline for comparison. As can be seen from the diagrams, both models use a BiLSTM at the text level that works on embeddings computed for the sentences of the text. However, while HAN uses the sequential order to compute an embedding, the baseline model averages word vectors, disregarding order, similar to bag-of-word representations. We also use a second baseline model: Average Word Embeddings (AWE), which simply encodes text by the average word embedding.

3.1 Sentence type tags for more structure
The hierarchical structure of text characterized by structure elements such as sections, paragraphs and sentences and labeling elements such as document titles and section titles reveals important information. Models without hierarchy such as plain RNN/LSTM models ignore this structure, which motivated HAN. HAN uses an LSTM with attention to create encodings of each sentence separately and combines this with a second LSTM with attention on top to transform these into an encoding of the entire text. The hierarchical structure of HAN provides several advantages over flat sequential models, i.e. plain RNNs/LSTMs:

1. Trainability on long texts: using less steps for back-propagating gradients during training, HAN can process longer texts without running into vanishing/exploding gradient problems.
Abstract

We present an algorithmic framework for learning multiple related tasks.

Introduction

When two NLP systems are run on the same data, we expect certain constraints to hold between their outputs.

Figure 2: Example of structure-tags for a paper from in the PeerRead computation and language arXiv dataset.

HAN preserves high-resolution when forming sentence-level encodings.

2. Computational efficiency: the structure of HAN makes computations better parallelizable, since its sentence encoding LSTMs can process their inputs in parallel.

3. Interpretability of predictions: visualizing HAN attention facilitates some qualitative insight into what inputs are important for making predictions at a sentence and word level.

Despite these large advantages, HAN in its normal application still remains limited in its use of structure. In particular, while HAN encodes sentences in a hierarchical way, it does so while using the same LSTM encoder for every sentence without structure context. In this work we introduce a way to overcome these problems by adding sentence type tags encoding the role of a sentences or other information, which is then directly available to the BiLSTM when encoding the sentences. This is illustrated in Figure 2. First the input is segmented into a list of sentences, as is also done in preprocessing for regular HAN. Then the role of each sentence is added at the beginning and end of each sentence. In our current experiments the roles are restricted to three options: TITLE, ABSTRACT, BODY_TEXT, however, the idea is general enough to include much more specific tags as well as tags encoding relative or absolute sentence position information; to be explored in future work. We will refer to this system as hierarchical attention network with structure tags (HAN_ST). The advantages of the tag-base approach over other possible solutions, such as using different BiLSTMs for different types of sentences are simplicity and scalability. Equally important, using tags allows the BiLSTM to only specialize its functioning to specific types of sentences where needed, while effectively sharing what can be generalized independent of sentence type.

4 Accept/Reject prediction on PeerRead

The first scholarly document quality prediction task we test our methods on is accept/reject prediction on arXiv papers from the PeerRead dataset (Kang et al., 2018). This dataset is chosen because of the large amount of earlier work in the literature reporting results on it, allowing comparison against the state-of-the-art on a well studied task.

The full PeerRead dataset holds 14784 papers in total, each of which contains implicit or explicit accept/reject labels. Furthermore, PeerRead contains different subsets of papers. The largest subset consists of arXiv papers (11778) in three computer-science sub-domains: machine learning (cs.LG), computation and language (cs.CL), artificial intelligence (cs.AI), and has only accept/reject labels; this is the dataset that we use. A part of the papers also include reviews (3006 papers) and a subset of the latter also contains aspect scores (586 papers). However, of these papers with reviews, the large majority is from NIPS (2420 papers), and those papers are all accepted. As the arXiv portion is relatively larger, and accept/reject labeled, most work has focused on the task of accept/reject prediction for the papers in this set.

Table 1 shows the sizes of the different subsets of the arXiv PeerRead dataset and their respective division in number of accept and reject examples. Note that this division is imbalanced for each of the three domains, with the least imbalance for the machine learning subset and the most imbalance for the artificial intelligence subset, in which around 90% of the examples is rejected. These imbalances in the number of examples for each of the classes make learning harder, but can be partly overcome by using strategies such as re-sampling.

4.1 Experimental Setup

In our experiments we tried to stay close to the experimental setup used by (Shen et al., 2019), while deviating from their settings when necessary. We used PyTorch for our code and a single GeForce RTX 2080 Ti GPU for our experiments. Table 2 gives an overview of the used hyperparameters that are shared across experiments, as well as the hyperparameters that are specific to the accept/reject
Table 1: Data sizes and division between the ratio of accepted and rejected papers for the arXiv subsets.

<table>
<thead>
<tr>
<th>Subset</th>
<th>Training</th>
<th>Validation</th>
<th>Testing</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>num</td>
<td>acc:rej</td>
<td>num</td>
<td>acc:rej</td>
</tr>
<tr>
<td>machine learning</td>
<td>4543</td>
<td>36.4% : 63.6%</td>
<td>252</td>
<td>36.5% : 63.5%</td>
</tr>
<tr>
<td>computation &amp; language</td>
<td>2374</td>
<td>24.3% : 75.7%</td>
<td>132</td>
<td>22.0% : 78.0%</td>
</tr>
<tr>
<td>artificial intelligence</td>
<td>3682</td>
<td>10.5% : 89.5%</td>
<td>205</td>
<td>8.3% : 91.7%</td>
</tr>
</tbody>
</table>

Table 2: Hyperparameters used in the experiments.

<table>
<thead>
<tr>
<th>Feature</th>
<th>PeerRead classification</th>
<th>S2ORC regression</th>
</tr>
</thead>
<tbody>
<tr>
<td>optimizer, learning rate</td>
<td>Adam, 0.005</td>
<td></td>
</tr>
<tr>
<td>maximum input characters</td>
<td>20000</td>
<td></td>
</tr>
<tr>
<td>vocabulary size</td>
<td>10000</td>
<td></td>
</tr>
<tr>
<td>weight initialization general</td>
<td>Xavier uniform</td>
<td></td>
</tr>
<tr>
<td>lstm bias</td>
<td>XGtav uniform</td>
<td></td>
</tr>
<tr>
<td>word embeddings</td>
<td>GloVe</td>
<td></td>
</tr>
<tr>
<td>loss function</td>
<td>cross entropy</td>
<td>MAE</td>
</tr>
<tr>
<td>dropout probability</td>
<td>0.5</td>
<td>0.2</td>
</tr>
<tr>
<td>BiLSTM hidden size</td>
<td>256</td>
<td>100</td>
</tr>
<tr>
<td>batch size</td>
<td>4</td>
<td>64</td>
</tr>
<tr>
<td>embedding size</td>
<td>50</td>
<td>300</td>
</tr>
</tbody>
</table>

Table 3: Total trainable parameters per model.

<table>
<thead>
<tr>
<th>Task</th>
<th>AWE</th>
<th>BiLSM</th>
<th>HAN/HANST</th>
</tr>
</thead>
<tbody>
<tr>
<td>PeerRead</td>
<td>500202</td>
<td>1657222</td>
<td>3235206</td>
</tr>
<tr>
<td>citation prediction</td>
<td>3000901</td>
<td>3402801</td>
<td>3644801</td>
</tr>
</tbody>
</table>

Learning rate 0.0001 gave poor results in our experiments.

Using the full text as input is in theory preferred over using only selected text, in order not to lose information prematurely. In practice however, this is not feasible with high resolution deep learning models such as HANs, which take input that starts at the word level. To save memory and computation, models may instead start out from the sentence level, using embeddings directly as inputs. But with simple sentence embeddings, this leads to a substantial loss of input information, which may hamper performance. Even so, (Shen et al., 2019) apply this strategy in a basic way by computing the average word embedding for each sentence, and using a BiLSTM model on top of that. Nevertheless, they still use a limit on the input length, by allowing only a maximum of 350 sentences. With HAN, which uses more memory and computation-intense sentence-level encodings, limiting the input length is even more crucial. However, rather than limiting the number of sentences, we limited the maximum number of characters, set to 20000. We found that with HANs this gives better results, even though on average it corresponds to less words. This is explained by the distribution over the number of words per example for each of the two length cutoff policies, see Table 4. Fixing the number of sentences causes large variance in the number of words per example, likely caused by writing style differences across authors. In contrast, fixing the number of characters by definition assures a con-
stant input length, and hence a more constant number of words (which is proportional to number of characters). We believe this more constant amount of information in the input aides learning.

### 4.1.2 Results

Table 5 shows our best results on the PeerRead dataset, using HANST. The same table also shows the previous literature results of (Shen et al., 2019) and (Kang et al., 2018). Observe that in the computation & language domain, we gain 4.7% accuracy over the best of the these literature models (Joint), while on the machine learning domain and artificial intelligence domain datasets we lose 2.4% and 3.8% respectively in comparison to the best performing of the literature models on these domains (BiLSTM and Joint). In Table 6 we show the results for both our HAN models as well as for the other models. These results show a clear and consistent improvement from HANST over plain HAN: 1.5% accuracy for the computation & language domain and 2.1% for the machine learning domain and 0.7% for the machine learning domain. Table 6 also shows results for our own re-implementation of the BiLSTM model described by (Shen et al., 2019). This useful for comparison since we made some changes to the experimental setup, including the use of higher learning rate and use of re-sampling. We observe better results with our re-implementation of BiLSTM than in the original work for these datasets where re-sampling was helpful, showing its importance for imbalanced datasets. Statistical significance was tested with the exact two-sided McNemar’s test. Comparing to the AWE systems: At significance level 0.05, other systems improve significantly over AWE except for HAN on the machine learning domain. At significance level 0.01 improvements over AWE are only significant on the artificial intelligence domain for BiLSTM and HANST. Other differences are not statistically significant.

While HANST is competitive with the literature models on PeerRead, it benefits from larger training data, as is available for the task of number of citations prediction.

### 4.1.3 Effects of reducing the label set

To determine the importance of different structure tags, in particular the title marking, we performed ablation experiments in which we reduced the label set. We combined the title and abstract label into one, leaving a structure-tag set with only two tags. Table 7 shows the results. As can be seen, the smaller structure-tag set reduces performance in comparison to HAN with three structure tags on all three domains. In the computation & language domain, the model performs worse also than HAN without structure tags on both accuracy and AUC, and in the artificial intelligence domain it performs equal in terms of accuracy but worse still on AUC. In the machine learning domain the model also loses performance over HAN with three structure tags, but still outperforms plain HAN. The results suggest that the titles of articles contain information.

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5The scores of the 3 runs for one system are combined at example level by taking the mode/average, i.e. simple voting.
Table 7: Results of the HAN_{ST} model with a reduced structure-tag set of only two tags.

<table>
<thead>
<tr>
<th>domain metric</th>
<th>artificial intelligence &amp; language</th>
<th>machine learning</th>
</tr>
</thead>
<tbody>
<tr>
<td>accuracy</td>
<td>89.6 ± 1.57%</td>
<td>77.2 ± 1.21%</td>
</tr>
<tr>
<td>AUC</td>
<td>0.610 ± 0.067</td>
<td>0.759 ± 0.017</td>
</tr>
</tbody>
</table>

5 Number of citations prediction

The second task we test our models on is number of citations prediction. A key advantage of this task is that large datasets can be obtained relatively easy by leveraging public sources such as the Semantic Scholar Database. In contrast, obtaining accept/reject labels in large quantities typically requires having an agreement with publishers, and even then because of legal problems, it is hard to obtain and publish such data.6

Yet, how useful it is to predict the number of citations? More specifically: is the number of citations of a paper predictive of its quality? Intuitively one would expect this to be the case at least to some extent. Figure 3 shows histograms of the numbers of citations of articles from the PeerRead datasets for accepted and rejected papers.7 While there are some differences between the two domains, the main trend is the same in both cases: for rejected papers, the counts are peaked around zero citations and quickly decrease to one or zero for high citation counts. In contrast, the number of citations for accepted papers is two to three times higher on average, depending on the domain. Finally, we formally computed correlation in the form of the Spearman rank-order correlation coefficient (ρ) and associated p-value for both domains. For both domains, the value of ρ is high and the p-value extremely close to zero, which indicates significant correlation can be concluded at all p-levels of significance for a two-sided test. These histograms and numbers prove that there is indeed a strong correlation between acceptance/rejection and the number of citations. Therefore it makes sense to consider the number of citations as an imperfect but nonetheless useful proxy for the quality of scholarly documents.

5.1 The dataset

Recent works undertake the task of number of citations prediction based on the scholarly document text, but mostly do so while using relatively small datasets. As discussed in related work, some of the recent work adds review text to the input. However, creating models using reviewer comments limits their practical application to after reviewing and reduces available training data. These observations motivated us to rather aim for a relatively large dataset of ⟨paper, number of citations⟩ pairs. We selected a subset of papers in the computer science domain from the S2ORC (Lo et al., 2020) data, for which title, abstract and body text information is present; these are combined as the example text. We did this for papers in the year range 2000–2010, and counted the number of citations of citing papers that are published within 8 years after the publication of a paper. Randomly ordering the papers, from this we compiled a dataset with in total about 88K papers, and statistics as shown in Table 8.8

Note that the new S2ORC-derived log-citation-count prediction dataset, used in our experiments, is available from: https://github.com/gwenniger/s2orc-cc/.

8The new S2ORC-derived log-citation-count prediction dataset, used in our experiments, is available from: https://github.com/gwenniger/s2orc-cc/.

9Due to a misunderstanding of the S2ORC data format, which actually does contain longer text when combining all the text dictionaries, which got clarified after submission.
Figure 3: Histograms and global statistics of number of citations for accepted and rejected papers for the sub-domains of PeerRead. Histograms are truncated on the right at 100 citations. The table in 3c shows the formal correlation measure: average numbers of citations and Spearman rank-order correlation in the different domains.

full paper text for future work. The labels added to the examples consist of a function of the number of citations, as explained next.

5.2 Citation-score as a quality proxy

The number of citations follow of scholarly documents follow a Zipfian distribution (Silagadze, 1997). That is, most papers have little citations, but those that obtain more citations tend to get exponentially more. To account for this, we used the log of the number of citations to create a metric that aims to approximate a measure of quality on a linear scale. In practice, we use the function:

\[
\text{citation-score} = \log_e(n + 1)
\]

adding one to the number of citations \(n\) before taking the log, to make sure the function is well-defined even for papers with zero citations.

Comparison to alternative citation scores

What alternatives to our log-based metric have been explored in the literature? Li et al. (2019) map citation counts to the \([0,1]\) range, presumably by simply scaling them after the paper with the maximum and minimum number of citations in a dataset have been determined. But this approach transfers poorly to new data, since as the number of citations follows the Zipfian distribution, still higher citation counts in unseen data are likely. Furthermore, because of the Zipfian nature of the number of citations, this transformation will map the citation score of many papers to a number close to zero, drastically inflating the evaluation scores of predictions for this citation score. A better approach is to discretize the number of citations into a fixed number of ranges. To predict the impact of scientific papers, Plank and van Dale (2019) discretize time-normalized citation statistics into low, medium and high impact papers based on a boxplot and outlier analysis. In comparison however, our approach does not require discretization/binning, which has
advantages: 1) not committing to a fixed resolution, 2) avoiding problems for papers with a number of citations on the border of two bins, 3) allowing the predicted scores to be deterministically transformed back to a number of citations.

5.3 Loss function and evaluation metrics

The next important question is what loss function we should optimize when training our networks to predict the chosen citation score (1). Whereas mean-squared-error (MSE) is the default choice for regression problems, we found this loss function to perform poorly in combination with our score. In contrast, preliminary experiments showed that mean-absolute-error (MAE) facilitates effective and relatively stable optimization, so we decided to use this. Another important question is the choice of quality metrics. MSE and MAE are standard metrics for regression evaluation, so we report those. Additionally, we report the $R^2$ score, denoting the proportion of the variance in the dependent variable that is predictable from the independent variable(s), defined as:

$$R^2 = 1 - \frac{FVU}{\text{var}[Y]} = 1 - \frac{\text{MSE}(Y, Y')}{\text{var}[Y]}$$

(2)

With $Y'$ and $Y$ being the predicted and actual labels respectively, MSE being the mean-squared-error and $FVU$ the fraction of variance unexplained. This explains how the $R^2$ score normalizes for the relative difficulty for the task, by dividing by the variance of the labels in the test set. Another interpretation is that the $R^2$ score normalizes by the error obtained when always predicting the average of the test labels. Consequently, a $R^2$ score larger than 0 means performance better than this baseline, and below 0 means worse. This avoids the need to add scores for this baseline as reference, making the $R^2$ score more directly interpretable than MSE or MAE. As such, unlike the other metrics the $R^2$ score is also meaningfully comparable across datasets, which typically differ in test set variance.

5.4 Number of citations prediction results

Table 9 shows the results of our models trained on our new S2ORC number of citations prediction dataset. We observe that the HAN$_{ST}$ model outperforms the other models. Furthermore, the improvements of HAN$_{ST}$ over BiLSTM and AWE is statistically significant (wilcoxon signed-rank test), with p-value 0.008 in both cases.

6 Conclusion

This work showed the usefulness of HAN and rich context tags to the processing of scientific documents. Consistent improvements in prediction quality were obtained for both accept/reject estimation and number of citations prediction for HAN when adding structure-tags. A strong and significant correlation between accept/reject labels and number of citations was demonstrated, signaling the usefulness of the latter as a measure of scholarly document quality. With more training data, as available on the citation-score prediction task, HAN with structure-tags outperforms the strong and recently proposed scholarly document quality prediction models that we compared to in this study.

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