Research Paper Summarizer

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Abstract

In this report, we aim to improve accessibility of information presented in complex research papers by using Natural Language Processing (NLP) methods to automatically generate summaries of research papers that are easier to read than the average technical abstract. A dataset of research papers in the medical field which have a technical abstract as well as a plain language summary associated with each paper was used to finetune a pretrained T5 model. The research paper and the technical abstract were used as inputs to the model, and the plain language summary was used as the target output. The finetuned model was evaluated using the ROUGE metric for summarization as well as various readability metrics. Our code can be found here.

1 Introduction

Research papers are groundbreaking in terms of presenting new findings, theories, and perspectives, making them essential for scientific breakthroughs. However, the knowledge presented in research papers is often inaccessible to the general public for various reasons. Reasons include putting research papers behind paywalls, using dense technical language that the average person cannot understand, and sometimes simply being far too long. This discourages those in the general public who may be interested in the topic of the research paper or those who would like to enter the research area presented in the paper from reading the paper. Though the first issue is not one that can be solved without a change in the current publishing system, the latter two issues can be addressed by using Natural Language Processing (NLP) methods to automatically generate summaries of research papers that are easier to read than the average technical abstract. This would involve finetuning a pretrained model on a dataset to convert the paper and its technical abstract into a plain language summary, which, to some extent, would take down the barrier of technical language and length.

2 Dataset and Resources

2.1 Dataset

To train and evaluate the model, a dataset of research papers in the biomedical domain was used. This dataset contains 28,124 research papers, each with an associated technical abstract and a plain language summary. The technical abstract is the abstract that is typically found in research papers, but the plain language summary is a summary that is written by the author, where they are required to highlight how the work fits into a broader context in a simple manner without complex acronyms and terminology (Luo et al., 2022). The dataset was split into the train set, the validation set, and the test set, with 26,124, 1,000, and 1,000 papers respectively. The dataset can be found here.

2.2 Model Resources

To obtain the resources to load our pretrained model and finetune it, HuggingFace was used, which is an online hub where users and/or companies are able to publish their trained PyTorch or TensorFlow models for others to use. Hugging-Face provides a high-level API that allows us to load the model, load the dataset in a proper format, and train the model without worrying about the inner workings of the model.

In order to train the model efficiently, training on a local machine would not suffice. For this purpose, Google Colab was used, where the access of a GPU is provided for free for some limited amount of time.

3 Methodology

3.1 Model Selection

For the summarization task, a couple of model architectures were researched and explored. The final architecture that was chosen was an encoder-decoder transformer model; namely the T5 (Text-to-Text Transfer Transformer) model (Raffel et al., 2020). This model was pretrained on a mixture of self-supervised and supervised tasks over a large text corpus, which made it a well-performing model for a variety of tasks out of the box. All that had to be done was the prepend the input with the given task (e.g., for summarization: "summarize:", for translation: "translate English to German:"). The model was chosen because of its ease of use as well as its performance on the general summarization task.

3.2 Model Finetuning/Training

To finetune the model, first the data had to be prepared. The initial approach was to use the LongT5 model, which was specialized for tasks with long inputs, and provide as much of each research paper as possible as the input to the model (Guo et al., 2022). However, due to the limited resources provided by Google Colab, it became apparent that using whole research papers as the input to the model would not be possible without running out of GPU memory or truncating our inputs to the point that significant information was lost. Therefore, the approach was changed to use the base T5 model and provide the technical abstract as the input to hopefully create a readable summary.

The data was converted into input tokens that are digestable by the model using the Hugging-Face Tokenizer API. The model was then trained using the Hugging-Face Trainer API, which allows for easy training of a model on a dataset¹. The model was trained for 3 epochs, with a learning rate of 2e-5, a maximum input length of 1024 tokens, and a maximum output length of 256 tokens. The training for this would take around 5 hours in the Google Colab environment.

3.3 Summary Generation

Once the model was finetuned, it was used to generate summaries for the test set. Initially, the default parameters were used for the generation of the summaries; however, using these parameters led to the generation of summaries that were not very readable for many reasons. One main reason was the constant generation of repeated tokens as seen in the following example:

Schistosoma mansoni is one of the major tropical infectious diseases responsible for tropical infectious diseases caused by Schistosoma mansoni Schistosoma Mansoni.

This case happened quite often in the test set. To mitigate these issues, the following parameters were used for the generation of the summaries:

max_length=256, min_length=64,
no_repeat_ngram_size=2,
num_beams=4, early_stopping=True,
length_penalty=2.0.

Parameters like length_penalty and early_stopping prevented the model from generating summaries that were too long because the longer the summaries went, the more susceptible they were to becoming unreadable. The no_repeat_ngrame_size parameter was implemented to make sure that the model did not generate the same token over and over again. With the implementation of these parameters, the summaries generated by the model were far more readable and coherent.

4 Results and Evaluation

References

Mandy Guo, Joshua Ainslie, David Uthus, Santiago Ontanon, Jianmo Ni, Yun-Hsuan Sung, and Yinfei Yang. 2022. Longt5: Efficient text-to-text transformer for long sequences.

Zheheng Luo, Qianqian Xie, and Sophia Ananiadou. 2022. Readability controllable biomedical document summarization.

Colin Raffel, Noam Shazeer, Adam Roberts, Katherine Lee, Sharan Narang, Michael Matena, Yanqi Zhou, Wei Li, and Peter J. Liu. 2020. Exploring the limits of transfer learning with a unified text-to-text transformer.

¹Both of the API's can are found in the Transformers library: https://github.com/huggingface/transformers

A Group Work Split

For this project, Afnan took responsibility of formatting the data properly to be ingested by the model. He also took responsibility of training the model on Google Colab. Jaykumar took responsibility of generating the summaries for the test set, which also included tuning the parameters of the finetuned model to generate the most coherent model. He also worked on creating the demo for the class presentation. Both Afnan and Jaykumar worked on performing the evaluation of the summaries generated, and they both worked on the presentation and writing the