

# Lay It On Me. Generating Easy-to-Read Summaries for Non-Experts

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## Abstract

In this study, we present an extractive-abstractive lay summarization pipeline for biomedical papers aimed at generating accessible summaries for non-experts. To achieve this, we construct a sentence-level dataset optimized for maximizing ROUGE scores, utilizing both lay summaries and full articles. We employ a BERT-based classifier for identifying the most important sentences within each article. The extracted summaries are then input into two abstractive models, Clinical-Longformer and GPT-2, which paraphrase the summaries to enhance readability. We evaluate the performance of our models using the ROUGE metric, along with readability metrics such as Flesch-Kincaid Grade Level (FKGL), Gunning Fog Score, and Automated Readability Index (ARI). We find that a ROUGE-maximizing extractive summarization approach is effective for generating extractive summaries, with the Clinical-Longformer model achieving the best results for combined ROUGE and readability scores. Our approach demonstrates the potential for generating lay-friendly summaries of biomedical papers, bridging the gap between expert knowledge and public understanding.

## 1 Introduction

It can be challenging for individuals without expertise to comprehend scientific publications, particularly in biomedicine, where inaccuracies can directly impact health decisions (Islam et al. 2020). A possible remedy for this situation is to provide lay summaries, i.e., summaries in simpler terms, which are currently uncommon. Prior research on Automatic Text Summarisation (ATS) has neglected the biomedical domain owing to the absence of data (Chandrasekaran et al. 2020); however, two recently introduced datasets (PLOS and eLife) have emerged to tackle this issue (Goldsack et al. 2022).

## 2 Methods and Datasets

### 2.1 Dataset

The data we used is sourced from biomedical research articles in English published in the Public Library of Science (PLOS) and eLife (Goldsack et al. 2022). The datasets (Tables 1 and 2) contain technical abstracts and lay summaries written

by experts, which are part of BioLaySumm2023 shared task (Goldsack et al. 2023).

Dataset	Training	Validation
PLOS	24,773	1,376
eLife	4,346	241

Table 1: PLOS and eLife: number of articles

Dataset	Avg. Sentences	Avg. Tokens
PLOS	300	9,000
eLife	600	14,000

Table 2: PLOS and eLife: Dataset statistics

### 2.2 Extractor Network

Due to the extreme length of medical articles (e.g., eLife has an average of 600 sentences per article), it is not feasible to pass them directly as input to the abstractive models due to their limited maximum input size:

- i. **GPT-2** (Radford et al. 2019a): 1,024 tokens, and
- ii. **Clinical-Longformer** (Li et al. 2023): 4,096 tokens

To overcome this limitation, we use the BioClinicalBERT (Alsentzer et al. 2019) model, pre-trained on the MIMIC-III dataset (Johnson et al. 2016), to extract the most important sentences from the articles. For that purpose, we cast the extraction summarisation problem as supervised binary classification where the input is a sentence  $s$  and the output is a binary label indicating whether the sentence should be included in the summary  $c$  or not (i.e., 1 and 0, respectively). Due to the nature of the provided gold summaries (i.e., abstractive and lay), we generate our own sentence-level dataset by applying the ROUGE-maximisation technique (Zmandar et al. 2021; Nallapati, Zhai, and Zhou 2017) on the gold summaries and the whole articles. More formally, for each gold summary sentence  $s_i^k$ , we find the sentence  $s_j^k$  in article  $a_k$  that maximises the ROUGE-2 score between them. We then label  $s_j^k$  as 1 and the rest of the sentences in  $a_k$  as 0. Because the number of sentences in the articles is much larger than the number of sentences in the gold summaries:

- i. We base our extractive binary dataset on both eLife and PLOS data to maximise the number of training samples;

- ii. We further resolve the class imbalance problem by random under-sampling the majority class (i.e., 0) to match the number of samples in the minority class (i.e., 1);

Our final extractive dataset consists of 944,234 sentences with a completely balanced class distribution. Data is further split into 80-training, 10-validation and 10-testing datasets in a random stratified manner. We then fine-tune the extractive model with a batch size of 32 and a learning rate of  $2e-5$  following the guidance from BERT’s authors (Devlin et al. 2019) and find that the model starts to over-fit beyond 2 epochs (see Figures 1 and 2). We also report high F1 scores of 0.767 and 0.765 on the validation and test sets, respectively.

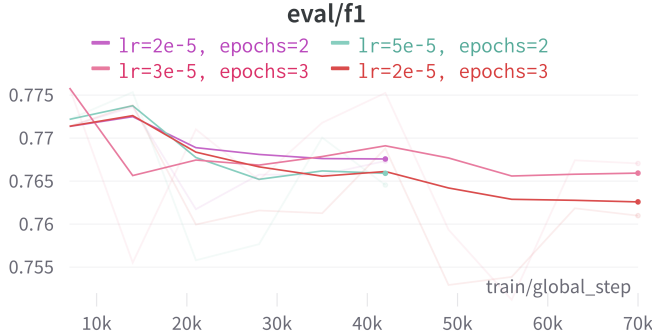


Figure 1: BioClinicalBERT: Evaluation F1

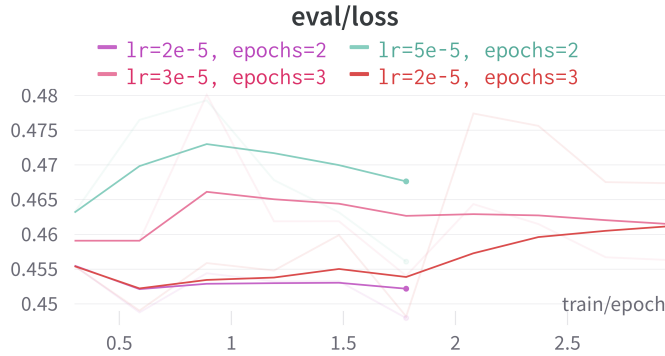


Figure 2: BioClinicalBERT: Evaluation Loss

We then use the BioClinicalBERT model to predict the probability of each sentence in the article being *summarizing*. The top 10 sentences with the highest probability are selected and concatenated to produce the final extractive summary. We arrive at this number after analysing the token distribution and finding that 10 sentences is a reasonable number to fit within the maximum input size of the GPT-2 abstractive model (i.e., 1,024 tokens split between the 10 sentences and their lay paraphrases). While we are aware that this can cause the *dangling anaphora phenomenon* (Lin 2009), we use the extracted text only as an intermediate step fed into the abstractive models which paraphrase it into lay language.

## 2.3 Abstractive Network

Once the extractive summary is generated, we train the abstractive models on the lay summaries and the extractive summaries. For this, we compare two models: GPT-2 (Radford et al. 2019a) and Clinical-Longformer (Li et al. 2022). We fine tune both models separately on eLife and PLOS. This is done due to the difference in structure and the average number of tokens in the lay summaries between the two datasets (i.e., 450 and 800 for PLOS and eLife, respectively). Hyperparameters are set based on widely used values in the literature (Li et al. 2022; Radford et al. 2019a; Devlin et al. 2019).

### 2.3.1 Clinical Longformer Abstractor

The Clinical Longformer (Li et al. 2023) is a transformer-based model that is pre-trained on the MIMIC-III dataset (Johnson et al. 2016) and can process up to 4,096 tokens in a single input sequence. This is achieved by the implementation of a sparse attention mechanism that allows more computationally efficient processing of long-range dependencies. We fine-tune the Clinical Longformer as a sequence-to-sequence task on pairs of (a) gold lay summaries and (b) ROUGE-maximising training data described in Section 2.2.

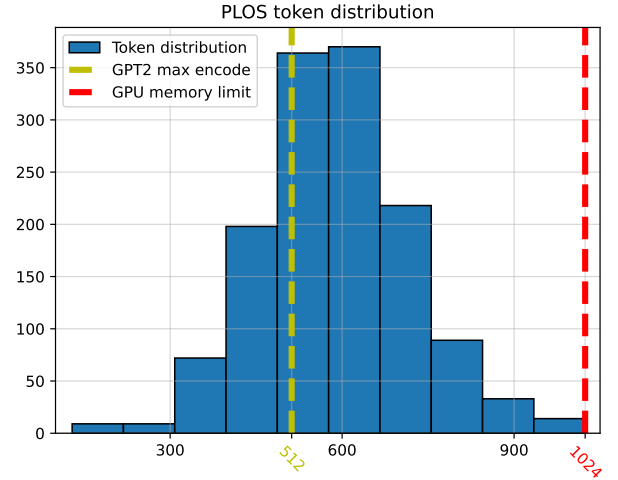


Figure 3: Token Distribution of Extracted Summaries

For the Longformer model, we experimented with window, batch, and input size to ensure that we would not run out of memory during training, as this is a common issue with Longformer models (Orzhenskii 2021). We found that a window size of 64, batch size of 1, and input size of 1,024 worked best for our dataset.

### 2.3.2 GPT-2 Abstractor

The GPT-2 is an autoregressive language model that was trained using a casual language modeling objective (Radford et al. 2019b). Given its extensive exposure to diverse text sources and natural language patterns, we hypothesize that GPT-2 would be particularly adept at generating lay summaries, making it a promising candidate for the abstractive summarization task. To fine-tune GPT-2 for this purpose, we utilize a “TL;DR” prompt, instructing the model to

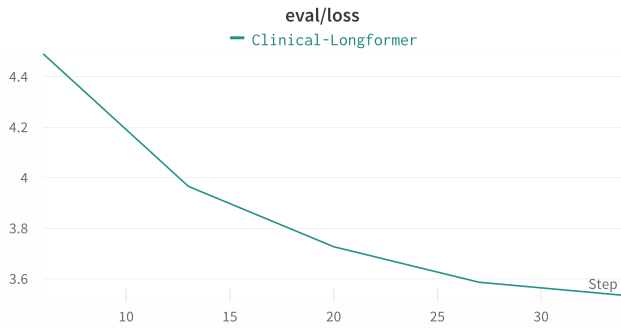


Figure 4: Longformer evaluation loss

generate concise and informative summaries.

Similar to the Longformer, we train GPT-2 on both eLife and PLOS datasets, adopting most hyperparameters from the existing literature to ensure optimal performance. Since GPT-2 can accommodate a total of 1024 tokens, we experimented with various splits between the number of tokens allocated for the extracted summary and the lay summary. Through experimentation, we determined that allocating 507 tokens for the article and 512 tokens for the summary, with 5 reserved for the “TL;DR” prompt, yielded the best results in terms of summary quality and model performance. The evaluation loss decrease during the fine-tuning process is illustrated in Figure 5.

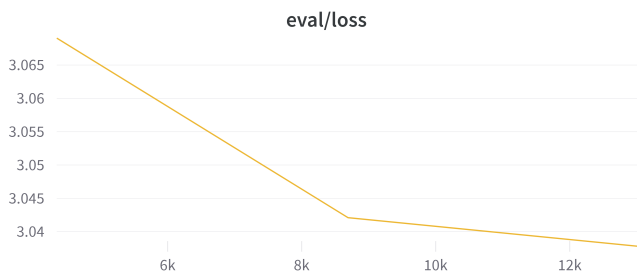


Figure 5: GPT 2 Evaluation Loss

In the evaluation phase, we compared the performance of the GPT-2 Abstractor against the Clinical Longformer Abstractor, as well as other summarization models. The results indicate that both models have their strengths and weaknesses, which we will discuss in further detail in the following sections.

### 3 Evaluation

### 4 Discussion and Conclusion

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Model	Rouge1	Rouge2	RougeL	FKGL	ARI	Gunning
Lexrank	0.334	0.085	0.164	33.59	15.41	18.50
Extractive	0.329	0.0998	0.163	10.6	25.01	26.22
GPT2	0	0	0	0	0	0
Longformer	0.289	0.062	0.143	27.33	16.89	18.44

Table 3: ROUGE and readability statistics.

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