

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

Ahmed Soliman Marc Wenzlawski Vladislav Yotkov

Abstract

In this project, we present an extractive-abstractive lay summarization pipeline for biomedical papers to generate accessible summaries for non-experts. To achieve this, we construct a sentence-level dataset optimized for maximizing ROUGE scores, utilizing lay summaries and complete articles. We employ a BERT-based classifier to identify each article’s most critical sentences. 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 and 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 practical 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

Comprehending biomedical scientific publications can be difficult for non-experts, potentially leading to misinformed health decisions (Islam et al. 2020). Lay summaries, simplified explanations of complex scientific content, could be a solution, but they are not always available. Despite past challenges in applying Automatic Text Summarisation (ATS) to biomedicine due to insufficient data (Chandrasekaran et al. 2020), two new datasets, PLOS and eLife, offer an opportunity to bridge this gap (Goldsack et al. 2022). This study investigates ATS techniques for generating biomedical lay summaries using these datasets.

2 Methods and Datasets

In this section outline the various ATS methodologies employed in this study and describe the PLOS and eLife datasets used for training and evaluation purposes.

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 2×10^{-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.

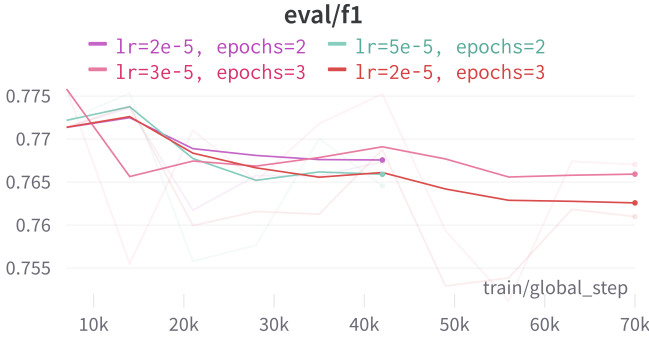


Figure 1: BioClinicalBERT: Evaluation F1

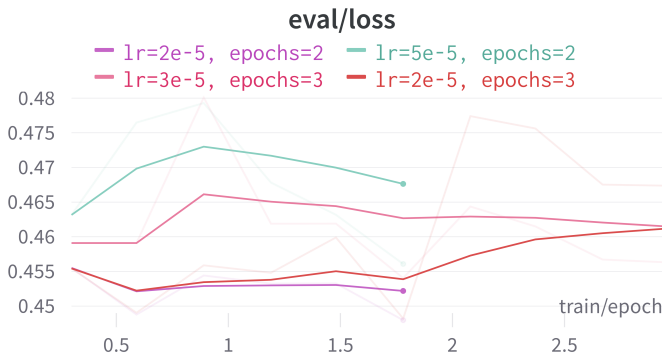


Figure 2: BioClinicalBERT: Evaluation Loss

We then use the BioClinicalBERT model to predict the probability of each sentence in the article being *summarising*. The top ten 10 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 ten 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, paraphrasing 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 implementing 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.

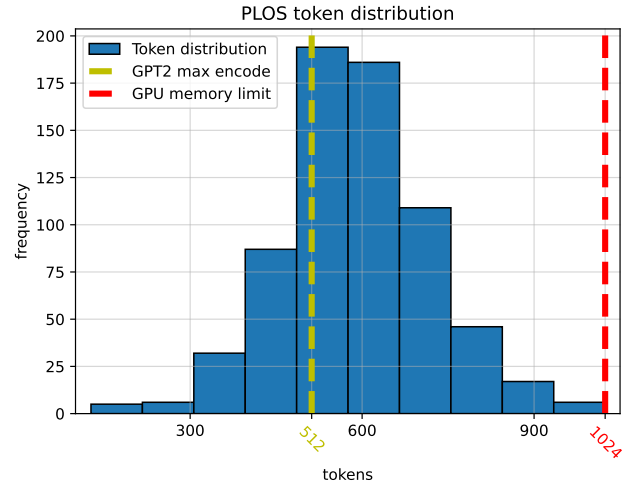


Figure 3: Token Distribution of Extracted Summaries

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 such models (Orzhenskii 2021). We found that a window size of 32, batch size of 1, and input size of 1,024 worked best for our dataset, resulting in an evaluation loss of 3.4 (Figure 4).

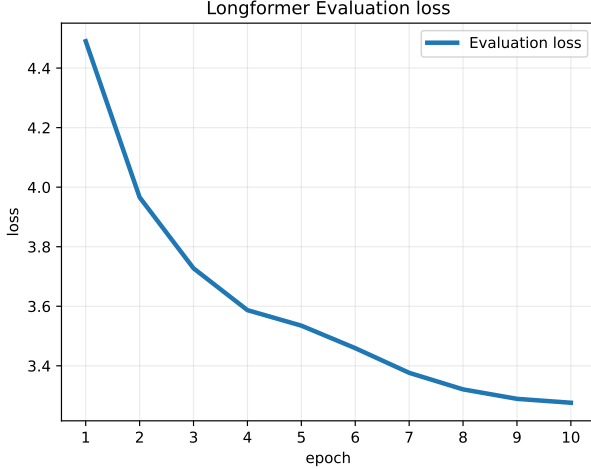


Figure 4: Longformer evaluation loss

2.3.2 GPT-2 Abstractor

The GPT-2 is an autoregressive language model that was trained using a casual language modelling 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 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 (Bajaj et al. 2021). Since GPT-2 can accommodate 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 regarding summary quality and model performance. Figure 5 illustrates the evaluation loss decrease during the fine-tuning process.

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

In this section, we evaluate the performance of the summarization models described in Section 2.

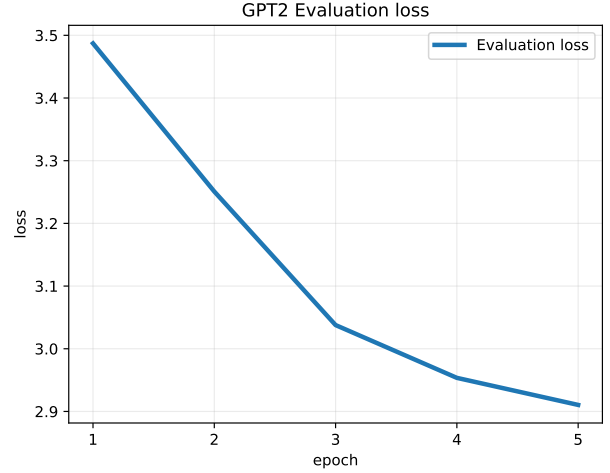


Figure 5: GPT 2 Evaluation Loss

Model	Rouge1	Rouge2	RougeL
Lexrank	0.34	0.09	0.16
Extractive	0.33	0.10	0.16
GPT2	0.18	0.02	0.09
Longformer (top-15)	0.28	0.07	0.15
Longformer (top-10)	0.29	0.06	0.14

Table 3: ROUGE F1 Scores.

Model	FKGL	ARI	Gunning
Lay	20.01	16.50	19.11
Lex (Baseline)	33.58	15.41	18.50
Extractive	10.60	25.01	26.22
GPT2	30.68	21.36	23.26
Longformer (top-15)	23.84	19.62	20.62
Longformer (top-10)	27.33	16.89	18.44

Table 4: Readability metrics.

FKGL - higher is better, ARI and Gunning - lower is better

3.1 Quantitative Evaluation

3.2 Qualitative Evaluation

•

4 Discussion and Conclusion

In this section, we discuss the performance of the proposed ATS approaches, their implications, and potential future research directions in the biomedical domain.

4.1 Limitations

We identify the following limitations of our work:

Readability Evaluation: Although, we are evaluating our models with the traditional metrics: FKGL (Kincaid et al. 1975), ARI (Senter and Smith 1967), and Gunning (Gunning 1952), they are insufficient for the estimation of text readability in scientific writing. Instead, what some researchers propose is to leverage masked language models (Martinc, Pollak, and Robnik-Šikonja 2021) like the noun-phrase BERT-based metric (Luo, Xie, and Ananiadou 2022) that computes the probability of technical jargon. We appreciate that this method would have provided a more thorough evaluation of our models, and we leave it as future work.

Limited input size: Due to the limited available computational resources (i.e., Tesla V100-SXM2-16GB) we had to restrict the input size of the Longformer to 1,024 tokens (i.e., 4 times less than the maximum size). Therefore, we could not make use of the full model capabilities in attending to long-range dependencies. This limitation propagates back to our extractor network, which produces only enough sentences to fit in the abstractor network. Thus, if we could increase the Longformer’s input size, we could do the same for the Extractor model.

4.2 Future Work

In light of the limitations discussed, we propose multiple venues for future work:

T5 Experimentation (Lehman and Johnson 2023): We aim to develop and assess the Clinical T5 model as a specialized counterpart to the Clinical Longformer. The T5, a transformer-based model, boasts unique features like a denoising autoencoder in its pretraining objective, which is adept at reconstructing corrupted input text. This makes it suitable for our extractive approach, utilising sentences from disparate article sections.

Clinical Longformer Enhancement (Li et al. 2022): Our goal is to augment the Clinical Longformer’s maximum token capacity by employing advanced hardware resources. This would facilitate experimentation with larger input dimensions and model training, potentially leading to superior summarisation performance and more precise lay summaries.

Feedback Integration: We suggest incorporating readability and factual correctness rewards into our summarisa-

tion pipeline using reinforcement learning methods (Scialom et al. 2019). This can be achieved by the combination of the RNPTC metric (Luo, Xie, and Ananiadou 2022) and the factual accuracy (Zhang et al. 2020) into a single reward function, optimised via the Reinforce algorithm (Williams 1992). This approach aspires to promote the generation of summaries that are not only more comprehensible for non-experts but also more correct with respect to the input article.

4.3 Conclusion

This project presents an effective two-step approach for generating lay summaries of biomedical research articles, incorporating an extraction step using the BioClinicalBERT model and an abstractive step with either the GPT-2 or Clinical Longformer models. By comparing the performance of these models on the PLOS dataset, we demonstrate the strengths and weaknesses of each model, finding that **TODO...**

Our results indicate that the GPT-2 Abstractor generally achieves higher ROUGE-2 F1 scores, while the Clinical Longformer Abstractor excels in ROUGE-1 and ROUGE-L F1 scores. This suggests that both models have unique advantages and can be further explored to develop more efficient and accurate summarization systems. Ultimately, our study contributes to the growing body of research aimed at bridging the gap between specialized biomedical knowledge and the general public, fostering better communication and understanding in the healthcare domain.

Bibliography

- Alsentzer, Emily, John Murphy, William Boag, Wei-Hung Weng, Di Jin, Tristan Naumann, and Matthew McDermott. 2019. “Publicly Available Clinical BERT Embeddings.” In *Proceedings of the 2nd Clinical Natural Language Processing Workshop*, 72–78. Minneapolis, Minnesota, USA: Association for Computational Linguistics. <https://doi.org/10.18653/v1/W19-1909>.
- Bajaj, Ahsaas, Pavitra Dangati, Kalpesh Krishna, Pradhiksha Ashok Kumar, Rheeja Uppaal, Bradford Windsor, Eliot Brenner, Dominic Dotterer, Rajarshi Das, and Andrew McCallum. 2021. “Long Document Summarization in a Low Resource Setting Using Pretrained Language Models.” In *Proceedings of the 59th Annual Meeting of the Association for Computational Linguistics and the 11th International Joint Conference on Natural Language Processing: Student Research Workshop*, 71–80. Online: Association for Computational Linguistics. <https://doi.org/10.18653/v1/2021.acl-srw.7>.
- Chandrasekaran, Muthu Kumar, Guy Feigenblat, Eduard Hovy, Abhilasha Ravichander, Michal Shmueli-Scheuer, and Anita de Waard. 2020. “Overview and Insights from the Shared Tasks at Scholarly Document Processing 2020: CL-SciSumm, LaySumm and LongSumm.” In *Proceedings of the First Workshop on Scholarly Document Processing*, 214–24. Online: Association for Computational

- Linguistics. <https://doi.org/10.18653/v1/2020.sdp-1.24>.
- Devlin, Jacob, Ming-Wei Chang, Kenton Lee, and Kristina Toutanova. 2019. “BERT: Pre-Training of Deep Bidirectional Transformers for Language Understanding.” *ArXiv abs/1810.04805*.
- Goldsack, Tomas, Zheheng Luo, Qianqian Xie, Carolina Scarton, Matthew Shardlow, Sophia Ananiadou, and Chenghua Lin. 2023. “Overview of the BioLaySumm 2023 Shared Task on Lay Summarization of Biomedical Research Articles.” In *Proceedings of the 22st Workshop on Biomedical Language Processing*. Toronto, Canada: Association for Computational Linguistics.
- Goldsack, Tomas, Zhihao Zhang, Chenghua Lin, and Carolina Scarton. 2022. “Making Science Simple: Corpora for the Lay Summarisation of Scientific Literature.” In *Proceedings of the 2022 Conference on Empirical Methods in Natural Language Processing*, 10589–604. Abu Dhabi, United Arab Emirates: Association for Computational Linguistics. <https://aclanthology.org/2022.emnlp-main.724>.
- Gunning, Robert. 1952. *The Technique of Clear Writing*. New York: McGraw-Hill.
- Islam, Md Saiful, Tonmoy Sarkar, Sazzad Hossain Khan, Abu-Hena Mostofa Kamal, Sarkar Mohammad Mursid Hasan, Alamgir Kabir, Dalia Yeasmin, et al. 2020. “COVID-19–Related Infodemic and Its Impact on Public Health: A Global Social Media Analysis.” *The American Journal of Tropical Medicine and Hygiene* 103 (4). <https://doi.org/https://doi.org/10.4269/ajtmh.20-0812>.
- Johnson, A. E. W., T. J. Pollard, L. Shen, L. W. Lehman, M. Feng, M. Ghassemi, B. Moody, P. Szolovits, L. A. Celi, and R. G. Mark. 2016. “MIMIC-III, a Freely Accessible Critical Care Database.” *Scientific Data* 3: 160035. <https://doi.org/10.1038/sdata.2016.35>.
- Kincaid, J. Peter, Robert P. Fishburne, Richard L. Rogers, and Brad S. Chissom. 1975. “Derivation of New Readability Formulas (Automated Readability Index, Fog Count and Flesch Reading Ease Formula) for Navy Enlisted Personnel.” In .
- Lehman, Eric, and Alistair Johnson. 2023. “Clinical-T5: Large Language Models Built Using MIMIC Clinical Text V1.0.0.” *Physionet.org*. <https://physionet.org/content/clinical-t5/1.0.0/>.
- Li, Yikuan, Ramsey M. Wehbe, Faraz S. Ahmad, Hanyin Wang, and Yuan Luo. 2022. “Clinical-Longformer and Clinical-BigBird: Transformers for Long Clinical Sequences.” <https://arxiv.org/abs/2201.11838>.
- Li, Yikuan, Ramsey M. Wehbe, Faraz S. Ahmad, Hanyin Wang, and Yuan Luo. 2023. “A Comparative Study of Pretrained Language Models for Long Clinical Text.” *Journal of the American Medical Informatics Association* 30 (2): 340–47.
- Lin, Jimmy. 2009. “Summarization.” In *Encyclopedia of Database Systems*, 2906–10. Heidelberg, Germany: Springer.
- Luo, Zheheng, Qianqian Xie, and Sophia Ananiadou. 2022. “Readability Controllable Biomedical Document Summarization.” In *Findings of the Association for Computational Linguistics: EMNLP 2022*, 4667–80. Abu Dhabi, United Arab Emirates: Association for Computational Linguistics. <https://aclanthology.org/2022.findings-emnlp.343>.
- Martinc, Matej, Senja Pollak, and Marko Robnik-Šikonja. 2021. “Supervised and Unsupervised Neural Approaches to Text Readability.” *Computational Linguistics* 47 (1): 141–79. https://doi.org/10.1162/coli_a_00398.
- Nallapati, Ramesh, Feifei Zhai, and Bowen Zhou. 2017. “SummaRuNNer: A Recurrent Neural Network Based Sequence Model for Extractive Summarization of Documents.” In *Proceedings of the Thirty-First AAAI Conference on Artificial Intelligence*, 3075–81. AAAI’17. San Francisco, California, USA: AAAI Press.
- Orzhenovskii, Mikhail. 2021. “T5-LONG-EXTRACT at FNS-2021 Shared Task.” In *Proceedings of the 3rd Financial Narrative Processing Workshop*, 67–69. Lancaster, United Kingdom: Association for Computational Linguistics. <https://aclanthology.org/2021.fnp-1.12>.
- Radford, Alec, Jeff Wu, Rewon Child, David Luan, Dario Amodei, and Ilya Sutskever. 2019a. “Language Models Are Unsupervised Multitask Learners.”
- Radford, Alec, Jeffrey Wu, Rewon Child, David Luan, Dario Amodei, and Ilya Sutskever. 2019b. *Language Models Are Unsupervised Multitask Learners*. https://cdn.openai.com/better-language-models/language_models_are_unsupervised_multitask_learners.pdf.
- Scialom, Thomas, Sylvain Lamprier, Benjamin Piwowarski, and Jacopo Staiano. 2019. “Answers Unite! Unsupervised Metrics for Reinforced Summarization Models.” In *Proceedings of the 2019 Conference on Empirical Methods in Natural Language Processing and the 9th International Joint Conference on Natural Language Processing (EMNLP-IJCNLP)*, 3246–56. Hong Kong, China: Association for Computational Linguistics. <https://doi.org/10.18653/v1/D19-1320>.
- Senter, R J, and E A Smith. 1967. “Automated Readability Index.” AMRL-TR-6620. Wright-Patterson Air Force Base: Aerospace Medical Research Laboratories (U.S.).
- Williams, Ronald J. 1992. “Simple Statistical Gradient-Following Algorithms for Connectionist Reinforcement Learning.” *Mach. Learn.* 8 (3–4): 229–56. <https://doi.org/10.1007/BF00992696>.
- Zhang, Yuhao, Derek Merck, Emily Tsai, Christopher D. Manning, and Curtis Langlotz. 2020. “Optimizing the Factual Correctness of a Summary: A Study of Summarizing Radiology Reports.” In *Proceedings of the 58th Annual Meeting of the Association for Computational Linguistics*, 5108–20. Online: Association for Computational Linguistics. <https://doi.org/10.18653/v1/2020.acl-main.458>.
- Zmandar, Nadhem, Abhishek Singh, Mahmoud El-Haj, and Paul Rayson. 2021. “Joint Abstractive and Ex-

tractive Method for Long Financial Document Summarization.” In *Proceedings of the 3rd Financial Narrative Processing Workshop*, 99–105. Lancaster, United Kingdom: Association for Computational Linguistics. <https://aclanthology.org/2021.fnp-1.19>.