MedExQA: Medical Question Answering Benchmark with Multiple Explanations

Anonymous ACL submission

Abstract

This paper introduces MedExQA, a novel benchmark in medical question-answering, to evaluate large language models' (LLMs) understanding of medical knowledge through explanations. By constructing datasets across five distinct medical specialties and further incorporating multiple explanations for each question-answer pair, we address a major gap in current medical QA benchmarks which is the absence of comprehensive assessments of LLMs' ability to generate nuanced medical explanations. Our work highlights the importance of explainability in medical LLMs, proposes an effective methodology for evaluating models beyond classification accuracy, and sheds light on one specific domain, speech language pathology, where current LLMs including GPT4 lack good understanding. Our results show generation evaluation with multiple explanations aligns better with human assessment, highlighting an opportunity for a more robust automated comprehension assessment for LLMs. To diversify open-source medical LLMs (currently mostly based on Llama2), this work also proposes a new medical model, MedPhi-2, based on Phi-2 (2.7B). The model outperformed medical LLMs based on Llama2-70B in generating explanations, showing its effectiveness in the resource-constrained medical domain. We will share our benchmark datasets and the trained model.

1 Introduction

Recent advancements in large language models (LLMs) have not only enhanced their understanding of medical domain text but also improved their ability to generate coherent text with correct medical knowledge (Tu et al., 2023; Singhal et al., 2023). Chatbots, powered by these advanced LLMs, have emerged as indispensable tools, offering unprecedented opportunities to enhance patient care, streamline clinical decision-making processes, and medical knowledge retrieval for the
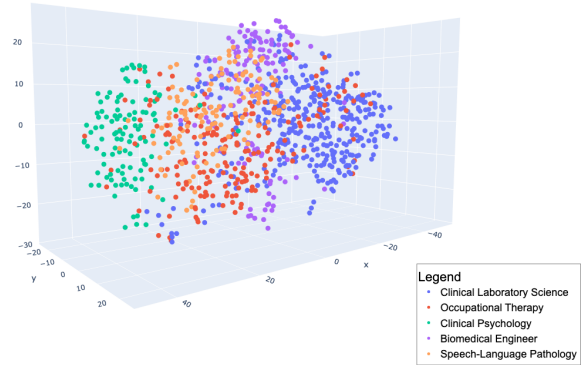

Figure 1: 3D tSNE plot for MedExQA specialties.

general public (Achiam et al., 2023; OpenAI, 2023; Groves et al., 2023). Moreover, open-source medical LLMs further enhance the usability of such technologies in hospitals by resolving the privacy concerns associated with patient data (Toma et al., 2023; Kweon et al., 2023; Chen et al., 2023).

This research in medical LLMs has been facilitated by the introduction of question-answering (QA) datasets that serve as benchmarks for evaluating the model's understanding of medical domain knowledge (Hendrycks et al., 2020; Jin et al., 2021; Pal et al., 2022; Singhal et al., 2023). The benchmark QA datasets typically consist of multiple-choice questions (MCQ), enabling researchers to readily assess the capabilities of LLMs in comprehending and responding to diverse medical inquiries. Thus, the diversity of medical scenarios and inquiries within these datasets is a key component in creating a rigorous assessment benchmark for complex medical concepts. Nonetheless, certain areas within the medical domain, such as speech language pathology, still remain uncovered by the current benchmark datasets.

As current medical QA benchmarks are often structured as MCQ, classification accuracy is used as an evaluation metric. However, classification accuracy alone may not adequately assess whether

LLMs possess the nuanced medical expertise required for reasoned responses. The explanation and rationale behind the selection of a particular choice by an LLM would provide a deeper understanding of the model's capabilities and limitations in generating responses to intricate medical questions. This comprehensive evaluation, delving into the explanation and rationale, is especially important in clinical settings where misleading information can have serious consequences. LLMs have the potential to produce convincing false medical information, known as hallucinations, leading to erroneous decisions that are challenging to rectify. Thus, understanding the rationale behind the model's responses is essential for mitigating potential harms and ensuring the safe deployment of LLMs in clinical practice.

In order to assess the quality of the model explainability, the dataset should include a golden explanation for the reasoning behind the answer. Additionally, since there are often multiple ways to express the same rationale in text, an ideal dataset would provide a multiple set of explanations for a single QA pair. However, current benchmark datasets are not focused on providing explanations as they often lack explanations entirely or only a subset of the dataset comes with an explanation (Hendrycks et al., 2020; Jin et al., 2021; Pal et al., 2022). This limitation highlights the need for improved datasets that are explicitly designed to include comprehensive explanations.

To address this issue, this paper presents a novel QA benchmark, MedExQA, with two sets of explanations, aiming to provide a more comprehensive evaluation of LLMs in the medical domain. The benchmark consists of five datasets that cover novel specialties within medical domains: biomedical engineering, clinical l laboratory science, clinical psychology, occupational therapy, and speech language pathology. In this work, the datasets were used to benchmark the performance of an extensive list of LLMs, including those trained with medical domain text. With this comprehensive benchmark evaluation, we explored the effects of medical domain-specific training. Additionally, to diversify the pool of open-source medical LLMs which are currently almost all based on the Llama2 model, we introduce our own trained model, MedPhi-2, a Phi-2 model trained with medical domain text. Our MedPhi-2 model outperformed medical LLMs based on Llama2-70B model in generating explanations for the rationale behind the answer.

The contributions of this paper are as follows:

1. **MedExQA novel datasets with explanations.** We constructed a benchmark with 5 distinct specialties within the medical domain. The datasets include two explanations for each question and answer pairs.

2. **Comprehensive Benchmark.** We evaluated an extensive list of models: 18 baseline open-source models with various sizes (from 2.7B to 70B), 3 OpenAI GPT models, as well as our model (detailed below). In terms of evaluation approach, classification accuracy, generated explanation performance, and human evaluations are considered. To highlight, this is the first benchmark using multiple explanations, and the results demonstrate that our benchmark dataset can better evaluate language models' understanding of medical domain knowledge.

3. **MedPhi-2 model.** We trained a small language model (SLM) based on Phi-2 model, with medical pretraining corpus and instruction-tuning datasets. The model outperformed medical LLMs based on Llama2 70B in generating explanations.

4. **Open source.** We will release the datasets, model weights, and codes to facilitate the research in medical large language modeling.

2 Related Works

2.1 MMLU

MMLU (Massive Multitask Language Understanding) (Hendrycks et al., 2020) is a benchmark designed to measure the model's ability in knowledge-intensive QA across 57 subjects. These subjects cover various levels of education: high school, college, and professional level. Questions in the dataset are structured as four-way MCQs, offering a standardized format for evaluation. Within the extensive list of subjects, there are nine healthcare-related subjects which are college medicine, professional medicine, clinical knowledge, anatomy, high school biology, college biology, medical genetics, nutrition, and virology. Collectively, these nine subjects comprise a total of 1,871 questions in the test set. However, it's worth noting that while MMLU provides a comprehensive set of questions, it lacks explanations for the answers, thereby limiting the dataset's evaluation to mere multiple-choice classification accuracy.

2.2 MedQA

MedQA (Jin et al., 2021) is an open-ended MCQ dataset made from professional medical doctor license exams. The dataset is available in three versions, one of which is an English version sourced from the United States Medical License Exams (USMLE). While MMLU's professional medicine subject also includes questions from USMLE practice examinations, MedQA's English version sets itself apart by incorporating questions drawn from both real exams and mock tests for USMLE. 1,273 USMLE-style questions are provided as the test dataset to benchmark the model's ability to answer medical questions at the professional level. Each question is accompanied by four or five answer choices and corresponding relevant document collections, intended to help models in generating accurate responses. As a result, similar to MMLU, MedQA does not include explanations for assessing the ability to generate rationale behind the answer.

2.3 MedMCQA

MedMCQA (Pal et al., 2022) is a benchmark with questions sourced from postgraduate-level Indian medical school entrance exams (AIIMS and NEET PG). Covering a breadth of medical specialties, the dataset has questions about 2,400 healthcare topics and 21 subjects within the medical domain. 4,183 MCQ, each offering four answer choices, are provided for evaluation. Although MedMCQA is known to have explanations, it is noteworthy that nearly half of the dataset lacks explanations, and instances of duplicate explanations are also observed. In fact, in their paper, accuracy is only reported as the evaluation metric, and explanation is not used in the evaluation process entirely. Therefore, MedMCQA is not primarily designed for the assessment of generating explanations.

3 MedExQA Datasets

We introduce MedExQA, a novel QA benchmark designed to tackle the limitations of existing benchmarks by incorporating two sets of explanations. This approach aims to offer a more thorough evaluation of performance in five novel specialties in the medical domain: Biomedical Engineering, Clinical Laboratory Science, Clinical Psychology, Occupational Therapy, and Speech Language Pathology.

Specialty	Questions
Biomedical Engineering	148
Clinical Laboratory Science	377
Clinical Psychology	111
Occupational Therapy	194
Speech Language Pathology	135
Total	965

Table 1: Number of questions for each specialty in **MedExQA**.

3.1 Datasets Preparation

The raw data was manually collected from diverse freely accessible online sources, including mock tests and online exams tailored to each medical professional specialty. The pass mark for the collected mock tests and online exams was 60 percent.

To ensure data integrity, rigorous preprocessing was conducted, including the removal of duplicate questions and explanations. Additionally, similar questions were identified and eliminated using BERT cosine similarity analysis. Questions containing keywords specific to laws or regulations were filtered out using a manually curated list of words. Following fair use regulations[1], answer options were systematically shuffled to maintain fairness and uphold the integrity of the dataset. Furthermore, to enhance the quality and coherence of the datasets, two sets of explanations as well as the questions underwent thorough human validation. This validation process aimed to ensure that the explanations exhibited distinct writing styles and provided comprehensible reasoning for the correct answer selection.

The resulting datasets have a total of 965 questions. Table 1 provides a detailed breakdown of the number of questions for each specialty. These datasets were split into a few-shot development set and a test set. Specifically, the few-shot development set has 5 questions per specialty, while the test set consists of 940 questions in total. It is noteworthy that each subject contains a minimum of 100 test examples, a length surpassing that of most exams tailored for human assessment.

An example of the dataset can be found in the appendix Figure 14. Word clouds can be found in the appendix Figure 8 for the diversity of the most frequent words across specialties.
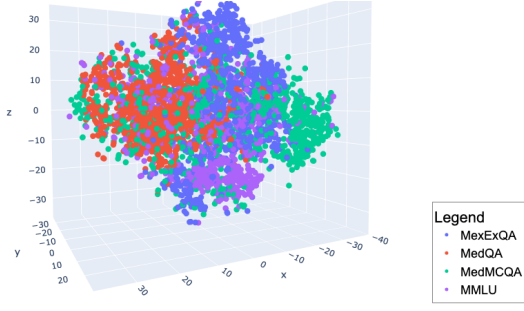
---

[1] https://www.copyright.gov/fair-use/more-info.html

Figure 2: 3D tSNE plot for MedExQA, MedQa, MedMCQA, and MMLU (Medicine Related 9 subjects) datasets.

3.2 Comparison of benchmark datasets in vector space with tSNE

In this analysis, MedQA specialty topics were visualized by their questions in the same vector space using t-distributed Stochastic Neighbour Embedding (tSNE). Each question is represented as a point in the vector space (see Table 1) and a vector representation was then obtained for each question by using the sklearn package tSNE. We used the 'all-mpnet-base-v2' sentence transformer model to retrieve vectors from questions and used 2 tSNE components. Figure 1 illustrates distinct clusters for each specialty, clearly highlighting the diversity in our dataset. Clinical Psychology demonstrates the strongest clustering away from the center. Occupational Theraphy, however, shows the greatest overlap of questions spanning across the center.

We also compare MedExQA with existing benchmark datasets by visualizing their questions in the same vector space. 965 questions were randomly sampled from each dataset. First, there is a cluster towards the top region mainly composed of questions from MedExQA, which clearly demonstrates its novelty compared to existing medical QA datasets. In addition, MedExQA questions span over a wide space from top to bottom in the center which reflects the diversity of our datasets. It is notable that MedMCQA also shows a wide topic coverage (the middle, disconnected horizontal space). Clearly, MedExQA and MedMCQA complement each other in terms of the distribution (i.e., semantics/topics of their questions). MedQA questions, on the other hand, overlap greatly with those of MedMCQA on the left, indicating their similarities. MMLU occupies a unique and rather condensed space at the bottom, indicating many of its questions have distinct and focused QA topics.
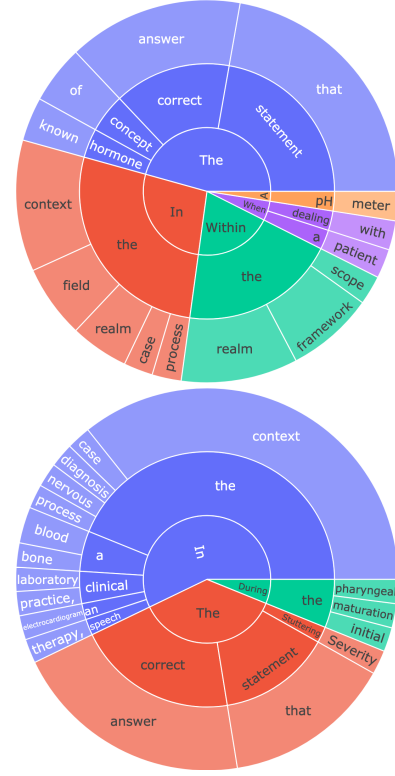

Figure 3: First three words combination of explanations. Top: Explanation 1. Bottom: Explanation 2.

3.3 First Three Words Frequency

In Figure 3, we present a detailed visualization of the lexical distribution within two distinct explanations from MedExQA datasets. For each pie chart, we combined the explanations from all five specialties. The pie chart encapsulates the hierarchical structure of the explanations, segmented into three concentric circles that correspond to the first, second, and third words of explanation, respectively. The top pie chart represents the word combination from explanation version 1, and the bottom pie chart represents the explanation version 2.

Upon examination, we note a convergence in linguistic choices, evidenced by recurring phrases such as "In the context" and "The correct answer." These phrases serve as linguistic anchors, providing a structured starting point for explanations. Despite this lexical overlap, the majority of the word choices exhibit significant variability. Some examples of this variability are "A pH meter" marked as orange and "When a patient" marked as purple on the top pie chart. By employing two versions of explanations that are semantically aligned yet lexically distinct, we aim to conduct a more holistic assessment of the model's generative outputs.

4

4 Methods

For all the experiments in this paper, both training and evaluation, we used 8 A6000 GPUs.

4.1 Baseline Models

We explored 18 baseline models with different sizes from 2.7B to 70B. Table 2 provides a comprehensive overview of the baseline models used in this paper, while more detailed descriptions of each model are available in the appendix. In cases where multiple sizes of a model are used, we distinguish each version by appending the model size to the model name. For example, the Llama2 models with sizes 7, 13, and 70B are denoted as **Llama2-7B**, **Llama2-13B**, and **Llama2-70B**, respectively. On the other hand, when a model has only one size, we refer to it solely by its name. For instance, **ClinicalCamel** denotes the ClinicalCamel 70B model.

Llama2 Variant Models	Model Size
Llama2 (Touvron et al., 2023)	7B, 13B, 70B
ClinicalCamel (Toma et al., 2023)	70B
Asclepius (Kweon et al., 2023)	7B, 13B
Med42 (M42)	70B
AlpaCare (Zhang et al., 2023)	7B, 13B
Meditron (Chen et al., 2023)	7B, 70B
Medinote (Bonnet and Boulenger)	7B, 13B
Other foundational models	**Model Size**
Mistral (Jiang et al., 2023)	7B
Yi (01.AI)	6B
Phi-2 (Microsoft)	2.7B
SOLAR (Kim et al., 2023)	10.7B
InternLm2 (Shanghai AI Lab)	7B

Table 2: Baseline Models. The models are sorted in the order of release dates.

4.2 Training MedPhi-2

As far as we know, all the publicly available open-source medical LLMs are based on Llama foundational models, we further extended our work to test the effect of medical domain training on a different foundational model. Phi-2 model was further trained using the medical datasets that are publicly available. We pretrained Phi-2 with a 110M medical-related corpus. We further finetuned the continued pretrained model with 239K instructions. We refer to the resulting model as **MedPhi-2** throughout our paper. Table 3 summarizes the detailed composition of our training dataset for continued pretraining and supervised fine-tuning. We used LLaMaFactory[2] and used Deep3 for efficient training. For both pretraining and finetuning, We trained the model with a batch size of 16 and a learning rate of 1e-5 with 3 epochs, which took 36 hours in total.

Pretrain	Tokens
Meditron Medical Guidelines[3]	48.3M
SNOMED CT descriptions	28.3M
Biomedical Article Abstracts	13.6M
Wikipedia Medical Terms	13.3M
PMC Patients Notes[4]	6.7M
Finetuning	**Instructions**
Asclepius Instruction[5]	158,114
AlpaCare Instruction[6]	52,002
NHS QA and Medical Task[7]	29,354

Table 3: MedPhi-2 training data. The number of tokens for pretraining data and the number of instructions for finetuning data are listed.

4.3 Evaluation

We evaluated all models with test datasets except for human evaluation, which was performed on the development datasets. For all the evaluations, we used zero-shot, a batch size of 1, temperature of 0. To benchmark the performance of closed source models we further extended to include OpenAI's GPT models. We used `GPT3.5_1106`, `GPT4.0_1106`, and `GPT4.0_0125` APIs[8].

4.3.1 Classification Accuracy - Logits

Classification accuracy of MCQ for generative models relies on classifying the next token using logits. In other words, the token with the highest logit value is selected as the model's predicted answer. This predicted answer is then compared to the expected answer to determine the classification accuracy. However, this approach cannot assess the model's understanding of the rationale behind the answer. We exclude `GPT3.5_1106`, `GPT4.0_1106`, and `GPT4.0_0125` for this evaluation, as we are not able to get the logit value for the next token.

---

[2] https://github.com/hiyouga/LLaMA-Factory
[3] https://huggingface.co/datasets/epfl-llm/guidelines
[4] https://huggingface.co/datasets/zhengyun21/pmc-patients
[5] https://huggingface.co/datasets/starmpcc/asclepius-synthetic-clinical-notes
[6] https://huggingface.co/datasets/casey-martin/medinstruct
[7] https://github.com/CogStack/OpenGPT
[8] https://platform.openai.com/docs/models

4.4 Classification Accuracy - Chat

We utilize string-matching to assess the model's proficiency in generating accurate textual responses. This approach involves searching for the specific phrase corresponding to the answer choice or the choice letter within the generated response, enabling a more realistic evaluation of the model's performance. We implement the string matching using regular expressions and `thefuzz` package.

4.4.1 Explanation Generation

The quality of generated explanations is further assessed using a combination of general lexical metrics. **BLEU (Papineni et al., 2002).** measures the geometric mean of precision scores of the generated explanations compared to reference explanations based on n-gram matches. **ROUGE (Lin, 2004).** assesses the similarity between generated and reference explanations, with ROUGE-L, providing a score that combines precision and recall based on the longest common subsequence. **METEOR (Banerjee and Lavie, 2005).** considers the semantic similarity and lexical variations with WordNet. **BERTScore (Zhang et al., 2019).** uses contextual embeddings, scibert embedding (Beltagy et al., 2019) for our work, to capture nuances in the semantics of the explanations. All the metrics are calculated using `evaluate` package.

We propose an enhanced methodology for evaluating models' understanding of medical domain knowledge by incorporating classification accuracy based on string matches into calculating these metrics. We assign a score of 0 to responses with incorrect answers based on string-matching classification results.

4.5 Evaluation - Human Evaluation

For human evaluation, three human annotators participated in assessing the quality of generated explanations. The evaluation process involved assigning a score for each explanation-answer pair based on the following rules:

1. **Score 0** the answer was incorrect, no explanation was provided, and/or the explanation is fully irrelevant.
2. **Score 0.5** the answer was correct, but the explanation or rationale was incorrect. Also, an incomplete explanation that ended with an incomplete sentence.
3. **Score 1.0** when both the answer and explanation were correct.

Although this human evaluation was performed on a small scale (development dataset: 5 samples for each specialty), this systematic evaluation process ensured a comprehensive assessment of the models' performance in providing accurate and coherent explanations.

5 Results and Discussion

5.1 Classification Accuracy - Logits
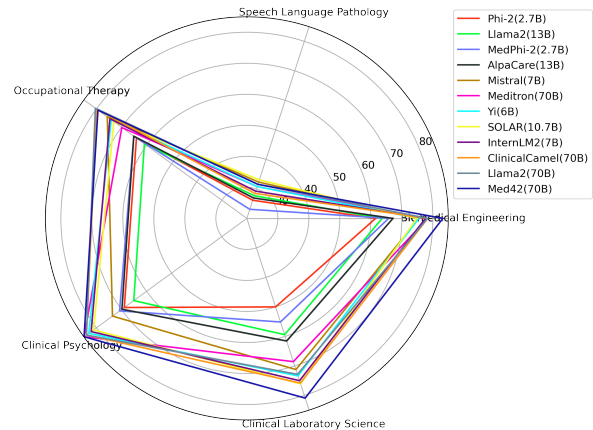
Figure 4: Multiple-choice accuracy using logits, which scales from 0 to 100. The models in the legend are ordered by macro average from lowest to highest. Only models passed (60 or above) in at least one of the specialties are included.

Figure 4 shows the conventional multiple-choice accuracy using logits across different specialties within MedExQA for the models that passed, with an accuracy above 60%, in at least one of the specialties. The models that did not pass this mark are primarily medical LLMs based on Llama-7B and 13B models. Smaller language models demonstrated relatively lower accuracy across specialties compared to larger models, as a result of their reduced capacity to capture complexity and nuances in language.

Med42 showed the best overall performance. It showed outstanding performance in Biomedical Engineering and Clinical Laboratory Science (83.22% and 84.91% respectively). It performed on par with Meditron-70B in Clinical Psychology (84.91%). In Occupational Therapy, Llama2-70B showed the highest accuracy (80.42%). All models underperformed in Speech Language Pathology, with SOLAR performing the best (33.08%). See Appendix Table 4 for detailed results of all models.

The effect of continued training is observed only in some models. MedPhi-2 demonstrated better

performance than Phi-2, and this improvement was also found in AlpaCare-13B compared to Llama2-13B and Med42 compared to Llama2-70B. However, ClinicalCamel and Meditron-70B performed worse than Llama2-70B. This drop in performance could be due to task-specific challenges as some models may not effectively handle varied levels of specificity in MedExQA.
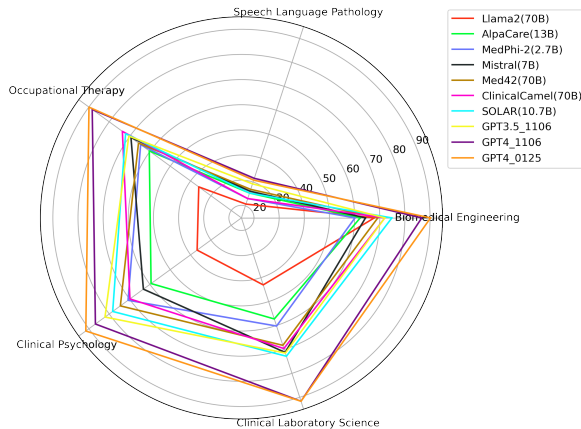

Figure 5: Multiple-choice accuracy using chat. The Figure style is the same as Figure 4.

5.2 Classification Accuracy - Chat

Accuracy based on the string match of the generated response shows slight changes in the performance of models (Figure 5). Most models performed worse in this classification accuracy than the conventional classification accuracy using logits. Phi-2, Llama2-13B, Yi, InternLM2, and Meditron-70B did not pass the pass mark indicating these models are not robust. Meditron-70B showed the biggest performance drop by 29.28%. Llama2-70B also showed a significant performance drop in this testing by 28.54%, although it passed in Biomedical Engineering. Of the 70B models we tested, ClinicalCamel was the most robust model (7.67% decrease), and it scored higher than Med42 by 0.71%.

Our model, MedPhi-2 was the most robust model among the passed ones (0.2% decrease), and it outperformed AlpaCare-13B, Meditron-70B, Llama2-70B. This result highlights the importance of the supervised finetuning with in-domain instructions of high quality as more robust models, such as AlpaCare, ClinicalCamel, and MedPhi-2, were instruction-tuned with medical domain data, while Meditron-70B was just further pretrained.

Closed-source models such as OpenAI's GPT4_0125, GPT4_1106, and GPT3.5_1106 out-performed all the open-source models. Even with the addition of high-performing closed-source models, there is still a universal failure in performance for Speech Language Pathology. See Appendix Table 5 for detailed results of all models.

5.3 Combining Classification Accuracy with Generated Explanation Performance

Figure 6 shows the relationship between model size and accuracy achieved in both MCQ (using logits) and generation performance. Generally, larger models tend to exhibit better performance as 70B models perform better than most of the other smaller models. However, interestingly, all medical LLMs with 13B (AlpaCare, Asclepius, and Meditron) exhibit worse performance in both MCQ accuracy and generation performance compared to their 7B counterparts. In fact, **Medinote-13B** is the worst-performing model. Also, 70B models do not always perform better than smaller models as Meditron-70B and Llama2-70B performed worse than many smaller models including AlpaCare and our model in the generation of reasonable explanations.

This result confirms the findings of Section 5.2 which highlighted the importance of supervised finetuning with in-domain instructions. In fact our model, **MedPhi-2** has performed exceptionally well in generating reasonable explanations and out-performed all medical LLMs including 70B models.

5.4 Evaluation - Human Evaluation

Human evaluation of generated responses reveals that MedPhi-2 has the best quality among the open-source models (Figure 7). Our model was the only open-source model that passed (a score of 3 or above) in all specialties in MedExQA. In fact, **MedPhi-2** on par with **GPT3.5_1106** in Biomedical Engineering and Clinical Laboratory Science, and with **GPT4_1106** in Occupational Therapy.

Figure 14 provides an example of generated responses of the models, in the context of Speech-Language Pathology questions. **MedPhi-2** and **GPT3.5_1106** generated the most coherent and accurate responses. However, other models generated irrelevant sentences or failed to provide explanations. **Medinote-13B** generated a case study example instead of answering the question and providing an explanation and **Asclepius-13B** hallucinated and provided an option for the answer that was not present and generated further incorrect ex-
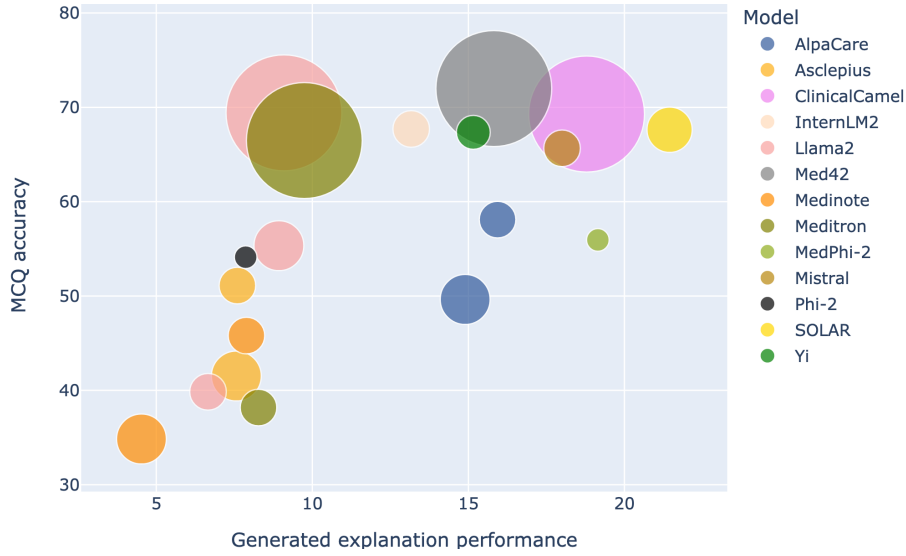
Figure 6: Scatter plot of model performance. The Y-axis is the macro average of accuracy based on logits (Table 4). The X-axis is the average score of generated explanations (Table 6). The dot size is proportional to the model size.
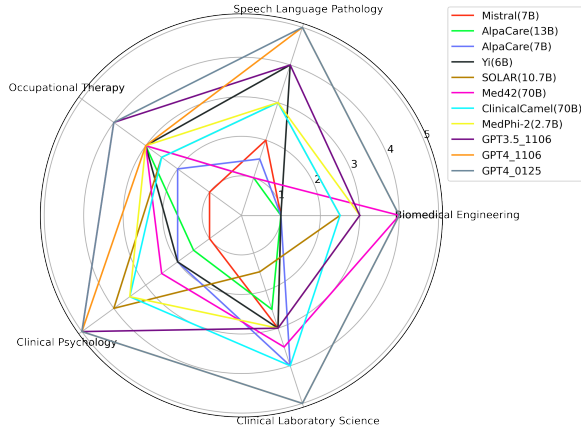

Figure 7: Human evaluation on the generated explanations, which scales from 0 to 5. The Figure style is the same as Figure 4.

planations. Appendix Table 7 shows the detailed results for all the models.

The effect of adding additional explanation was confirmed by analyzing the Pearson correlation between human evaluation and generation performance. When we used just one set of explanations the correlation was 0.9347, and this correlation increased to 0.9385 when we used two versions of explanations together. This indicates generation evaluation with multiple explanations aligns better with human evaluation, which is usually treated as the gold standard.

6 Conclusion

Our MedExQA benchmark demonstrates the significant potential of LLMs in enhancing medical QA with explanations. The findings reveal that the generation of coherent and accurate explanations remains a challenging frontier for the current medical LLMs. We also find that the 'Speech Language Pathology' dataset posed challenges for all language models, including GPT4. Through the development and evaluation of our MedPhi-2 model, we underscore the importance of targeted pretraining and fine-tuning strategies in improving explanation quality. Our benchmark and model will set the foundation for future advancements in medical research by facilitating the development and evaluation of medical LLMs.

Limitation

While MedExQA provides a robust benchmark for evaluating LLMs in the context of the medical domain, the current version only tests the model's ability in QA task, limiting its applicability in real-world clinical scenarios to a few applications. This limitation results from the manual collection process. Future work will extend our benchmark to include tasks such as summarizing clinical notes with accompanying explanations.

Though we performed the human evaluation of generated explanations of different LLMs through three authors, we performed this at a small scale, at 5 samples per specialty. Future work will seek to increase both the volume of samples and the number of annotators to provide a more robust method of assessing models' performance.

Broader Impacts and Ethics Statement

We will release MedExQa under a Creative Commons Attribution-Non Commercial-ShareAlike 4.0 International License. MedPhi-2 will follow the MIT license as it is based on Phi-2. License and copyright information and Terms of Use will be shared when the dataset and model are released. The dataset may be used for non-commercial purposes and any models trained using the dataset should be used only for research purposes.

Our work does not raise any major ethical concerns. All LLMs tested, including Phi-2, were used for research purposes only. While MedPhi-2 outperformed all medical variants of Llama2 models in generating accurate medical answers and explanations, MedPhi-2 is not rigorously tested for use in real-world clinical applications or scenarios. Thus, MedPhi-2 is not suitable for use in the clinical decision making process. This restriction of usage in clinical care is to mitigate any potential risks or harms such as wrong decisions from hallucinations which can lead to unwanted situations.

References

01.AI. Yi. building the next generation of open-source and bilingual llms.

Josh Achiam, Steven Adler, Sandhini Agarwal, Lama Ahmad, Ilge Akkaya, Florencia Leoni Aleman, Diogo Almeida, Janko Altenschmidt, Sam Altman, Shyamal Anadkat, et al. 2023. Gpt-4 technical report. *arXiv preprint arXiv:2303.08774*.

Satanjeev Banerjee and Alon Lavie. 2005. Meteor: An automatic metric for mt evaluation with improved correlation with human judgments. In *Proceedings of the acl workshop on intrinsic and extrinsic evaluation measures for machine translation and/or summarization*, pages 65–72.

Iz Beltagy, Kyle Lo, and Arman Cohan. 2019. Scibert: A pretrained language model for scientific text. *arXiv preprint arXiv:1903.10676*.

Antoine Bonnet and Paul Boulenger. Medinote.

Zeming Chen, Alejandro Hernández Cano, Angelika Romanou, Antoine Bonnet, Kyle Matoba, Francesco Salvi, Matteo Pagliardini, Simin Fan, Andreas Köpf, Amirkeivan Mohtashami, et al. 2023. Meditron-70b: Scaling medical pretraining for large language models. *arXiv preprint arXiv:2311.16079*.

Emily Groves, Minhong Wang, Yusuf Abdulle, Holger Kunz, Jason Hoelscher-Obermaier, Ronin Wu, and Honghan Wu. 2023. Benchmarking and analyzing in-context learning, fine-tuning and supervised learning for biomedical knowledge curation: a focused study on chemical entities of biological interest. *arXiv preprint arXiv:2312.12989*.

Dan Hendrycks, Collin Burns, Steven Basart, Andy Zou, Mantas Mazeika, Dawn Song, and Jacob Steinhardt. 2020. Measuring massive multitask language understanding. *arXiv preprint arXiv:2009.03300*.

Albert Q Jiang, Alexandre Sablayrolles, Arthur Mensch, Chris Bamford, Devendra Singh Chaplot, Diego de las Casas, Florian Bressand, Gianna Lengyel, Guillaume Lample, Lucile Saulnier, et al. 2023. Mistral 7b. *arXiv preprint arXiv:2310.06825*.

Di Jin, Eileen Pan, Nassim Oufattole, Wei-Hung Weng, Hanyi Fang, and Peter Szolovits. 2021. What disease does this patient have? a large-scale open domain question answering dataset from medical exams. *Applied Sciences*, 11(14):6421.

Dahyun Kim, Chanjun Park, Sanghoon Kim, Wonsung Lee, Wonho Song, Yunsu Kim, Hyeonwoo Kim, Yungi Kim, Hyeonju Lee, Jihoo Kim, et al. 2023. Solar 10.7 b: Scaling large language models with simple yet effective depth up-scaling. *arXiv preprint arXiv:2312.15166*.

Sunjun Kweon, Junu Kim, Jiyoun Kim, Sujeong Im, Eunbyeol Cho, Seongsu Bae, Jungwoo Oh, Gyubok Lee, Jong Hak Moon, Seng Chan You, et al. 2023. Publicly shareable clinical large language model built on synthetic clinical notes. *arXiv preprint arXiv:2309.00237*.

Chin-Yew Lin. 2004. Rouge: A package for automatic evaluation of summaries. In *Text summarization branches out*, pages 74–81.

M42. Med42 - clinical large language model.

Microsoft. Phi-2: The surprising power of small language models.

OpenAI. 2023. Chatgpt.

Ankit Pal, Logesh Kumar Umapathi, and Malaikannan Sankarasubbu. 2022. Medmcqa: A large-scale multi-subject multi-choice dataset for medical domain question answering. In *Conference on Health, Inference, and Learning*, pages 248–260. PMLR.

Kishore Papineni, Salim Roukos, Todd Ward, and Wei-Jing Zhu. 2002. Bleu: a method for automatic evaluation of machine translation. In *Proceedings of the 40th annual meeting of the Association for Computational Linguistics*, pages 311–318.

Shanghai AI Lab. Internlm.

Karan Singhal, Tao Tu, Juraj Gottweis, Rory Sayres, Ellery Wulczyn, Le Hou, Kevin Clark, Stephen Pfohl, Heather Cole-Lewis, Darlene Neal, et al. 2023. Towards expert-level medical question answering with large language models. *arXiv preprint arXiv:2305.09617*.

Augustin Toma, Patrick R Lawler, Jimmy Ba, Rahul G Krishnan, Barry B Rubin, and Bo Wang. 2023. Clinical camel: An open-source expert-level medical language model with dialogue-based knowledge encoding. *arXiv preprint arXiv:2305.12031*.

Hugo Touvron, Louis Martin, Kevin Stone, Peter Albert, Amjad Almahairi, Yasmine Babaei, Nikolay Bashlykov, Soumya Batra, Prajjwal Bhargava, Shruti Bhosale, et al. 2023. Llama 2: Open foundation and fine-tuned chat models. *arXiv preprint arXiv:2307.09288*.

Tao Tu, Shekoofeh Azizi, Danny Driess, Mike Schaekermann, Mohamed Amin, Pi-Chuan Chang, Andrew Carroll, Chuck Lau, Ryutaro Tanno, Ira Ktena, et al. 2023. Towards generalist biomedical ai. *arXiv preprint arXiv:2307.14334*.

USMLE. United states medical licensing examination.

Tianyi Zhang, Varsha Kishore, Felix Wu, Kilian Q Weinberger, and Yoav Artzi. 2019. Bertscore: Evaluating text generation with bert. *arXiv preprint arXiv:1904.09675*.

Xinlu Zhang, Chenxin Tian, Xianjun Yang, Lichang Chen, Zekun Li, and Linda Ruth Petzold. 2023. Alpacare: Instruction-tuned large language models for medical application. *arXiv preprint arXiv:2310.14558*.

Appendix

.1 Example of MedExQA

Our dataset has the format of multiple-choice question and 2 sets of explanations. Figure 14 shows an example of Speech Language Pathology specialty dataset.

.2 WordCloud

Word clouds generated from each dataset highlight the most frequent terms or phrases. Figure 8 demonstrates the different word clouds for the 5 specialties present in the dataset. These word clouds were generated from the analysis of both the questions and answers of the datasets. It is clear to see that in MedExQA, words adjacent to the relevant specialty are the most common, with terms such as 'muscle' highly prevalent for Occupational Therapy orientated questions and 'cell' for Clinical Laboratory Science related questions. Compared to the word clouds of other datasets, seen in the Appendix. MedExQA shows a greater breadth of topic coverage.

Figure 8: WordClouds for each specialties in MedExQA

Figure 9: WordCloud for MedExQA Dataset

Figure 10: WordCloud for MMLU Dataset

Figure 11: WordCloud for MedMCQA Dataset

Figure 12: WordCloud for MedQA Dataset

Figure 13: Word Count Distribution Plots for Explanations. Top: Explanation 1. Bottom: Explanation 2.

11

Question	**Which of the following is a correct description of a language learning disability?** A It shows difficulties in reading and writing. B It does not show problems with higher-level language abilities. C It occurs due to visual or motor disabilities. D It often accompanies autism spectrum disorder. **Correct Answer: A.** It shows difficulties in reading and writing.
Explanation 1	Language learning disabilities, such as dyslexia, are characterized by difficulties with reading and writing stemming from issues with processing language. These challenges are not directly caused by visual or motor impairments. Although language learning disabilities may occur alongside other disorders like autism spectrum disorder, there is no one-to-one correlation between the two; individuals with language learning disabilities may not have autism, and vice versa. …
Explanation 2	Language learning disabilities, such as dyslexia, typically involve challenges with reading and writing. These difficulties are due to problems with language processing, not necessarily related to visual or motor disabilities. While language learning disabilities can co-occur with other conditions, including autism spectrum disorder, not all individuals with language learning disabilities have autism, and not all individuals with autism have language learning disabilities. …
Medinote-13B **Score: 0**	A 10-year-old boy with a language-learning disability was assessed...
Asclepius-13B **Score: 0**	E. It is a type of specific learning disorder (SLD). …
AlpaCare-13B **Score: 0**	The answer is A) It shows difficulty in reading & writing.
Med42 **Score: 0**	So, the correct answer is: A. It showing difficulties in readings and writings.
Meditron-70B **Score: 0**	The following is an open-ended question. You should answer it in your own words…
ClinicalCamel **Score: 0.5**	\end{blockquote} think the answer is A. … It is not a problem with higher level language abilites. Visual or motor difficulties can cause learning difficulties, but they are not the cause of language learning difficulties. Language learning difficulties often accompany autism, but this is not the correct description. Therefore, the correct answer is option A.
GPT3.5_1106 **Score: 1**	A. It shows difficulties in reading and writing. Explanation: A language learning disability typically involves difficulties in reading and writing, such as dyslexia. … While other options may also be associated with learning disabilities, the primary characteristic of a language learning disability is the difficulty in reading and writing.
Phi-2 **Score: 0**	Answer: D. It is often accompanied by autism spectrum disorders. Explanation: …
MedPhi-2 **Score: 1**	The answer is A.… Additionally, language learning disorders are not typically caused by visual or hearing impairments, but rather by difficulties with processing and using linguistic information. Finally, while language learning deficits can be associated with autism spectrum disorders, not all individuals with autism have language learning problems, ...

Figure 14: Example of data in Speech Language Pathology and Qualitative Analysis Example. Two sets of explanations, Explanation 1 and Explanation 2, are provided. The score given by humans is provided beneath the model name. The response with no explanations has a grey font color. Red shows the irrelevant or wrong sentences or phrases. Yellow demonstrates incoherent phrases or errors. Green highlights coherent and medically correct sentences.

.3 Frequency Plots

We use frequency plots to demonstrate the word count distribution for the two different types of explanations in each dataset. For MedExQA, as shown in Figure 13, the average length of the words in the first set of explanations is 82.50 and in the second set of explanations, is 83.17.

.4 Baseline Models

.4.1 Llama2 variants

Llama2 We use Llama2 HuggingFace weights released on the Hugging Face model repository[9]. 7B, 13B, and 70B models without chat optimization are used in this work to assess the effect of continued training of the following Llama2 medical models with medical domain text. These models are trained on 2 trillion pretraining tokens in the general domain and have a context length of 4,096.

ClinicalCamel We use ClinicalCamel 70B weights from the Hugging Face model repository. It is a finetuned **Llama2-70B** model with instruction-tuning datasets made from medical articles and MedQA. It uses QLoRA for finetuning. The instruction tuning datasets are not released.

Asclepius We use Asclepius Llama2 weights released on the Hugging Face model repository. We use both 7B and 13B models which are further finetuned Llama2 models using instruction tuning dataset made from synthetic clinical notes. The synthetic clinical notes are generated from PMC-patients using GPT3.5 and turned into instruction-tuning datasets using GPT3.5. The synthetic clinical notes are used due to the privacy concerns of the real clinical notes. This training dataset is released.

Med42 We use Med42 70B weights from the Hugging Face model repository. The details of the training dataset and training method are not available. The only detail available is that it was continued trained **Llama2-70B** model with medical domain text.

AlpaCare We use AlpaCare Llama2 weights from the Hugging Face model repository. Llama2 7B and 13B models were further finetuned on a medical self-instruct dataset made from the clinical seed set. The dataset is released.

Meditron We use Meditron weights released on the Hugging Face model repository. Both 7B and 70B models are used in this work. Meditron models are continued pretrained with clinical guidelines, medical articles abstracts, and full text of the articles. A subset of clinical guidelines are released.

Medinote We use Medinote weights released on the Hugging Face model repository. Both 7B and 13B models are used in this work. These models are further finetuned from the Meditron models to generate clinical notes from doctor and patient dialogues. Their training dataset is a synthetic dialog generated with ChatGPT from PMC-patients data.

.4.2 Other baseline models

We extended our baseline models to other general domain baseline models with various sizes.

Mistral We use Mistral-7B-v0.1 weight released on the Hugging Face model repository. The details of the training dataset remain unknown. However, this model is known to use Grouped Query Attention, which **Llama2-70B** also uses, and Sliding Window Attention. The model size is known to be 7.24B parameters, and this is slightly larger than **Llama2-7B**, 6.74B.

Yi We use Yi-6B weight released on the Hugging Face model repository. The model is trained on 3 trillion pretraining tokens in the general domain and has a context length of 4,096. The model size is known to be 6.06B parameters, which is smaller than other 7B models.

Phi-2 We use Phi-2 model weight released on the Hugging Face model repository. It has 2.78B parameters and is trained on the augmented textbook corpus, 1.4 trillion tokens. This is the smallest model in our paper.

SOLAR We use SOLAR-10.7B-v1.0 model weight released on the HuggingFace model repository. The model size is 10.7 billion parameters. It uses depth-wise scaling called Depth up-scaling and continued pretraining of the scaled model. However, the pretraining dataset details are unknown.

InternLM2 We use InternLM2-7b model weight from the HuggingFace model repository. The details of the training method and data are unknown.

.5 Result Tables

---

[9]https://huggingface.co/meta-llama

Model	Size (B)	BE	CP	SLP	OT	CLS	MAvg	MiAvg
Medinote	7	33.57	34.91	23.08	38.1	44.62	34.85	37.55
Meditron	7	37.76	46.23	20.77	42.86	43.28	38.18	39.57
Llama2	7	41.96	47.17	22.31	40.21	47.58	39.85	41.7
Asclepius	7	44.76	47.17	27.69	42.86	45.16	41.53	42.45
Medinote	13	46.15	52.83	28.46	49.21	52.42	45.81	47.55
AlpaCare	7	53.15	53.77	26.92	59.79	54.57	49.64	51.49
Asclepius	13	57.34	56.6	25.38	59.79	56.45	51.11	52.98
Phi-2	2.7	61.54	68.87	26.15	64.02	50	54.12	53.4
Llama2	13	63.64	65.09	27.69	60.85	59.41	55.34	56.6
MedPhi-2	2.7	65.73	70.75	23.08	65.08	55.11	55.95	56.06
AlpaCare	13	67.13	69.81	26.92	65.08	61.56	58.1	59.26
Mistral	7	75.52	73.58	32.31	75.66	71.24	65.66	67.66
Meditron	70	78.32	**84.91**	30.77	69.84	68.55	66.48	66.91
Yi	6	75.52	83.02	30.77	74.07	73.39	67.35	69.04
SOLAR	10.7	74.83	81.13	**33.08**	73.02	76.08	67.63	69.8
InternLM2	7	77.62	82.08	29.23	74.6	75	67.71	69.79
ClinicalCamel	70	78.32	83.96	28.46	79.89	75.81	69.29	71.38
Llama2	70	78.32	83.96	31.54	**80.42**	72.85	69.42	70.7
Med42	70	**83.22**	**84.91**	31.54	79.37	**80.91**	**71.99**	**74.57**

Table 4: MCQ accuracy using logits. "BE": Biomedical Engineering; "CP": Clinical Psychology; "CLP": Speech Language Pathology; "OT": Occupational Therapy; "CLS": Clinical Laboratory Science; "MAvg": Macro Average; "MiAvg": Micro Average.

Model	Size (B)	BE	CP	SLP	OT	CLS	MAvg	MiAvg
Medinote	13	27.27	22.64	23.85	21.16	32.26	25.44	27.02
Phi-2	2.7	25.87	30.19	28.46	20.63	25	26.03	25.32
Asclepius	13	36.36	23.58	21.54	25.4	33.6	28.1	29.57
Medinote	7	28.67	26.42	29.23	29.63	33.06	29.4	30.43
Asclepius	7	33.57	30.19	26.15	27.51	31.72	29.83	30.21
Llama2	7	32.87	37.74	23.85	27.51	30.11	30.41	30
Meditron	7	30.07	30.19	23.08	32.28	36.56	30.43	32.13
Llama2	13	37.06	22.31	**43.92**	32.08	41.94	35.46	37.77
Meditron	70	41.96	41.51	25.38	32.8	44.35	37.2	38.72
InternLM2	7	52.45	43.4	23.85	38.62	41.4	39.94	40.32
Llama2	70	67.83	36.79	20.77	35.98	43.01	40.88	41.6
Yi	6	55.24	54.72	31.54	53.44	56.18	50.22	51.91
AlpaCare	7	59.44	55.66	33.08	56.08	54.03	51.66	52.55
AlpaCare	13	62.24	59.43	25.38	60.32	57.26	52.93	54.47
MedPhi-2	2.7	60.14	70.75	23.08	64.55	60.22	55.75	57.13
Mistral	7	64.34	63.21	26.15	69.31	71.24	58.85	62.66
Med42	70	69.23	74.53	26.92	65.61	68.28	60.91	62.87
ClinicalCamel	70	72.03	69.81	23.08	73.54	69.62	61.62	64.36
SOLAR	10.7	74.83	78.3	25.38	71.96	72.85	64.66	67.02
GPT3.5_1106	-	72.03	82.08	29.23	70.37	71.51	65.04	66.7
GPT4_1106	-	86.71	86.79	31.54	88.36	**91.67**	77.01	81.38
GPT4_0125	-	**90.21**	**91.51**	30.77	**89.95**	**91.67**	**78.82**	**82.66**

Table 5: MCQ accuracy using chat generation. "BE": Biomedical Engineering; "CP": Clinical Psychology; "CLP": Speech Language Pathology; "OT": Occupational Therapy; "CLS": Clinical Laboratory Science; "MAvg": Macro Average; "MiAvg": Micro Average.

Model	Size (B)	ROUGE-L	METEOR	BLEU	BERTScore	AVG
Medinote	13	1.88	2.79	0.46	12.96	4.52
Llama2	7	4.92	4.03	0.16	17.52	6.66
Asclepius	13	6.12	6.12	0.32	17.70	7.56
Asclepius	7	6.07	5.61	0.22	18.48	7.60
Phi-2	2.7	5.77	7.51	1.76	16.41	7.86
Medinote	7	4.78	7.82	2.14	16.81	7.89
Meditron	7	5.15	7.96	2.56	17.43	8.27
Llama2	13	6.65	6.89	1.37	20.80	8.93
Llama2	70	6.41	6.71	1.40	21.84	9.09
Meditron	70	7.42	8.32	1.63	21.59	9.74
InternLM2	7	10.30	12.20	3.89	26.28	13.17
AlpaCare	13	11.56	11.97	2.77	33.29	14.90
Yi	6	10.97	13.25	4.79	31.62	15.16
Med42	70	11.03	12.88	3.46	35.89	15.82
AlpaCare	7	12.43	14.19	3.64	33.47	15.94
Mistral	7	12.59	17.49	5.28	36.66	18.00
ClinicalCamel	70	13.45	17.38	5.52	38.80	18.79
MedPhi-2	2.7	15.26	17.75	6.13	37.45	19.15
SOLAR	10.7	16.45	20.17	6.72	42.46	21.45
GPT3.5_1106	-	21.71	25.99	14.07	46.59	27.09
GPT4_1106	-	23.08	**35.74**	14.40	**54.50**	31.93
GPT4_0125	-	**24.83**	35.21	**16.71**	54.40	**32.79**

Table 6: Explanation Generation performance (average across the 5 subjects for each evaluation metric).

Model	Size (B)	BE	CP	SLP	OT	CLS	AVG
Llama2	13	0	0	0	0	0	0
Meditron	70	0	0	0.5	0.5	0.5	0.3
Asclepius	13	0	1.5	0	0	1	0.5
Medinote	13	0.5	0.5	0.5	0	1	0.5
Meditron	7	0.5	0	1	1	0	0.5
Llama2	7	0	1	0	1.5	1	0.7
Llama2	70	0.5	0	1	0.5	2	0.8
Asclepius	7	1	1.5	0	0	2	0.9
Medinote	7	1.5	0.5	1	0	1.5	0.9
InternLM2	7	2	2	1	0	1.5	1.3
Phi-2	2.7	2	2	0	1	2	1.4
Mistral	7	1	1	2	1	3	1.6
AlpaCare	13	1	1.5	1	3	2.5	1.8
AlpaCare	7	1	2	1.5	2	4	2.1
Yi	6	1	2	4	3	3	2.6
SOLAR	10.7	2.5	4	3	2.5	1.5	2.7
Med42	70	4	2.5	1	3	3.5	2.8
ClinicalCamel	70	2.5	3.5	3	2.5	4	3.1
MedPhi-2	2.7	3	3.5	3	3	3	3.1
GPT3.5_1106	-	3	5	4	4	3	3.8
GPT4_1106	-	4	5	5	3	5	4.4
GPT4_0125	-	**4**	**5**	**5**	**4**	**5**	**4.6**

Table 7: Explanation Generation performance (human evaluation). "BE": Biomedical Engineering; "CP": Clinical Psychology; "CLP": Speech Language Pathology; "OT": Occupational Therapy; "CLS": Clinical Laboratory Science; "AVG": Average score.