Automatic Evaluation of Healthcare LLMs Beyond Question-Answering

Abstract

Current Large Language Models (LLMs) benchmarks are often based on open-ended or close-ended QA evaluations, avoiding the requirement of human labor. Close-ended measurements evaluate the factuality of responses but lack expressiveness. Open-ended capture the model's capacity to produce discourse responses but are harder to assess for correctness. Both these approaches are being used, sometimes together sometimes separately, while the relationship between both remains unclear. This work is focused on the healthcare domain, where both factuality and discourse matter greatly. It introduces a comprehensive, multi-axis suite for healthcare LLM evaluation, exploring correlations between open and close benchmarks and metrics. Findings include blind spots and overlaps in current methodologies. As an updated sanity check, we release a new medical benchmark –ANONIMIZED– , with both open and closed variants. Finally, we propose a novel metric for open-ended evaluations –Relaxed Perplexity– to mitigate the identified limitations.

1 Introduction

The growing use of large language models (LLMs) in public domains, such as healthcare, shows promise for improving global quality of life (et al., 2023). At the same time, the reliability and evaluation of LLMs in such sensitive topics requires extreme caution due to the potential impact on people's rights and well-being.

LLM evaluation today is approached through various perspectives, which consider different types of LLM assessment: automatic evaluation (scalable and factual), user evaluation (utility and usability) (et al., 2024b), and expert evaluation (support and coherence) (Chen et al., 2023). While each of these evaluation perspectives serves distinct roles that contribute to a holistic assessment, automatic evaluation remains the most prevalent one due to its lack of dependency on human effort.

Within automatic evaluation, there are two types of tests. Those which include closed-ended responses (Bedi et al., 2024), namely multiple-choice question answering (MCQA), and those which have open-ended responses (Dada et al., 2024). Close-ended MCQA validation enables the automatic verification of response factuality, but it does not reflect the complex nature of real world situations (*e.g.*, clinical settings (Hager et al., 2024; Zhou et al., 2023)). As such, MCQA alone often fails to identify critical short-coming of model performance (Umapathi et al., 2023; Ahmad et al., 2023; Pezeshkpour and Hruschka, 2023; et al., 2024a; Zheng et al., 2023). To incorporate a broader range of tasks relevant to the medical field (Dada et al., 2024; Kanithi et al., 2024) one typically has to rely on open-ended answers. That is, reference responses that are not the only valid outputs. Since these cannot be completely assessed for factuality without human expert supervision, approximate measures based on n-grams, and model perplexity remain in place, which limits the reliability of these automatic evaluations.

Efforts have been dedicated to analyze the relation between automatic evaluations and either user or expert evaluations, showing a lack of direct correspondence (Fleming et al., 2024; Nimah et al., 2023). This is explained by the difference in the model features these assess (*e.g.*, factuality vs usability vs support capacity), pointing at their complementary nature. Nonetheless, a similar analysis within the family of automatic evaluations is still pending; a study of the relations between open-ended and close-ended benchmarks and metrics, to understand which of these tests should be used, and when. For that purpose, we focus on the healthcare domain, providing the following contributions:

- A correlation-based, empirical analysis of

Tasks	Metrics	Datasets	
Multiple choice questions	Accuracy	· MedMCQA (et al., 2022)
· MedQA (et al., 2020b)
· ANONIMIZED-Close	· PubMedQA (et al., 2019)
· MMLU (et al., 2020a)
Prescriptions writing	"	· Prescription	
Medical text classification	"	· Medical Text for classification (Schopf et al., 2023)
· Medical Transcriptions	
Relation extraction	"	· BioRED (Luo et al., 2022)	

	Open-ended		
Open-ended medical questions	BLEU, BLEURT, ROUGE, BERTScore, MoverScore, Prometheus, Perplexity	· MedDialog Raw (Zeng et al., 2020)
· MEDIQA2019 (Ben Abacha et al., 2019)
· ANONIMIZED-Open	
Making diagnosis and treatment recommendations	"	· MedText	
Clinical note-taking	"	· MTS-Dialog (Ben Abacha et al., 2023)
· ACI-Bench (Yim et al., 2023)	
Medical factuality	"
+ Relaxed Perplexity	· OLAPH (Jeong et al., 2024)	
Summarization	"
+ F1-RadGraph	· MIMIC-III (Johnson et al., 2016)	
Question entailment	"	· Meddialog Qsumm (Zeng et al., 2020)	

Table 1: This table presents the tasks implemented in this paper. The first column specifies the different tasks. The second details the metrics used to evaluate performance in each task. Perplexity includes Bits per Byte, Byte Perplexity, and Word Perplexity. The third column outlines the different benchmarks used for each task.

open-ended and close-ended tasks, benchmarks, and metrics.

- A novel medical benchmark (ANONIMIZED) featuring both closed- and open-ended formats for the verification of our findings.
- A new metric for open-ended evaluations (Relaxed Perplexity) which fills a gap identified in existing benchmarks.

2 Methodology

This study considers five different close-ended healthcare tasks, which include nine different datasets (*e.g.*, MedQA). These are all assessed using the accuracy metric. At the same time, six open-ended tasks are studied, based on nine distinct datasets (*e.g.*, MedText). In this case, eleven different metrics are extracted. Further details are shown in Table 1. To assess the consistency within tasks, datasets and metrics, this work considers up to 14 different open LLMs, both specifically tuned for healthcare and general purpose, motivated by previous work (Shoham and Rappoport, 2024; Kanithi et al., 2024).

2.1 ANONIMIZED: A Novel Benchmark

Updated benchmarks are necessary to prevent both data drift (as human knowledge evolves), and data contamination (as train data crawling efforts scale).

To validate the integrity and consistency of existing tests, this work introduces a new benchmark for automatic evaluation, ANONIMIZED, available in both closed-ended and open-ended format.

ANONIMIZED originates from the Spanish Specialised Healthcare Training (MIR) exams by the Spanish *Ministry of Health*. The close-ended version is a MCQA including 5,621 QA pairs across six categories: medicine, nursing, biology, chemistry, psychology, and pharmacology, sourced from the 2020 to 2024 exam editions. ANONIMIZED is available in both English and Spanish.

The open-ended version (English only) was created by rephrasing the questions from the close-ended version using the Qwen2.5-72B-Instruct model. After the rephrasing process, the number of suitable questions was reduced to 3,730 QA pairs. This set retains the same categories as the closed-ended version. See Appendix A for further details.

2.2 Metrics

For close-ended evaluations, the metric of choice is accuracy. In contrast, for open-ended queries, there is a variety of metrics which provide different insights into model performance. This work considers eleven of those, which are sorted into four distinct categories:

- **N-gram based metrics** evaluate the over-

lap of n-grams between the generated and reference answers. This category includes: ROUGE-1, ROUGE-2, ROUGE-L and BLEU.

- **Semantic similarity metrics** evaluate the semantic similarity between the generated text and reference text, often leveraging embeddings or deep learning models. This includes: BERTScore, BLEURT and MoverScore.

- **Perplexity metrics** assess the predictive capabilities of the model by measuring how well it can predict a sequence of words. This includes: Word Perplexity, Bits per Byte and Byte Perplexity.

- **LLM-judge**: In this category we use the Prometheus (Kim et al., 2024) model to grade responses based on specific scoring criteria.

3 Experimentation

3.1 Correlation of open-ended vs close-ended

The first experiment conducted studies the correlation between open-ended and close-ended tasks, as detailed in Table 1. Specifically, we compare the weighted average of accuracy from the various MCQA benchmarks against every other close-ended open-ended task and metric in Figure 1.

Of all close and open-ended tasks, only clinical note-taking correlates positively with MCQA, and even in this case, correlation is rather weak. In contrast, summarization, question entailment and the remaining close-ended benchmarks correlate negatively with MCQA (the rest show a generalized lack of correlation). The negative correlation is explained by the lack of medical expertise needed for summarizing and entailing (as information is available in the input), and by the diverse nature of close-ended tasks. At metric level, all open alternatives correlate very weakly with MCQA.

These findings illustrate the relevance of the benchmarks chosen for evaluation, as well as the complementary nature of MCQA, when considering other tasks like summarization or clinical note-taking. Further details in Appendix B.1.

3.2 Correlation of open-ended benchmarks

The previous section locates open-ended tasks with a variable degree of correlation with close-ended tasks (*e.g.*, clinical note-taking, summarization). Let us know analyze correlations within the open-ended category. Details on this are shown in Appendix B.3.

Figure 1: Correlation between medical multiple choice question benchmarks (Medical MCQA) and the rest of open-ended benchmarks.

Notably, no consistently high correlation is observed for any benchmark or task. This suggests that each benchmark measures distinct aspects of model performance. This is the case even for benchmarks tackling the same task (*e.g.*, ACI-Bench and MTS-Dialog), illustrating the importance of benchmark source (*i.e.*, who crafted the benchmark and in which context). This underscores the need for specialized evaluations for downstream tasks, as generalization cannot be assumed.

3.3 Correlation of open-ended metrics

To assess whether the metrics used in the open evaluation are correlated among themselves, and to simplify future analyses for practitioners, we conduct a correlation analysis for each of the metrics detailed in §2.2 across all implemented open-ended benchmarks (more details in Appendix B.2).

This analysis identifies three distinct clusters of highly correlated metrics. The first cluster includes the perplexity metrics, (*i.e.*, Word Perplexity, Bits per Byte, and Byte Perplexity) all of which show a correlation above 0.96 across all analyzed benchmarks. Noticeably, these metrics are all based on probabilistic prediction (perplexity) and informa-

tion efficiency (Bits per Byte). The results obtained from Prometheus (an LLM judge) can be considered a distinct cluster of evaluation, illustrating how an external model provides a different and rather unique perspective on model performance. Finally, the third cluster includes all n-gram-based metrics, together with semantic similarity metrics (*i.e.*, BERTScore, BLEURT, and MoverScore). A strong correlation among these metrics is consistently observed across benchmarks, which can be attributed to their shared focus on content and overall text quality.

3.4 Metrics resilience to rephrasing

A limitation of open-ended evaluations is their sensitivity to rewording. Let us now analyze the different metrics under this Opensetup, to better understand their reliability. To do so, the model's output are rephrased, and evaluation recomputed. Six rephrased versions are produced using Qwen2.5-72B-Instruct.

Results show most n-gram-based metrics (*i.e.*, ROUGE1, ROUGE2, and ROUGEL) are resilient to rephrasing (unlike BLEU). This difference may arise because ROUGE metrics rely on surface-level word matching, making them less sensitive to phrasing changes as long as the core vocabulary remains intact. *i.e.*, in healthcare texts, key terms like 'diagnosis,' 'treatment,' or medication names often stay consistent, allowing these metrics to maintain a high overlap. In contrast, Prometheus (LLM judge) is the most affected by rewording, which is reasonable considering that, for this evaluation, correct punctuation and formatting in the answers greatly improve scores. This metric is followed by BLEURT and BERTScore (model similarity based) as the least resilient. More details on this analysis can be found in Appendix C.

4 Relaxed perplexity: A novel metric

By being optimized for next token prediction on the ground truth, LLM's are optimized for perplexity. However, as seen before, this does not necessarily entail good performance on open or close-ended downstream tasks. Additionally, perplexity can be greatly impacted by instruct-tuning and alignment techniques (Lee et al., 2024).

To evaluate factuality in open-ended benchmarks, with no dependence on confounders or exact formulation, we propose Relaxed-Perplexity.

Given a *question* and a *target*, we wish to estimate

$$\mathbb{P}(target \sim \text{model} \mid question) =$$
$$= \mathbb{P}(A_0) + \ldots + \mathbb{P}(A_n \mid B_n)$$

that is, the probability that the target is sampled from the model given the prompt, at any time in the completion. We denote the events $A_n \equiv \{target \sim \text{model}(question + seq_n)\}$ and $B_n \equiv \{seq_n \sim \text{model}(question)\}$ for any seq_n of n tokens that comes from the model before the target. B_n represents the probability distribution of all n-token model answers. We can estimate this as $\mathbb{P}(A \mid B_n) \sim \mathbb{P}(A|seq_n^{i_1}) + \ldots + \mathbb{P}(A|seq_n^{i_\ell})$ for the ℓ more likely n-token sequences sampled from the model given *question*, because the events seq_n^i and seq_n^j are mutually exclusive. Using this, we can define Relaxed Perplexity as

$$\text{Relaxed-Perplexity}(target, question, model) =$$
$$= \exp\left(-\frac{1}{n + len(target)} \sum_{i=0}^{n} logP(A_i|B_i)\right)$$

This allows to evaluate correctness in the model's answers probability distribution, with no regard for the exact formulation. We thus test it with the Olaph (Jeong et al., 2024) medical factuality dataset. In contrast to perplexity, we observe that Relaxed-Perplexity assigns higher scores to models fine-tuned on healthcare datasets. More details on the implementation and results of Relaxed perplexity can be found in Appendix D.

5 Conclusions

This study finds very weak correlations between close-ended and open-ended benchmarks. Even within benchmarks of the same task, correlations are weak. This points at the limited insights provided by individual tests, and advocates for broader evaluation setups. The experimentation shown identifies three clusters of metrics that should be considered when evaluating medical models: (1) perplexity metrics, (2) n-gram-based metrics combined with semantic similarity metrics, and (3) LLM-as-a-judge metrics. Noticeably, none of them correlates strongly with the close-ended MCQA evaluation.

The introduced `ANONIMIZED` benchmark, featuring both closed- and open-ended formats, serves as a sanity check of existing tests, while the proposed Relaxed-Perplexity metric fills a gap in evaluation, by focusing on factuality while being resistant to exact formulations in an open-ended setting.

6 Limitations

Since this study is based on specific models, the findings may not generalize to other LLM architectures. Additionally, the quality and diversity of the datasets used for evaluation are limited, meaning these benchmarks may not fully capture the performance of LLMs across the broader healthcare landscape. While metrics and benchmarks can indicate how well LLMs perform on certain tasks, they may not reflect the complexities of integrating LLMs into real-world healthcare practices.

Regarding the new benchmark introduced, although subject matter experts created the original exam materials, which underwent public scrutiny, `ANONIMIZED` has not been subjected to formal bias assessment. Consequently, it may not adequately represent the full spectrum of medical knowledge or encompass all possible patient demographics. Furthermore, the open-ended version has not undergone thorough evaluation by healthcare experts, raising the possibility of errors or biases introduced by the LLM used to rephrase the questions. Therefore, we advise users to exercise caution when interpreting and generalizing the results.

All experiments are conducted on English benchmarks, and generalization to other languages has not been considered. To enable reproducibility, all resources (datasets and code) are made available in ANONIMIZED_URL.

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A Novel Benchmarks

A.1 ANONIMIZED (close-ended)

ANONIMIZED is a novel benchmark for evaluating healthcare Large Language Models (LLMs) through multiple-choice question answering. ANONIMIZED was created by collecting exam materials in PDF format from the official Spanish government website. These documents were automatically parsed and then underwent post-processing to ensure data quality. This process involved removing 23 inaccurately parsed instances and excluding officially impugned questions. To enhance global accessibility, the original Spanish questions were translated into English using GPT-4, with a subset of 60 instances manually reviewed for translation accuracy.

Each ANONIMIZED sample contains metadata including a numeric exam identifier, full question text, four answer options, correct answer, exam year, and specialization category. The dataset is available in both Spanish and English, facilitating cross-lingual research. Examples of ANONIMIZED samples are provided in Figure 2 and Table 2.

Figure 2: ANONIMIZED example from Medicine category.

While ANONIMIZED shares its source with HeadQA in the Spanish Specialised Healthcare Training (MIR) exams, there is no overlap between the datasets. ANONIMIZED expands upon its predecessor, covering the years 2020 to 2024 and comprising 5,621 question-answer test pairs, compared to HeadQA's 2,742 test pairs from 2013 to 2017. The dataset's composition is illustrated in Figure 3, showing the category distribution by year to reveal potential temporal trends in exam content.

Table 3 presents additional information about the dataset, including the total number of questions per category, the longest and average question and answer lengths (in tokens), and the overall vocabulary size. This comprehensive overview of ANONIMIZED's structure and content demonstrates its potential as a valuable resource for evaluating and improving healthcare-focused language models.

A.2 ANONIMIZED (open-ended)

To create the open-ended dataset, we started with the closed-ended version of the ANONIMIZED dataset, specifically focusing on the English set. The first step was to filter out questions that contained terms such as "incorrect," "except," "false," "not correct", or "NOT", as these terms indicate that the questions focus on identifying incorrect answers among the provided options. After this filtering, we rephrased the remaining questions into an open-ended format using the Qwen2.5-72B-Instruct model, specifically instructing it to only rephrase questions that could be effectively transformed. This process excluded questions that explicitly ask for incorrect options or require a selection from the provided answers. We employed two different prompts for rephrasing, followed by a selection process to determine the best rephrased version or to discard the question if neither was suitable.

Initially, the close-ended ANONIMIZED contained 5,621 QA pairs, but after the rephrasing process, the number of suitable questions for the openeended version was reduced to 3,730 QA pairs. This new dataset retains the same categories as the closed-ended version, including medicine, nursing, biology, chemistry, psychology, and pharmacology. And it encompasses questions sourced from official exam materials covering the years 2020 to 2024. Figure 4 illustrates the distribution of the 3,730 QA pairs in the open-ended version and examples of QA pairs from both the close-ended and open-ended versions of the ANONIMIZED dataset are shown in Table 4. Both datasets are publicly available at [1]

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[1] ANONIMIZED

Question	Option 1	Option 2	Option 3	Option 3	Year	Category
The Glisson's capsule covers:	Spleen.	**Liver.**	Kidney.	Lung.	2024	Biology
Cardiolipin is a:	Sphingolipid.	**Phosphoglyceride.**	Steroid.	Ganglioside.	2020	Biology
The cinnamic acid is a:	Terpene.	Fatty acid.	Flavonoid.	**Phenylpropanoid.**	2021	Chemistry
Which of the following acids is strongest?:	HCl.	**HI.**	H2SO4.	HNO3.	2023	Chemistry
Indicate the ketogenic amino acid:	Cysteine.	Glutamine.	Methionine.	**Lysine.**	2020	Pharmacology
O2 and O3 are examples of:	Isotopes.	**Allotropes.**	Isomers.	Conformers.	2023	Pharmacology
Malignant hyperthermia is not related to:	Succinylcholine.	Desflurane.	**Propofol.**	Sevoflurane.	2024	Medicine
The most common benign tumors of the esophagus are:	Fibrovascular polyps.	**The leiomyomas.**	Squamous papillomas.	The hemangiomas.	2021	Medicine
Which opioid presents a higher analgesic potency?	Morphine.	Methadone.	Meperidine.	**Fentanyl.**	2023	Nursing
Indicate the antidote for ethylene glycol:	Methylene blue.	**Fomepizole.**	Carnitine.	Dimercaprol.	2024	Nursing
Olfactory hallucinations are more common in:	Delirium.	Manic episode.	**Epilepsy.**	Alcoholic hallucinosis.	2022	Psychology
What kind of drug is quetiapine?	A benzodiazepine.	An anxiolytic.	An antidepressant.	**An antipsychotic.**	2020	Psychology

Table 2: Examples of `ANONIMIZED` samples. Correct options are marked in bold. Questions were selected based on length for space reasons.

Figure 3: Category distribution per Category and Year.

	ANONIMIZED					
	QA Pairs	**Max Q tokens**	**Avg Q tokens**	**Max A tokens**	**Avg A tokens**	**Vocab**
Medicine	857	202	48.57	43	9.65	9626
Nursing	923	96	24.61	70	12	9113
Pharmacology	969	147	18.94	56	8.51	7906
Biology	966	51	12.82	48	6.6	6300
Psychology	962	208	22.60	67	9.92	7573
Chemistry	944	81	16.88	47	8.2	6022

Table 3: `ANONIMIZED` dataset statistics, where Q and A represents the Question and Answer respectively.

Figure 4: Category distribution per Category and Year.

	Close-ended	**Open-ended**	**Category**
Question	The best way to estimate the relative strength of hydrogen bonds between the molecules of halogen hydrides, H-X, is by measuring:	What is the best way to estimate the relative strength of hydrogen bonds between the molecules of halogen hydrides, H-X?	Chemistry
Answer	The enthalpies of vaporization	The enthalpies of vaporization.	
Question	Taking into account the general principles regarding the minimum interval between the non-simultaneous administration of vaccines, identify the minimum interval between 2 attenuated vaccines:	What is the minimum interval recommended between the non-simultaneous administration of two attenuated vaccines, according to general principles?	Nursing
Answer	Four weeks.	Four weeks.	
Question	We evaluated in the emergency room an adult person who is irritable, yawning, complaining of muscle pain and cramps. They are nauseous and have notable tearing. The pupils are dilated. Which of the following is the most probable diagnosis?	An adult patient presents to the emergency room with irritability, yawning, muscle pain and cramps, nausea, notable tearing, and dilated pupils. What is the most probable diagnosis based on these symptoms?	Medicine
Answer	Opioid abstinence.	Opioid abstinence.	

Table 4: Examples of QA pairs: On the left, the close-ended version from `ANONIMIZED`, and on the right, the open-ended version.

B Correlations

B.1 Correlations between MCQA and Elo results

We perform a correlation analysis on the performance results of the medical MCQA benchmarks listed in Table 1. Additionally, we include Elo scores from the Chatbot Arena[2], a crowdsourcing platform that collects pairs of model-generated answers in response to user prompts, where the user selects the winning model based on their criteria.

We conducted a correlation analysis using both small and medium models. The small models used for the correlation shown in Figure 5 are as follows: gemma-2-9b-it (Team, 2024), Meta-Llama-3.1-8B-Instruct(AI@Meta, 2024), Mistral-7B-Instruct-v0.2, Mistral-7B-Instruct-v0.3, Phi-3-mini-4k-instruct, Phi-3-medium-4k-instruct, Qwen1.5-7B-Chat, Starling-LM-7B-beta, Starling-LM-7B-beta and Yi-1.5-9B-Chat. And the medium models used in Figure 6 are as follows: Athene-70B(Frick et al., 2024), tulu-2-dpo-70b(Ivison et al., 2023), Yi-1.5-34B-Chat, gemma-2-27b-it, Llama-3.1-70B-Instruct, Mixtral-8x7B-Instruct-v0.1, Qwen2-72B-Instruct(Yang et al., 2024), and WizardLM-70B-V1.0

From this analysis, we found that MedQA, MedMCQA, `ANONIMIZED`, and MMLU are highly correlated with one another. However, PubMedQA exhibits a noticeably lower correlation with the other medical benchmarks, particularly in smaller models.

Regarding the Elo scores, we observe a moderate correlation with the MCQA benchmarks, with the correlation being significantly stronger for larger models. This is likely due to larger models' ability to produce more coherent responses. Non-expert evaluators, such as those in the Elo scoring system, may favor responses that are well-structured and fluent, even if they lack precise medical accuracy. As a result, this preference for more polished answers could lead to a higher correlation with MCQA performance.

Figure 5: Comparison of correlations between MCQA benchmarks and ELO results for small models.

Figure 6: Comparison of correlations between MCQA benchmarks and ELO results for medium models.

[2]https://lmarena.ai/

Figure 7: This correlation matrix illustrates the relationships among the different open-ended metrics used to evaluate the benchmark for diagnosis and treatment recommendations. Three distinct clusters of metrics are identified: (1) perplexity metrics, (2) n-gram and semantic similarity metrics, and (3) Prometheus metrics.

B.2 Correlation between metrics

In this correlation analysis, we fix the open-ended benchmark and examine the correlations across the various computed metrics. Figure 7, presents the correlation matrix for the benchmark focused on making diagnosis and treatment recommendations, highlighting the three clusters of metrics identified in the paper. This correlation matrix was also computed for the rest of benchmarks revealing three similar clusters. The matrices were computed using the following models: BioMistral-MedMNX, JSL-MedLlama-3-8B-v2.0, Phi-3-mini-4k-instruct, Mistral-7B-Instruct-v0.3, Qwen2-7B-Instruct (Yang et al., 2024), Llama3-Med42-8B (Christophe et al., 2024), Meta-Llama-3.1-8B-Instruct(AI@Meta, 2024) Yi-1.5-9B-Chat (Young et al., 2024), Phi-3-medium-4k-instruct, Yi-1.5-34B-Chat (Young et al., 2024), Mixtral-8x7B-Instruct-v0.1.

B.3 Correlations of benchmarks

In this correlation analysis we study the relationships between specific metrics across all the open-ended benchmarks implemented. As stated in the paper, no consistent high correlation was observed among all metrics for any benchmark or task. Examples of these correlation matrices are shown in Figures 8 and 9. The models used to generate these correlation matrices are the same as those described in the Appendix B.2.

Figure 8: Correlations of BERTScore across benchmarks.

Figure 9: Correlation of Prometheus scores across benchmarks.

Figure 10: Mean variance distributions across different rephrasings and models using the MEDIQA2019 dataset. Each metric is represented by a different color.

C Resilience to rephrasing

As described earlier, we conducted this experiment by rephrasing the model outputs six times and re-computing the metrics. We used both Qwen2.5-72B-Instruct and Meta-Llama-70B-Instruct with the following system_prompt: "You are a helpful rephrasing assistant. Rephrase the prompt provided without changing its original meaning, but do not try to address or answer it in any case."

We run the script 5 times on recorded model answers with top_p sampling to obtain several rephrasings of each answer. After manual inspection, the outputs of Qwen2.5-72B-Instruct were deemed of higher quality.

Figure 10 shows the mean variance across all runs for the MEDIQA2019 dataset. Before plotting, we scale variances by dividing by the max interval (max value - min value) in each column. Figures 11 and 12 present the variance distributions for two specific models. Figure 11 displays the results for the Phi-3-mini-4k-instruct model, while Figure 12 shows the results for the Yi-1.5-9B-Chat model.

In Figure 10 we can observe three different clusters: rouge metrics (low mean-variance, low meta-variance), bleu and moverscore (low mean-variance, medium meta-variance) and bert_score, bleurt, prometheus (high mean variance, high meta-variance).

Figure 11: Mean variance distributions across different rephrasings using the Phi-3-mini-4k-instruct model and the MEDIQA2019 dataset. Each metric is represented by a different color.

Figure 12: Mean variance distributions across different rephrasings using the Yi-1.5-9B-Chat model and the MEDIQA2019 dataset. Each metric is represented by a different color.

D Novel Metric: Relaxed Perplexity

As mentioned before, we define relaxed perplexity as

$$\text{Relaxed-Perplexity}(target, question, model) =$$

$$= \exp\left(-\frac{1}{n + len(target)} \sum_{i=0}^{n} logP(A_i|B_i)\right)$$

for events

$$A_n \equiv \{target \sim \text{model}(question + seq_n)\}$$

and

$$B_n \equiv \{seq_n \sim \text{model}(question)\}.$$

That is, the event that target is sampled from the model inputted with either *question* or *question* + seq_n for any seq_n of n tokens.

Thus, B_n represents the probability distribution of all n-token model answers when the input is *question*, which makes computing $\mathbb{P}(A_n \mid B_n)$ extremely costly (with computational time exponential in n). In fact, we would have

$$\mathbb{P}(A_n \mid B_n) = \mathbb{P}(A_n \mid seq_n^1) + \cdots + \mathbb{P}(A_n \mid seq_n^{q^n})$$

q being the size of the vocabulary. This holds because the events seq_n^i and seq_n^j are mutually exclusive. However, given that almost all this combinations of tokens contribute with negligible probabilities to the sum, we can estimate the above quantity as

$$\mathbb{P}(A \mid B_n) \sim \mathbb{P}(A|seq_n^{i_1}) + \ldots + \mathbb{P}(A|seq_n^{i_\ell})$$

for the ℓ more likely n-token sequences sampled from the model given *question*, which can be computed efficiently using beam search, diverse beam search (Vijayakumar et al., 2016) or top-p sampling.

Relaxed Perplexity is specifically designed to evaluate factuality in the answers, with no regard for the exact formulation. We thus test it with the Olaph (Jeong et al., 2024) dataset, and note that for more effective evaluation of other open-ended benchmarks, some preprocessing of the ground truths must be carried out.

For our experiments we use top-p sampling, selecting the $m \in \{5, 10\}$ best sentences in a search space of $s \in \{10, 100\}$. We observe similar results with all combinations, and so fix $m = 5$ and $s = 10$ for better performance.

Figure 13: Correlation between OLAPH - Relaxed Perplexity and the rest of benchmarks

The implementation is built using vllm[3], which provides tools for efficient LLM inference (Kwon et al., 2023). It remains as future work to implement Relaxed Perplexity with beam search.

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[3]https://github.com/vllm-project/vllm

Figure 14: Ranking results for the various metrics are presented, where the top position is ranked as 1 and the lowest as 11. Different models are represented in distinct colors.

Question	Must have	Nice to have	Benchmark	Relaxed-Logprobs
A 50-year-old male presents with a history of recurrent kidney stones and osteopenia. He has been taking high-dose vitamin D supplements due to a previous diagnosis of vitamin D deficiency. Laboratory results reveal hypercalcemia and hypercalciuria. What is the likely diagnosis, and what is the treatment?	Vitamin D toxicity	Stop vitamin D supplementation	Medtext	[-2.055, -8.229] - [-2.639, -4.142]
Are benign brain tumors serious?	Benign brain tumors are not cancerous and do not spread or invade surrounding tissues.	Benign brain tumors grow slowly and often have clear boundaries.	OLAPH	[-12.825, -15.7796] -[-11.208, -16.580]
We evaluated in the emergency room an adult person who is irritable, yawning, complaining of muscle pain and cramps. They are nauseous and have notable tearing. The pupils are dilated. What is the most probable diagnosis?	Opioid withdrawal.	Possibly other substance withdrawal symptoms.	ANONIMIZED-Open	[-4.2512, -24.7192] -[-5.812, -26.883]

Table 5: Open-ended evaluation using Relaxed Perplexity on samples from MedText, OLAPH and ANONIMIZED-Open on Mistral-7B-Instruct-v0.3 (first) and JSL-MedLlama-3-8B-v2.0 (second).

| Model | Open-ended Medical Questions | | | | | | | | |
| | ANONIMIZED-Open | | | MedDialog Raw | | | MediQA2019 | | |
	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓
BioMistral-MedMNX	1.385	2.612	688.279	1.043	2.060	74.760	0.416	1.335	6.044
JSL-MedLlama-3-8B-v2.0	1.761	3.389	4050.635	1.179	2.265	131.509	0.517	1.431	9.312
Llama3-Med42-8B	1.331	2.516	534.246	1.069	2.097	83.115	0.405	1.324	5.754
Meta-Llama-3.1-70B-Instruct	1.274	2.418	406.877	0.993	1.991	60.907	0.245	1.185	2.886
Meta-Llama-3.1-8B-Instruct	1.346	2.543	573.612	1.060	2.085	80.124	0.430	1.347	6.407
Mistral-7B-Instruct-v0.3	1.411	2.659	777.203	1.073	2.104	84.603	0.420	1.338	6.145
Mixtral-8x7B-Instruct-v0.1	4.405	21.184	nan	1.028	2.039	70.258	0.300	1.232	3.662
Phi-3-medium-4k-instruct	1.298	2.458	455.98	1.068	2.097	82.957	0.410	1.329	5.884
Phi-3-mini-4k-instruct	1.299	2.460	458.481	1.082	2.117	87.936	0.444	1.360	6.796
Qwen2-7B-Instruct	1.438	2.710	885.33	1.044	2.063	75.218	0.447	1.363	6.895
Yi-1.5-34B-Chat	1.512	2.852	1254.136	1.101	2.145	95.042	0.485	1.399	8.112
Yi-1.5-9B-Chat	1.603	3.037	1921.553	1.123	2.178	104.205	0.532	1.446	9.968

Table 6: Perplexity results for Open-ended Medical Questions.

| Model | Clinical Note-taking | | | | | |
| | ACI Bench | | | MTS Dialog | | |
	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓
BioMistral-MedMNX	0.601	1.517	13.894	1.059	2.083	132.827
JSL-MedLlama-3-8B-v2.0	0.703	1.628	21.725	1.099	2.143	160.188
Llama3-Med42-8B	0.485	1.399	8.357	1.060	2.085	133.416
Meta-Llama-3.1-70B-Instruct	-	-	-	0.984	1.978	93.943
Meta-Llama-3.1-8B-Instruct	0.612	1.529	14.618	1.074	2.105	142.211
Mistral-7B-Instruct-v0.3	0.596	1.512	13.628	1.053	2.074	129.076
Mixtral-8x7B-Instruct-v0.1	0.566	1.481	11.933	1.046	2.064	125.070
Phi-3-medium-4k-instruct	0.642	1.560	16.600	0.971	1.960	88.447
Phi-3-mini-4k-instruct	0.599	1.514	13.754	0.972	1.962	89.163
Qwen2-7B-Instruct	0.619	1.535	15.009	1.063	2.089	135.111
Yi-1.5-34B-Chat	0.728	1.657	24.270	1.099	2.143	160.265
Yi-1.5-9B-Chat	0.711	1.636	22.456	1.180	2.265	232.073

Table 7: Perplexity results for clinical note-taking.

| Model | Making treatment recommendations | | | Question Entailment | | | Summarization | | |
| | MedText | | | MedDialog Qsumm | | | Mimic-III | | |
	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓	Bits per Byte ↓	Byte Perplexity ↓	Word Perplexity ↓
BioMistral-MedMNX	0.499	1.413	10.605	1.471	2.772	275.846	1.771	3.413	4697.580
JSL-MedLlama-3-8B-v2.0	0.556	1.470	13.868	1.715	3.282	699.785	2.035	4.099	16607.943
Llama3-Med42-8B	0.455	1.370	8.593	1.359	2.564	179.527	1.839	3.577	6489.224
Meta-Llama-3.1-70B-Instruct	0.447	1.364	8.298	1.280	2.428	132.988	-	-	-
Meta-Llama-3.1-8B-Instruct	0.534	1.448	12.501	1.371	2.587	188.513	1.826	3.545	6106.099
Mistral-7B-Instruct-v0.3	0.510	1.424	11.163	1.447	2.727	251.938	1.790	3.457	5138.524
Mixtral-8x7B-Instruct-v0.1	0.491	1.405	10.194	1.370	2.586	187.912	1.679	3.202	3028.534
Phi-3-medium-4k-instruct	0.423	1.341	7.400	1.332	2.517	162.163	2.084	4.239	20901.351
Phi-3-mini-4k-instruct	0.438	1.355	7.956	1.311	2.481	149.718	1.902	3.737	8784.663
Qwen2-7B-Instruct	0.527	1.441	12.106	1.383	2.608	197.167	1.878	3.676	7839.132
Yi-1.5-34B-Chat	0.556	1.470	13.875	1.437	2.708	242.427	2.202	4.600	36704.322
Yi-1.5-9B-Chat	0.559	1.473	14.052	1.470	2.771	275.222	2.341	5.067	71436.330

Table 8: Perplexity results for the following tasks: making diagnosis and treatment recommendation, question entailment and summarization tasks.

Model	Treatment recommendations MedText	Question Entailment MedDialog Qsumm	Summarization Mimic-III	Clinical Note-Taking ACI Bench	Clinical Note-Taking MTS Dialog
			Prometheus ↑		
BioMistral-MedMNX	0.225 ± 0.063	0.163 ± 0.005	0.330 ± 0.016	0.273 ± 0.027	0.297 ± 0.009
JSL-MedLlama-3-8B-v2.0	0.263 ± 0.084	0.087 ± 0.004	0.298 ± 0.017	0.365 ± 0.031	0.172 ± 0.008
Llama3-Med42-8B	0.138 ± 0.062	0.241 ± 0.007	0.213 ± 0.016	0.157 ± 0.024	0.130 ± 0.008
Meta-Llama-3.1-70B-Instruct	0.062 ± 0.043	0.314 ± 0.007	0.342 ± 0.016	0.313 ± 0.026	0.281 ± 0.009
Meta-Llama-3.1-8B-Instruct	0.188 ± 0.063	0.156 ± 0.005	0.263 ± 0.015	0.245 ± 0.027	0.237 ± 0.008
Mistral-7B-Instruct-v0.3	0.050 ± 0.029	0.194 ± 0.006	0.187 ± 0.015	0.087 ± 0.018	0.055 ± 0.005
Mixtral-8x7B-Instruct-v0.1	0.075 ± 0.036	0.112 ± 0.005	0.252 ± 0.016	0.090 ± 0.017	0.198 ± 0.009
Phi-3-medium-4k-instruct	0.175 ± 0.064	0.168 ± 0.005	0.358 ± 0.017	0.190 ± 0.023	0.219 ± 0.008
Phi-3-mini-4k-instruct	0.125 ± 0.057	0.126 ± 0.005	0.376 ± 0.016	0.287 ± 0.027	0.280 ± 0.009
Qwen2-7B-Instruct	0.125 ± 0.052	0.177 ± 0.006	0.267 ± 0.014	0.255 ± 0.026	0.144 ± 0.007
Yi-1.5-34B-Chat	0.287 ± 0.069	0.179 ± 0.006	0.372 ± 0.016	0.342 ± 0.030	0.420 ± 0.008
Yi-1.5-9B-Chat	0.138 ± 0.067	0.405 ± 0.007	0.550 ± 0.015	0.362 ± 0.026	0.397 ± 0.008
gemma-2-27b-it	0.000 ± 0.000	0.001 ± 0.000	0.002 ± 0.001	0.000 ± 0.000	0.000 ± 0.000
gemma-2-9b-it	0.000 ± 0.000	0.002 ± 0.001	0.011 ± 0.004	0.000 ± 0.000	0.000 ± 0.000

Table 9: Prometheus results for the following tasks: making diagnosis and treatment recommendations, question entailment, summarization and clinical note-taking.

Model	Open-ended Medical Questions		
	MedDialog Raw	MediQA2019	ANONIMIZED-Open
		Prometheus ↑	
BioMistral-MedMNX	0.535 ± 0.005	0.342 ± 0.007	-
JSL-MedLlama-3-8B-v2.0	0.304 ± 0.005	0.459 ± 0.008	-
Llama3-Med42-8B	0.138 ± 0.062	0.241 ± 0.007	-
Meta-Llama-3.1-70B-Instruct	0.293 ± 0.005	0.326 ± 0.008	0.296 ± 0.006
Meta-Llama-3.1-8B-Instruct	0.375 ± 0.005	0.229 ± 0.007	-
Mistral-7B-Instruct-v0.3	0.476 ± 0.005	0.384 ± 0.008	0.087 ± 0.004
Mixtral-8x7B-Instruct-v0.1	0.543 ± 0.005	0.361 ± 0.008	0.030 ± 0.002
Phi-3-medium-4k-instruct	0.249 ± 0.005	0.281 ± 0.008	0.308 ± 0.007
Phi-3-mini-4k-instruct	0.353 ± 0.005	0.328 ± 0.008	0.299 ± 0.006
Qwen2-7B-Instruct	0.541 ± 0.005	0.267 ± 0.007	0.427 ± 0.007
Yi-1.5-34B-Chat	0.508 ± 0.005	0.347 ± 0.009	0.488 ± 0.007
Yi-1.5-9B-Chat	0.288 ± 0.005	0.417 ± 0.009	0.612 ± 0.006
gemma-2-27b-it	0.001 ± 0.000	0.000 ± 0.000	0.003 ± 0.001
gemma-2-9b-it	0.002 ± 0.000	0.003 ± 0.001	0.006 ± 0.001

Table 10: Prometheus results for the open-ended medical questions.

Model	Clinical Note-taking													
	ACI Bench							MTS Dialog						
	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑
BioMistral-MedMNX	0.839 ± 0.007	0.012 ± 0.005	-0.834 ± 0.057	0.537 ± 0.006	0.171 ± 0.016	0.039 ± 0.009	0.130 ± 0.014	0.800 ± 0.001	0.001 ± 0.000	-1.304 ± 0.006	0.493 ± 0.001	0.040 ± 0.000	0.003 ± 0.000	0.036 ± 0.001
JSL-MedLlama-3-8B-v2.0	0.853 ± 0.011	0.033 ± 0.016	-0.810 ± 0.143	0.549 ± 0.013	0.212 ± 0.050	0.083 ± 0.026	0.173 ± 0.040	0.801 ± 0.001	0.002 ± 0.000	-1.279 ± 0.007	0.492 ± 0.001	0.048 ± 0.001	0.006 ± 0.001	0.043 ± 0.001
Llama3-Med42-8B	0.863 ± nan	0.059 ± 0.019	-0.608 ± nan	0.564 ± nan	0.285 ± nan	0.114 ± nan	0.224 ± nan	0.803 ± 0.001	0.003 ± 0.001	-1.290 ± 0.011	0.495 ± 0.001	0.048 ± 0.002	0.007 ± 0.001	0.043 ± 0.002
Meta-Llama-3.1-70B-Instruct	0.852 ± nan	0.019 ± nan	-0.613 ± nan	0.548 ± nan	0.201 ± nan	0.056 ± nan	0.154 ± nan	0.798 ± 0.001	0.000 ± 0.000	-1.350 ± 0.007	0.492 ± 0.001	0.041 ± 0.001	0.002 ± 0.000	0.038 ± 0.001
Meta-Llama-3.1-8B-Instruct	0.829 ± 0.011	0.017 ± 0.007	-0.870 ± 0.068	0.538 ± 0.006	0.188 ± 0.024	0.047 ± 0.013	0.138 ± 0.019	0.797 ± 0.001	0.001 ± 0.000	-1.364 ± 0.007	0.490 ± 0.001	0.044 ± 0.001	0.003 ± 0.000	0.040 ± 0.001
Mistral-7B-Instruct-v0.3	0.812 ± nan	0.000 ± nan	-1.138 ± nan	0.522 ± nan	0.046 ± nan	0.004 ± nan	0.037 ± nan	0.800 ± 0.001	0.000 ± 0.000	-1.322 ± 0.007	0.491 ± 0.001	0.042 ± 0.001	0.002 ± 0.000	0.039 ± 0.001
Mixtral-8x7B-Instruct-v0.1	0.832 ± nan	0.013 ± 0.006	-0.881 ± nan	0.540 ± nan	0.168 ± nan	0.038 ± nan	0.119 ± nan	0.800 ± 0.001	0.001 ± 0.000	-1.349 ± 0.007	0.492 ± 0.001	0.042 ± 0.001	0.003 ± 0.000	0.039 ± 0.001
Phi-3-medium-4k-instruct	0.824 ± 0.007	0.014 ± 0.014	-1.005 ± 0.067	0.528 ± 0.005	0.111 ± 0.023	0.023 ± 0.011	0.086 ± 0.017	0.800 ± 0.001	0.001 ± 0.000	-1.346 ± 0.007	0.494 ± 0.001	0.040 ± 0.001	0.003 ± 0.000	0.037 ± 0.001
Phi-3-mini-4k-instruct	0.821 ± nan	0.015 ± 0.007	-1.026 ± nan	0.529 ± nan	0.135 ± nan	0.035 ± nan	0.111 ± nan	0.800 ± 0.001	0.000 ± 0.000	-1.312 ± 0.007	0.494 ± 0.001	0.039 ± 0.001	0.002 ± 0.000	0.036 ± 0.001
Qwen2-7B-Instruct	0.841 ± nan	0.015 ± 0.007	-0.861 ± nan	0.538 ± nan	0.167 ± nan	0.051 ± nan	0.133 ± nan	0.798 ± 0.001	0.000 ± 0.000	-1.334 ± 0.006	0.489 ± 0.001	0.040 ± 0.001	0.002 ± 0.000	0.037 ± 0.001
Yi-1.5-34B-Chat	0.840 ± 0.009	0.015 ± 0.009	-0.814 ± 0.085	0.533 ± 0.007	0.163 ± 0.024	0.046 ± 0.015	0.126 ± 0.019	0.806 ± 0.001	0.004 ± 0.001	-1.266 ± 0.011	0.498 ± 0.001	0.063 ± 0.003	0.012 ± 0.001	0.056 ± 0.002
Yi-1.5-9B-Chat	0.836 ± nan	0.030 ± 0.024	-0.892 ± nan	0.531 ± nan	0.159 ± nan	0.063 ± nan	0.140 ± nan	0.803 ± 0.001	0.003 ± 0.001	-1.320 ± 0.009	0.494 ± 0.001	0.053 ± 0.002	0.007 ± 0.001	0.048 ± 0.002
gemma-2-27b-it	0.815 ± nan	0.000 ± nan	-1.016 ± nan	0.521 ± nan	0.077 ± nan	0.000 ± nan	0.077 ± nan	0.790 ± 0.004	0.000 ± 0.000	-1.226 ± 0.036	0.504 ± 0.005	0.026 ± 0.008	0.000 ± 0.000	0.026 ± 0.008
gemma-2-9b-it	0.789 ± 0.005	0.000 ± 0.000	-1.081 ± 0.041	0.506 ± 0.004	0.080 ± 0.017	0.000 ± 0.000	0.061 ± 0.011	0.795 ± 0.001	0.000 ± 0.000	-1.231 ± 0.010	0.491 ± 0.001	0.041 ± 0.002	0.000 ± 0.000	0.040 ± 0.002

Table 11: Clinical note-taking results.

| Model | Making Treatment Recommendations | | | | | | |
| | Medtext | | | | | | |
	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑
BioMistral-MedMNX	0.855 ± 0.001	0.013 ± 0.001	-0.650 ± 0.007	0.547 ± 0.001	0.177 ± 0.002	0.037 ± 0.001	0.136 ± 0.002
JSL-MedLlama-3-8B-v2.0	0.856 ± 0.001	0.021 ± 0.002	-0.652 ± 0.012	0.546 ± 0.001	0.185 ± 0.003	0.045 ± 0.002	0.146 ± 0.003
Llama3-Med42-8B	0.865 ± 0.001	0.018 ± 0.002	-0.546 ± 0.015	0.557 ± 0.001	0.204 ± 0.005	0.052 ± 0.003	0.158 ± 0.004
Meta-Llama-3.1-70B-Instruct	0.859 ± 0.001	0.022 ± 0.002	-0.644 ± 0.008	0.547 ± 0.001	0.196 ± 0.003	0.048 ± 0.002	0.150 ± 0.002
Meta-Llama-3.1-8B-Instruct	0.843 ± 0.001	0.010 ± 0.001	-0.839 ± 0.007	0.535 ± 0.001	0.155 ± 0.002	0.032 ± 0.001	0.120 ± 0.002
Mistral-7B-Instruct-v0.3	0.870 ± 0.002	0.038 ± 0.005	-0.467 ± 0.022	0.562 ± 0.002	0.230 ± 0.008	0.072 ± 0.006	0.183 ± 0.007
Mixtral-8x7B-Instruct-v0.1	0.868 ± 0.001	0.029 ± 0.002	-0.502 ± 0.011	0.559 ± 0.001	0.220 ± 0.003	0.060 ± 0.002	0.172 ± 0.003
Phi-3-medium-4k-instruct	0.869 ± 0.001	0.033 ± 0.002	-0.504 ± 0.011	0.560 ± 0.001	0.231 ± 0.004	0.069 ± 0.003	0.182 ± 0.003
Phi-3-mini-4k-instruct	0.863 ± 0.001	0.027 ± 0.002	-0.551 ± 0.009	0.555 ± 0.001	0.213 ± 0.003	0.060 ± 0.002	0.165 ± 0.003
Qwen2-7B-Instruct	0.859 ± 0.001	0.021 ± 0.002	-0.634 ± 0.012	0.547 ± 0.001	0.193 ± 0.004	0.049 ± 0.002	0.147 ± 0.003
Yi-1.5-34B-Chat	0.867 ± 0.001	0.033 ± 0.002	-0.580 ± 0.008	0.559 ± 0.001	0.245 ± 0.003	0.074 ± 0.002	0.189 ± 0.002
Yi-1.5-9B-Chat	0.863 ± 0.000	0.022 ± 0.001	-0.513 ± 0.006	0.555 ± 0.001	0.213 ± 0.002	0.054 ± 0.002	0.163 ± 0.002
gemma-2-27b-it	nan ± nan	0.000 ± 0.000	-1.190 ± 0.024	0.495 ± 0.002	0.071 ± 0.007	0.003 ± 0.001	0.062 ± 0.005
gemma-2-9b-it	0.819 ± 0.001	0.000 ± 0.000	-1.146 ± 0.009	0.498 ± 0.001	0.077 ± 0.003	0.002 ± 0.000	0.067 ± 0.002

Table 12: Making diagnosis and treatment recommendations results.

| Model | Open-ended medical questions | | | | | | |
| | ANONIMIZED-Open | | | | | | |
	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑
BioMistral-MedMNX	0.835 ± 0.001	0.003 ± 0.000	-1.040 ± 0.007	0.517 ± 0.001	0.069 ± 0.001	0.017 ± 0.001	0.057 ± 0.001
JSL-MedLlama-3-8B-v2.0	0.844 ± 0.001	0.007 ± 0.001	-0.908 ± 0.010	0.527 ± 0.001	0.083 ± 0.002	0.025 ± 0.001	0.070 ± 0.002
Llama3-Med42-8B	0.839 ± 0.001	0.005 ± 0.001	-0.989 ± 0.012	0.521 ± 0.001	0.090 ± 0.003	0.022 ± 0.002	0.074 ± 0.002
Meta-Llama-3.1-70B-Instruct	0.843 ± 0.001	0.007 ± 0.001	-0.969 ± 0.012	0.526 ± 0.001	0.134 ± 0.003	0.030 ± 0.002	0.113 ± 0.003
Meta-Llama-3.1-8B-Instruct	0.830 ± 0.001	0.003 ± 0.000	-1.128 ± 0.007	0.514 ± 0.001	0.065 ± 0.001	0.017 ± 0.001	0.052 ± 0.001
Mistral-7B-Instruct-v0.3	0.849 ± 0.002	0.006 ± 0.002	-0.892 ± 0.023	0.529 ± 0.002	0.145 ± 0.007	0.038 ± 0.004	0.129 ± 0.006
Mixtral-8x7B-Instruct-v0.1	0.814 ± 0.001	0.000 ± 0.000	-1.264 ± 0.007	0.492 ± 0.001	0.051 ± 0.002	0.002 ± 0.000	0.045 ± 0.002
Phi-3-medium-4k-instruct	0.847 ± 0.001	0.008 ± 0.001	-0.883 ± 0.013	0.529 ± 0.001	0.145 ± 0.003	0.034 ± 0.002	0.122 ± 0.003
Phi-3-mini-4k-instruct	0.839 ± 0.001	0.004 ± 0.001	-0.986 ± 0.009	0.520 ± 0.001	0.115 ± 0.003	0.022 ± 0.001	0.097 ± 0.002
Qwen2-7B-Instruct	0.844 ± 0.001	0.006 ± 0.001	-0.896 ± 0.009	0.527 ± 0.001	0.133 ± 0.003	0.028 ± 0.001	0.111 ± 0.002
Yi-1.5-34B-Chat	0.845 ± 0.001	0.008 ± 0.001	-0.887 ± 0.009	0.526 ± 0.001	0.141 ± 0.003	0.034 ± 0.001	0.119 ± 0.002
Yi-1.5-9B-Chat	0.843 ± 0.000	0.007 ± 0.000	-0.906 ± 0.007	0.525 ± 0.001	0.136 ± 0.002	0.030 ± 0.001	0.113 ± 0.002

Table 13: Results for ANONIMIZED-Open

| Model | Open-ended Medical Questions | | | | | | | | | | | | | |
| | MedDialog | | | | | | | MEDIQA2019 | | | | | | |
	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑
BioMistral-MedMNX	0.833 ± 0.001	0.001 ± 0.000	-0.898 ± 0.012	0.526 ± 0.001	0.113 ± 0.003	0.010 ± 0.001	0.088 ± 0.002	0.850 ± 0.002	0.005 ± 0.002	-0.660 ± 0.024	0.547 ± 0.002	0.169 ± 0.007	0.032 ± 0.003	0.132 ± 0.005
JSL-MedLlama-3-8B-v2.0	0.832 ± 0.001	0.000 ± 0.000	-0.875 ± 0.015	0.524 ± 0.001	0.109 ± 0.003	0.009 ± 0.001	0.087 ± 0.002	0.849 ± 0.002	0.008 ± 0.002	-0.688 ± 0.027	0.543 ± 0.002	0.164 ± 0.006	0.030 ± 0.003	0.130 ± 0.005
Llama3-Med42-8B	0.834 ± 0.001	0.000 ± 0.000	-0.887 ± 0.019	0.527 ± 0.001	0.108 ± 0.004	0.010 ± 0.001	0.085 ± 0.003	0.850 ± 0.003	0.008 ± 0.003	-0.646 ± 0.043	0.546 ± 0.004	0.166 ± 0.012	0.026 ± 0.005	0.129 ± 0.010
Meta-Llama-3.1-70B-Instruct	0.835 ± 0.001	0.000 ± 0.000	-0.875 ± 0.014	0.525 ± 0.001	0.115 ± 0.003	0.011 ± 0.001	0.089 ± 0.002	0.856 ± 0.002	0.010 ± 0.003	-0.630 ± 0.030	0.547 ± 0.002	0.176 ± 0.008	0.037 ± 0.004	0.139 ± 0.007
Meta-Llama-3.1-8B-Instruct	0.824 ± 0.001	0.000 ± 0.000	-1.013 ± 0.011	0.521 ± 0.001	0.096 ± 0.003	0.008 ± 0.001	0.074 ± 0.002	0.843 ± 0.002	0.005 ± 0.001	-0.775 ± 0.024	0.538 ± 0.002	0.154 ± 0.007	0.028 ± 0.003	0.117 ± 0.005
Mistral-7B-Instruct-v0.3	0.841 ± 0.001	0.000 ± 0.000	-0.762 ± 0.024	0.530 ± 0.001	0.121 ± 0.005	0.014 ± 0.002	0.095 ± 0.004	0.852 ± 0.004	0.016 ± 0.008	-0.661 ± 0.061	0.541 ± 0.005	0.158 ± 0.016	0.046 ± 0.011	0.132 ± 0.015
Mixtral-8x7B-Instruct-v0.1	0.838 ± 0.001	0.001 ± 0.000	-0.819 ± 0.020	0.529 ± 0.001	0.119 ± 0.004	0.012 ± 0.001	0.093 ± 0.003	0.846 ± 0.004	0.006 ± 0.003	-0.837 ± 0.058	0.536 ± 0.004	0.135 ± 0.015	0.022 ± 0.009	0.110 ± 0.014
Phi-3-medium-4k-instruct	0.837 ± 0.001	0.001 ± 0.000	-0.854 ± 0.016	0.528 ± 0.001	0.121 ± 0.003	0.013 ± 0.001	0.093 ± 0.003	0.859 ± 0.003	0.011 ± 0.004	-0.552 ± 0.042	0.551 ± 0.004	0.197 ± 0.013	0.049 ± 0.009	0.157 ± 0.012
Phi-3-mini-4k-instruct	0.834 ± 0.001	0.000 ± 0.000	-0.891 ± 0.015	0.526 ± 0.001	0.103 ± 0.003	0.009 ± 0.001	0.082 ± 0.002	0.850 ± 0.003	0.008 ± 0.004	-0.682 ± 0.036	0.543 ± 0.003	0.163 ± 0.011	0.032 ± 0.007	0.129 ± 0.009
Qwen2-7B-Instruct	0.833 ± 0.001	0.000 ± 0.000	-0.939 ± 0.015	0.526 ± 0.001	0.109 ± 0.004	0.010 ± 0.001	0.084 ± 0.003	0.851 ± 0.002	0.005 ± 0.002	-0.673 ± 0.031	0.542 ± 0.002	0.155 ± 0.008	0.029 ± 0.005	0.120 ± 0.007
Yi-1.5-34B-Chat	0.839 ± 0.001	0.000 ± 0.000	-0.785 ± 0.014	0.529 ± 0.001	0.131 ± 0.004	0.016 ± 0.001	0.101 ± 0.003	0.858 ± 0.002	0.008 ± 0.002	-0.524 ± 0.031	0.551 ± 0.003	0.185 ± 0.009	0.039 ± 0.005	0.147 ± 0.008
Yi-1.5-9B-Chat	0.837 ± 0.001	0.001 ± 0.000	-0.804 ± 0.012	0.528 ± 0.001	0.123 ± 0.003	0.014 ± 0.001	0.096 ± 0.002	0.857 ± 0.002	0.011 ± 0.003	-0.540 ± 0.026	0.549 ± 0.002	0.197 ± 0.007	0.043 ± 0.004	0.159 ± 0.006
gemma-2-27b-it	0.815 ± 0.003	0.000 ± 0.000	-1.209 ± 0.045	0.502 ± 0.003	0.049 ± 0.009	0.000 ± 0.000	0.045 ± 0.008	0.829 ± nan	0.000 ± 0.000	-1.134 ± nan	0.515 ± nan	0.093 ± nan	0.000 ± nan	0.085 ± nan
gemma-2-9b-it	0.818 ± 0.002	0.000 ± 0.000	-1.115 ± 0.018	0.510 ± 0.001	0.066 ± 0.004	0.003 ± 0.001	0.057 ± 0.004	0.822 ± 0.004	0.000 ± 0.000	-1.108 ± 0.045	0.513 ± 0.004	0.084 ± 0.010	0.004 ± 0.002	0.070 ± 0.007

Table 14: Open-ended medical questions results.

Model	Question Entailment						
	MedDialog						
	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑
BioMistral-MedMNX	0.839 ± 0.000	0.005 ± 0.000	-1.056 ± 0.003	0.520 ± 0.000	0.093 ± 0.001	0.018 ± 0.001	0.081 ± 0.001
JSL-MedLlama-3-8B-v2.0	0.840 ± 0.000	0.004 ± 0.000	-0.967 ± 0.004	0.522 ± 0.000	0.085 ± 0.001	0.013 ± 0.001	0.074 ± 0.001
Llama3-Med42-8B	0.845 ± 0.000	0.004 ± 0.000	-1.020 ± 0.005	0.521 ± 0.000	0.099 ± 0.002	0.019 ± 0.001	0.084 ± 0.001
Meta-Llama-3.1-70B-Instruct	0.849 ± 0.000	0.008 ± 0.001	-1.013 ± 0.005	0.525 ± 0.000	0.120 ± 0.002	0.029 ± 0.001	0.102 ± 0.001
Meta-Llama-3.1-8B-Instruct	0.836 ± 0.000	0.005 ± 0.000	-1.097 ± 0.004	0.518 ± 0.000	0.091 ± 0.001	0.017 ± 0.001	0.078 ± 0.001
Mistral-7B-Instruct-v0.3	0.852 ± 0.001	0.010 ± 0.001	-0.966 ± 0.007	0.526 ± 0.001	0.122 ± 0.003	0.031 ± 0.002	0.106 ± 0.002
Mixtral-8x7B-Instruct-v0.1	0.848 ± 0.001	0.004 ± 0.000	-0.984 ± 0.006	0.525 ± 0.000	0.099 ± 0.002	0.020 ± 0.001	0.086 ± 0.002
Phi-3-medium-4k-instruct	0.839 ± 0.000	0.004 ± 0.000	-1.086 ± 0.004	0.522 ± 0.000	0.093 ± 0.001	0.017 ± 0.001	0.081 ± 0.001
Phi-3-mini-4k-instruct	0.840 ± 0.000	0.003 ± 0.000	-1.041 ± 0.004	0.521 ± 0.000	0.083 ± 0.001	0.012 ± 0.001	0.072 ± 0.001
Qwen2-7B-Instruct	0.844 ± 0.000	0.006 ± 0.001	-1.007 ± 0.004	0.524 ± 0.000	0.102 ± 0.002	0.020 ± 0.001	0.088 ± 0.001
Yi-1.5-34B-Chat	0.842 ± 0.001	0.006 ± 0.001	-1.010 ± 0.005	0.522 ± 0.000	0.100 ± 0.002	0.021 ± 0.001	0.087 ± 0.002
Yi-1.5-9B-Chat	0.852 ± 0.000	0.010 ± 0.001	-0.979 ± 0.004	0.525 ± 0.000	0.128 ± 0.001	0.033 ± 0.001	0.109 ± 0.001
gemma-2-27b-it	0.817 ± 0.002	0.000 ± 0.000	-1.151 ± 0.016	0.513 ± 0.001	0.052 ± 0.006	0.002 ± 0.001	0.049 ± 0.006
gemma-2-9b-it	0.821 ± 0.001	0.000 ± 0.000	-1.135 ± 0.006	0.510 ± 0.001	0.057 ± 0.002	0.002 ± 0.000	0.053 ± 0.002

Table 15: Question entailment results.

Model	Summarization							
	MIMIC-III							
	F1-RadGraph ↑	BERTScore ↑	BLEU ↑	BLEURT ↑	MoverScore ↑	ROUGE1 ↑	ROUGE2 ↑	ROUGEL ↑
BioMistral-MedMNX	0.089 ± 0.001	0.837 ± 0.000	0.009 ± 0.000	-0.796 ± 0.003	0.551 ± 0.000	0.130 ± 0.001	0.031 ± 0.001	0.110 ± 0.001
JSL-MedLlama-3-8B-v2.0	0.079 ± 0.002	0.841 ± 0.000	0.014 ± 0.001	-0.780 ± 0.005	0.556 ± 0.001	0.143 ± 0.002	0.041 ± 0.001	0.124 ± 0.002
Llama3-Med42-8B	0.093 ± 0.002	0.843 ± 0.000	0.013 ± 0.001	-0.729 ± 0.005	0.557 ± 0.001	0.152 ± 0.002	0.041 ± 0.001	0.129 ± 0.002
Meta-Llama-3.1-70B-Instruct	0.059 ± 0.002	0.836 ± 0.000	0.009 ± 0.001	-0.811 ± 0.005	0.547 ± 0.001	0.130 ± 0.002	0.031 ± 0.001	0.110 ± 0.002
Meta-Llama-3.1-8B-Instruct	0.065 ± 0.001	0.830 ± 0.000	0.007 ± 0.000	-0.834 ± 0.004	0.542 ± 0.000	0.115 ± 0.001	0.025 ± 0.001	0.097 ± 0.001
Mistral-7B-Instruct-v0.3	0.082 ± 0.002	0.845 ± 0.000	0.013 ± 0.001	-0.753 ± 0.005	0.558 ± 0.000	0.157 ± 0.002	0.044 ± 0.001	0.134 ± 0.002
Mixtral-8x7B-Instruct-v0.1	0.088 ± 0.002	0.844 ± 0.000	0.015 ± 0.001	-0.762 ± 0.004	0.557 ± 0.000	0.157 ± 0.002	0.044 ± 0.001	0.134 ± 0.002
Phi-3-medium-4k-instruct	0.038 ± 0.002	0.838 ± 0.001	0.010 ± 0.001	-0.771 ± 0.008	0.550 ± 0.001	0.137 ± 0.003	0.034 ± 0.001	0.116 ± 0.002
Phi-3-mini-4k-instruct	0.066 ± 0.002	0.836 ± 0.000	0.008 ± 0.001	-0.767 ± 0.005	0.548 ± 0.001	0.123 ± 0.002	0.029 ± 0.001	0.104 ± 0.002
Qwen2-7B-Instruct	0.078 ± 0.001	0.843 ± 0.000	0.009 ± 0.000	-0.761 ± 0.004	0.555 ± 0.000	0.142 ± 0.002	0.035 ± 0.001	0.120 ± 0.001
Yi-1.5-34B-Chat	0.065 ± 0.001	0.839 ± 0.000	0.009 ± 0.001	-0.775 ± 0.004	0.550 ± 0.000	0.137 ± 0.002	0.033 ± 0.001	0.116 ± 0.001
Yi-1.5-9B-Chat	0.080 ± 0.002	0.840 ± 0.000	0.012 ± 0.001	-0.806 ± 0.005	0.554 ± 0.001	0.136 ± 0.002	0.035 ± 0.001	0.117 ± 0.002
gemma-2-27b-it	0.000 ± 0.000	0.802 ± 0.001	0.000 ± 0.000	-1.139 ± 0.014	0.519 ± 0.002	0.038 ± 0.004	0.001 ± 0.000	0.033 ± 0.004
gemma-2-9b-it	0.000 ± 0.000	0.806 ± 0.001	0.000 ± 0.000	-1.051 ± 0.007	0.509 ± 0.001	0.046 ± 0.002	0.001 ± 0.000	0.042 ± 0.002

Table 16: Summarization results.

Model	Close-ended										
	MedMCQA ↑	MedQA ↑	ANONIMIZED ↑	multimedqa ↑	PubMedQA ↑	MedText (easy) ↑	MedText (hard) ↑	BioRED ↑	Med prescriptions (easy) ↑	Med prescriptions (hard) ↑	MMLU ↑
BioMistral-MedMNX	0.495 ± 0.008	0.515 ± 0.014	0.629 ± 0.006	0.547 ± 0.006	0.776 ± 0.019	0.202 ± 0.011	0.050 ± 0.010	0.216 ± 0.013	0.989 ± 0.001	0.942 ± 0.002	0.6784 ± 0.034
JSL-MedLlama-3-8B-v2.0	0.613 ± 0.008	0.617 ± 0.014	0.671 ± 0.006	0.648 ± 0.006	0.742 ± 0.020	0.191 ± 0.010	0.082 ± 0.012	0.254 ± 0.014	0.993 ± 0.001	0.937 ± 0.002	0.7739 ± 0.0305
Llama3-Med42-8B	0.603 ± 0.008	0.626 ± 0.014	0.684 ± 0.006	0.642 ± 0.006	0.772 ± 0.019	0.202 ± 0.011	0.072 ± 0.012	0.203 ± 0.013	0.978 ± 0.001	0.933 ± 0.002	0.7525 ± 0.0315
Meta-Llama-3.1-70B-Instruct	0.722 ± 0.007	0.798 ± 0.011	0.836 ± 0.005	0.764 ± 0.005	0.800 ± 0.018	0.145 ± 0.003	-	0.515 ± 0.016	0.987 ± 0.001	0.934 ± 0.002	0.8711 ± 0.0236
Meta-Llama-3.1-8B-Instruct	0.593 ± 0.008	0.637 ± 0.013	0.703 ± 0.006	0.638 ± 0.006	0.752 ± 0.019	0.161 ± 0.003	-	0.232 ± 0.013	0.986 ± 0.001	0.929 ± 0.002	0.7621 ± 0.031
Mistral-7B-Instruct-v0.3	0.482 ± 0.008	0.523 ± 0.014	0.607 ± 0.007	0.538 ± 0.006	0.774 ± 0.019	0.178 ± 0.010	0.074 ± 0.012	0.358 ± 0.015	0.990 ± 0.001	0.941 ± 0.002	0.661 ± 0.0345
Mixtral-8x7B-Instruct-v0.1	0.564 ± 0.008	0.614 ± 0.014	0.725 ± 0.006	0.622 ± 0.006	0.796 ± 0.018	0.207 ± 0.011	0.062 ± 0.011	0.352 ± 0.015	0.982 ± 0.001	0.937 ± 0.002	0.7766 ± 0.0304
Phi-3-medium-4k-instruct	0.623 ± 0.007	0.596 ± 0.014	0.769 ± 0.006	0.661 ± 0.006	0.782 ± 0.018	0.048 ± 0.002	-	0.261 ± 0.014	0.990 ± 0.001	0.933 ± 0.002	0.8237 ± 0.0275
Phi-3-mini-4k-instruct	0.572 ± 0.008	0.537 ± 0.014	0.700 ± 0.006	0.604 ± 0.006	0.752 ± 0.019	0.192 ± 0.003	-	0.262 ± 0.014	0.983 ± 0.001	0.921 ± 0.003	0.7398 ± 0.0321
Qwen2-7B-Instruct	0.551 ± 0.008	0.570 ± 0.014	0.681 ± 0.006	0.596 ± 0.006	0.742 ± 0.020	0.225 ± 0.011	0.058 ± 0.010	0.197 ± 0.013	0.976 ± 0.001	0.934 ± 0.002	0.7337 ± 0.032
Yi-1.5-34B-Chat	0.575 ± 0.008	0.614 ± 0.014	0.732 ± 0.006	0.628 ± 0.006	0.774 ± 0.019	0.301 ± 0.012	0.345 ± 0.007	0.543 ± 0.016	0.972 ± 0.002	0.938 ± 0.002	0.7806 ± 0.0298
Yi-1.5-9B-Chat	0.488 ± 0.008	0.515 ± 0.014	0.649 ± 0.006	0.546 ± 0.006	0.774 ± 0.019	0.227 ± 0.011	0.046 ± 0.009	0.537 ± 0.016	0.985 ± 0.001	0.934 ± 0.002	0.7007 ± 0.0329
gemma-2-27b-it	0.616 ± 0.008	0.669 ± 0.013	0.781 ± 0.006	0.670 ± 0.006	0.800 ± 0.018	-	-	-	0.220 ± 0.004	0.331 ± 0.004	0.8163 ± 0.0278
gemma-2-9b-it	0.566 ± 0.008	0.629 ± 0.014	0.745 ± 0.006	0.625 ± 0.006	0.766 ± 0.019	0.209 ± 0.011	0.246 ± 0.019	0.536 ± 0.016	0.230 ± 0.004	0.399 ± 0.005	0.7796 ± 0.0303

Table 17: Close-ended results.