Are Clinical T5 Models Better for Clinical Text?

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Abstract

Large language models with a transformerbased encoder/decoder architecture, such as T5 (Raffel et al., 2023), have become standard platforms for supervised tasks. To bring these technologies to the clinical domain, recent work has trained new (Lehman et al., 2023) or adapted existing (Lu et al., 2022) models to clinical data. However, the evaluation of these clinical T5 models and comparison to other models has been limited. Are the clinical T5 models better choices than FLAN-tuned (Chung et al., 2022a) generic T5 models? Do they generalize better to new clinical domains that differ from the training sets? We comprehensively evaluate these models across several clinical tasks and domains. We find that clinical T5 models provide marginal improvements over existing models, and perform worse when evaluated on different domains. Our results inform future choices in developing clinical LLMs.

Keywords: Language Models, Domain Generalization, Pre-training

Data and Code Availability We do not release any new data or models as part of this study. The code in available on GitHub: https://github.com/yli-z/ml4h_are_clinical_t5_models_better_for_clinical_text.

Institutional Review Board (IRB) Our clinical datasets were obtained from the PhysioNet website under the PhysioNet Credentialed Health Data Use Agreement and from an anonymized Hospital System following IRB approval.

1. Introduction

There is an ongoing conversation in the community about the best strategy for developing Large Language Models (LLMs) for specialized domains. general-purpose LLMs, trained on massive amounts of diverse text, can generalize remarkably well to new domains and tasks (Brown et al., 2020; Touvron et al., 2023; Bubeck et al., 2023), especially with a small amount of domain-specific alignment data (Singhal et al., 2022). At the same time, specialized LLMs trained using domain-specific data can outperform their generic counterparts (Wu et al., 2023b; Taylor et al., 2022; Lehman et al., 2023; Lu et al., 2022). For supervised tasks, Text-to-Text Transfer Transformer (T5) models are a popular choice as they provide the benefits of both extensive pre-training and task generalization(Raffel et al., 2023); however, there are few studies on the benefits of domain-specialized T5 training.

Consider clinical text from Electronic Health Records (EHR). Access to this data is severely restricted and the data differs substantially from popular pre-training data sets. Recent work has explored developing T5 models for this domain. One approach adapts an existing T5 model first to the biomedical domain (journal articles) and then to the clinical domain using a small amount of clinical text (Lu et al., 2022). Another approach starts from scratch, developing a new vocabulary and model focused on clinical data (Lehman et al., 2023). As these works were completed concurrently, they do not include a direct comparison between the approaches, leaving the best strategy for T5 models an open question.

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Furthermore, while a specialized clinical T5 model sounds attractive, it has several potential shortcomings. First, the severe limitation of available EHR pre-training text means specialized T5 models are not only trained on less data overall, but also that the pre-training text is unlikely to adequately represent the true diversity of the domain. In contrast, pre-trained T5 models benefit from massive amounts of diverse data, which may yield better generalization abilities. Second, existing general T5 models may make use of extensive supervised datasets for FLAN tuning (Chung et al., 2022a), potentially leading to better generalization to new supervised tasks. Specialized clinical T5 models do not have the same opportunity nor benefit. Does FLAN training mean better generalization, even without access to EHR pre-training?

We thereby present a series of evaluations of existing general-purpose and clinical T5 models to answer two questions – 1) What existing T5 model should clinical NLP practitioners utilize when training a supervised system, and 2) What training strategy is more effective in the clinical domain with severe data limitations nowadays?

We make the following contributions.

- We evaluate 2 clinical, 2 non-clinical, and 1 FLAN-tuned T5 variants across 7 clinical and biomedical tasks. We independently reproduce prior results (Lu et al., 2022; Lehman et al., 2023) and extend the evaluation to new clinical datasets. We find that general T5 models achieve similar results as the clinical variants.
- We investigate model performance under datalimited conditions and find that FLAN-T5 excels on clinical tasks in the low-data settings.
- We evaluate clinical T5 models on new clinical domains and show they perform worse than general T5 models, suggesting that existing clinical models are overfitted to the limited clinical datasets available.

Taken together with other studies on clinical LLMs (Alsentzer et al., 2019; Lewis et al., 2020a; Singhal et al., 2022; Lu et al., 2022; Lehman et al., 2023), our findings suggest that future work should focus on adapting general T5 models to the clinical domain through new supervised training sets rather than training new methods from scratch.

2. Clinical T5 Models

LLMs have become the standard base models upon which NLP systems are built. Instruction following tasks typically leverage general-purpose decode LLMs (e.g., GPT, LlaMa) (Radford et al., 2019; Brown et al., 2020; Touvron et al., 2023; Bubeck et al., 2023), whereas supervised tasks typically rely on encoder/decoder (Raffel et al., 2023) or encoder models (Devlin et al., 2019).

Historically, clinical NLP systems have benefited from specialized versions of these models due to differences in the topic, style, and vocabulary of clinical text relative to other language domains. However, these advantages have been tempered under various circumstances (Gutiérrez et al., 2023; Lewis et al., 2020a), and are not necessarily guaranteed to hold given the existence of much larger contemporary LLMs. This uncertainty raises several key questions for clinical NLP practitioners.

What existing models should be used to develop a supervised clinical NLP system?

T5 (Raffel et al., 2023) and Flan-T5 (Chung et al., 2022a) do exceedingly well on a wide variety of supervised tasks, benefited from large amounts of pretraining and diverse task-specific fine-tuning, respectively. These models balance out-of-the-box zero-shot performance and supervised fine-tuning when a moderate amount of supervised data is available. Within the clinical domain, two T5-based models (Lehman et al., 2023; Lu et al., 2022) have been trained on clinical text from MIMIC (Johnson et al., 2016, 2023). Which model should we prefer and in what setting? Unfortunately, the publication of these two models was concurrent, so a head-to-head comparison is unavailable. Therefore, we conduct a direct comparison with the same training and evaluation settings to determine which model should be preferred.

Should we train from scratch on clinical data or adapt an existing pre-trained model?

Beyond determining which model is preferred, these two models represent different strategies for developing domain-specific models:

Lu et al. (2022) utilize continued pre-training to tune an existing model for the clinical setting, which allows them to benefit from large amounts of previous training on general-purpose and related data. They use MIMIC-III (Johnson et al., 2016) to train SciFive (Phan et al., 2021), which is itself adapted to the scientific domain from the base T5 model (Raffel et al., 2023). In contrast, Lehman et al. (2023) train a new T5 model from scratch on clinical text from MIMIC-III and MIMIC-IV (Johnson et al., 2016, 2023), which enables them to leverage a domain-specific vocabulary.

These models follow larger trends in developing domain-specific models, including training from scratch (Wu et al., 2023b; Taylor et al., 2022; Lewis et al., 2020a) and continued pre-training or aligning existing models (Alsentzer et al., 2019; Singhal et al., 2022). In the clinical domain specifically, encoder (masked) LLMs have been adapted from pre-trained models (e.g., BioBERT (Lee et al., 2019) and Clinical-BioBERT (Alsentzer et al., 2019) from BERT (Devlin et al., 2019)) and have also been trained from scratch (e.g., GatorTron was pre-trained on a combination of Wikipedia articles, PubMed publications, and deidentified clinical notes (Yang et al., 2022)). Decoder (GPT-style) LLMs have been adapted to the medical domain, but not necessarily the clinical domain (e.g., Med-PaLm (Singhal et al., 2022) from FLAN-PaLm (Chowdhery et al., 2022)).

Arguments have been made in support of both pretraining strategies. We are interested in understanding whether one strategy is preferable over another when a) there is a need to generalize to a new clinical domain, and b) when only a limited amount of supervised data in the target domain is available.

Should we prefer domain-specialized models over powerful, general-purpose models?

Domain-specificity might not be optimal: biomedical language models have been shown to be robust to domain-general tokenization (Gutiérrez et al., 2023), and pre-trained LLMs have been shown to possess significant knowledge about the medical domain despite not having been trained specifically on biomedical data (Agrawal et al., 2022; Singhal et al., 2023). In the clinical domain, pre-trained clinical BERT models do not consistently demonstrate a significant improvement over non-clinical variants of BERT (Lewis et al., 2020b; Harrigian et al., 2023a; Yue et al., 2020). This suggests that the benefits of in-domain pre-training may not always justify the additional complexity and computational cost.

Morover, in the case of T5 models, there exist highquality FLAN-tuned versions that do well on a variety of supervised tasks. While they are not clinically tuned, they do incorporate a lot of supervision. Should we prefer these general-purpose models to clinical models, especially in new clinical settings or those with limited supervision? We compare these models to clinical T5 models to determine which should be preferred in clinical settings.

Overall, our analysis provides practical guidance to practitioners, provides new evidence on LLM domain specialization methods, and explores the tradeoff between domain specialized and general-purpose LLMs for supervised tasks.

3. Experiment Setup

3.1. Models

Text-To-Text Transfer Transformer (T5) (Raffel et al., 2023) and its variants are sequence-to-sequence models utilizing an encoder/decoder Transformer architecture (Vaswani et al., 2017). We evaluate several existing pre-trained models, all being the large variant (\sim 770M parameters), by fine-tuning them on (x,y) pairs, where x and y are the input and output of a task, formatted as text.

Additional information about each model is included in Table 4.

T5 The original T5 model was trained in two stages. First, it was trained with a self-supervised text denoising objective on web crawl data (Raffel et al., 2023), and then with a conditional generation objective on labeled data from a diverse mixture of downstream tasks. To separate the effects of pretraining using non-clinical web data and performing supervised multitask pre-training, we evaluate two versions of T5: T5-Den which is trained only using the denoising objective and T5-Sup which is further pre-trained on supervised multi-task tasks.¹

FLAN-T5 FLAN-T5 (Chung et al., 2022a) took the LM-Adapted T5, which was trained for additional 100K steps on the LM objective from T5-Den (Raffel et al., 2023; Lester et al., 2021), and further trained with extensive supervised instruction finetuning. The instruction tuning process typically improves the model's generalization ability to new tasks (Longpre et al., 2023). We include FLAN-T5 to evaluate its generalization capabilities on clinical tasks.

Most studies only evaluate T5-Sup and call it a pre-trained only model.

Clinical-T5 We evaluate two concurrently developed Clinical-T5 models – similarly named, but pretrained on different corpora and with different weight initialization strategies. We refer to the model introduced by Lehman et al. (2023) as MIMIC-T5 since their model is only pre-trained on the union of all notes from MIMIC-III and MIMIC-IV (Johnson et al., 2016, 2023), including discharge summaries and radiology reports, from random initialization. We refer to the model introduced by Lu et al. (2022) as SciFive+MIMIC-T5 since their model is trained on MIMIC-III (Johnson et al., 2016) but initializes from a biomedically pre-trained model (SciFive), which gives it additional exposure to biomedical data. Sci-Five was itself trained from the base T5 initialization.

3.2. Datasets

We evaluate models on the union of datasets considered by each of the clinical T5 studies (Lehman et al., 2023; Lu et al., 2022), as well as clinical data not from MIMIC-III.

MIMIC-III Datasets We evaluate on CLIP (Mullenbach et al., 2021), a multi-label classification dataset, RadQA (Soni et al., 2022), a question-answering dataset, and MedNLI (Romanov and Shivade, 2018), a natural language inference dataset. Each dataset is comprised of instances from MIMIC-III (Johnson et al., 2016). Note that these datasets annotate data that comes from the same corpus used for clinical model pre-training.

Biomedical Datasets We evaluate on HOC (Baker et al., 2015), a multi-label classification dataset, and two named entity recognition datasets, BC5CDR-disease (Li et al., 2016a) and NCBI-disease (Doğan et al., 2014). HOC and NCBI-disease are datasets derived from PubMed abstracts ², and the BC5CDR-disease dataset is derived from full PubMed articles. Note that some of them annotate data that comes from the same corpus used for SciFive+MIMIC-T5 pre-training.

Clinical Stigmatizing Language Datasets Due to the extremely limited availability of clinical text data, public clinical NLP datasets are drawn from MIMIC (MIMIC-III (Johnson et al., 2016) in particular), the same dataset used to pre-train the clinical-T5 models. How well do these clinical models generalize to a novel clinical domain and task? We ob-

tain data from Hospital System (anonymized) that includes five medical specialties: Internal Medicine, Emergency Medicine, Pediatrics, OB-GYN, and Surgery. We evaluate the task of detecting stigmatizing language about patients by using the setup as Harrigian et al. (2023b). We also use their annotations from MIMIC-IV.

4. Are Clinical T5 Models Better for Clinical Text?

Both MIMIC-T5 and SciFive+MIMIC-T5 have demonstrated improved performance on clinical EHR text compared to a general-purpose model T5. Our first step is to replicate these findings with a more extensive and direct evaluation.

Methods We began by reproducing the results reported in Lehman et al. (2023) and Lu et al. (2022), adhering closely to each study's original evaluation methodology. Both studies only leveraged a single train/dev/test split for each respective task, and only Lehman et al. (2023) estimated uncertainty around their performance point estimates.³ To understand whether prior results were sensitive to the initial dataset split, we conducted an additional set of crossvalidation experiments. We merged existing training and development splits and randomly split them into 5 subsets to facilitate 5-fold cross-validation. Additionally, the original held-out test split was used to evaluate generalization and compare to prior work. We considered the same evaluation metrics used in prior work (Table 8), using a bootstrap procedure to estimate confidence intervals. For uniformity across the metrics (e.g., exact match, F1 score), differences in performance between models were assessed using a paired t-test (paired by fold). We referred to MIMIC-T5 and SciFive-T5's evaluation metrics on these datasets. Training and evaluation details are included in Appendix D.

Results Cross-validation results are reported in Table 1, while reproducibility efforts are reported alongside the originally reported performance metrics in Appendix F. Our results for MIMIC-T5 perform slightly worse than previously reported on MedNLI and RadQA, but better on CLIP. SciFive+MIMIC-T5 performs slightly better than previously reported on the three biomedical tasks. The average perfor-

^{2.} https://pubmed.ncbi.nlm.nih.gov/

^{3.} Lehman et al. (2023) used at least three random seeds in their training procedure to estimate uncertainty.

Dataset	Metrics	T5-Den	T5-Sup	FLAN-T5	MIMIC-T5	SciFive+MIMIC-T5
MedNLI	Acc.	85.9(85.3,86.2)	84.9(84.4,85.5)	86.0 _(85.7,86.4)	86.8 _(86.3,87.3)	85.6 _(85.1,86.4)
RadQA	EM F1	$52.6_{(51.9,53.3)} \\ 68.9_{(68.0,69.8)}$	$52.0_{(51.0,53.0)} \\ 68.7_{(67.8,69.8)}$	$53.0_{(51.8,54.0)} 70.6_{(69.8,71.3)}$	$54.5_{(54.0,55.1)} 73.4_{(72.8,74.0)}$	$53.4_{(52.7,54.2)} 70.4_{(69.4,71.5)}$
CLIP	Macro F1 Micro F1	$63.9_{(62.2,65.6)} 78.6_{(77.5,79.8)}$	$62.4_{(61.4,63.4)} 78.4_{(78.0,78.8)}$	$64.3_{(62.9,65.7)} 79.4_{(78.9,79.9)}$	$66.1_{(65.9,66.3)} 79.3_{(78.7,79.9)}$	$63.5_{(62.2,64.8)} \\78.2_{(77.9,78.5)}$
нос	F1 P R	84.9 _(84.4,85.6) 84.7 _(84.2,85.6) 85.1 _(84.7,85.6)	84.8 _(84.4,85.1) 84.8 _(84.4,85.1) 84.8 _(84.3,85.2)	84.8 _(84.2,85.3) 84.6 _(83.9,85.2) 85.0 _(84.5,85.4)	82.8 _(82.6,83.0) 82.9 _(82.8,83.1) 82.6 _(82.3,83.0)	$85.1_{(84.8,85.4)} \\ 85.0_{(84.7,85.5)} \\ 85.2_{(84.9,85.5)}$
BC5CDR	F1 P R	82.9 _(82.3,83.4) 81.7 _(80.8,82.5) 84.2 _(83.6,84.6)	82.9 _(82.5,83.5) 81.6 _(81.3,82.0) 84.3 _(83.6,85.0)	83.3 _(82.9,83.8) 82.1 _(81.5,82.7) 84.5 _(84.2,84.9)	$81.2_{(80.9,81.5)} \\ 80.6_{(80.2,80.9)} \\ 81.7_{(81.2,82.3)}$	$83.6_{(83.4,83.8)} \\ 83.1_{(82.5,83.6)} \\ 84.2_{(83.6,84.8)}$
NCBI	F1 P R	85.1 _(84.6,85.5) 83.5 _(82.8,84.2) 86.7 _(86.3,87.2)	84.6 _(84.2,85.0) 83.6 _(83.1,84.1) 85.6 _(85.2,86.2)	84.4 _(83.9,85.0) 82.2 _(81.6,83.0) 86.8 _(86.3,87.3)	79.1 _(78.3,79.9) 78.4 _(77.6,78.9) 79.9 _(78.7,81.3)	$85.4_{(85.0,86.2)} \\ 83.8_{(83.0,84.7)} \\ 87.1_{(86.9,87.5)}$

Table 1: Mean test performance and 95% confidence intervals across 5-fold cross-validation. MIMIC-T5 outperforms alternative models on clinical tasks, but struggles on biomedical tasks. Inclusion of biomedical data during pre-training is useful for the biomedical tasks.

mance over the cross-validation procedure mirrored the single train/dev/test split results.

Despite minor differences from the originallyreported results, we confirm the findings of Lehman et al. (2023) that MIMIC-T5 outperforms T5-Sup across the three clinical tasks at a statistically significant level. In comparison, SciFive+MIMIC-T5 either slightly underperformed or slightly outperformed the non-clinical T5 variants across the clinical tasks, albeit not at a statistically significant level. While the former outcome seems to corroborate the efficacy of domain-specific clinical pre-training, the latter outcome suggests the effect may be moderated inconsistently by pre-training on out-of-domain text. Given that MIMIC-T5 was pre-trained on text from the same corpus used for curating the evaluation datasets, the observed improvement may be a consequence of overfitting, which we explore below. Moreover, we note that the absolute increase in performance (~ 1.0 to ~ 1.5 points across clinical tasks) falls within a range that is reasonable to expect when using domain adaptive pre-training (i.e., continued pretraining) (Gururangan et al., 2020).

Results on biomedical (journal article) datasets highlight the differences between the adaptation and train from scratch strategies. SciFive+MIMIC-T5 does better than MIMIC-T5 on these tasks, some-

times by more than 4 points, presumably since it was adapted from SciFive which is trained on similar data, whereas MIMIC-T5 has only seen clinical data. However, SciFive+MIMIC-T5 only slightly outperforms the general-purpose models, and does not do so consistently at a statistically significant level. It is possible that the general-purpose models have already seen data similar enough to the biomedical data such that further pre-training has minimal effect.

Finally, we observe the benefits of supervised instruction fine-tuning. FLAN-T5 performs comparably to MIMIC-T5 on the three clinical datasets and to SCIFIVE+MIMIC-T5 on the three biomedical datasets. MIMIC-T5 only outperforms FLAN-T5 on RadQA dataset, and only under 1 metric (F1 score), with statistical significance. Recent work suggests that training on relevant target tasks, either through instruction tuning or multitask training, has broad downstream benefits (Chung et al., 2022b; Mueller et al., 2022). T5-Den and T5-Sup perform similarly on biomedical tasks, but not the clinical tasks, where T5-Den outperforms T5-Sup slightly, albeit not at a statistically significant level. Whether the remaining gap in performance between non-clinical and clinical T5 models can be decreased via supervised pretraining on related tasks or extensive pre-training in general is an important open area of study, especially given that non-clinical task datasets are significantly more abundant than clinical ones.

5. Do Clinical T5 Models Generalize to New Clinical Text?

A key limitation in developing clinical NLP systems is the lack of publicly available data due to PHI concerns. Mostly public available clinical models, including MIMIC-T5 and SCIFIVE+MIMIC-T5, and evaluation sets utilize MIMIC, so we do not know if they generalize to new clinical text sources. The weaker performance of MIMIC-T5 on the biomedical datasets may suggest its limited ability to generalize, or perhaps its benefits remain when focused on clinical EHR text. Since the clinical pre-training and evaluation sets represent only a single medical domain from a single institution, we ask: do clinical T5 models generalize to new clinical text sources and tasks better than general-purpose models?

Methods To evaluate generalization, we consider a new task for which we can evaluate on both a MIMIC dataset and a Hospital System dataset. The task is to characterize stigmatizing language in medical records, which includes three multi-class classification tasks that operate similarly to word sense disambiguation. The two datasets were curated using the annotation guidelines set forth by Harrigian et al. (2023b). We perform 5-fold cross-validation experiments as above on these two datasets, and compare the in-domain (MIMIC-IV) and out-of-domain (Hospital System (anonymized)) performance.

Results Table 2 shows the performance of the T5 models on the two stigmatizing language datasets. On the MIMIC-IV dataset, T5-Sup, MIMIC-T5, and SciFive+MIMIC-T5 achieve similar levels of performance. For this task, the benefits of MIMIC-T5 do not materialize, despite the MIMIC-IV dataset being within the distribution of its pre-training data.⁴

When we consider the out-of-distribution Hospital System (anonymized) data, MIMIC-T5 falls behind these other two models, perhaps either because MIMIC-T5 has been overfit to MIMIC data, or because it lacks the large-scale pre-training of the other models. SciFive+MIMIC-T5 performs slightly better than MIMIC-T5 on these two clinical datasets.

However, neither MIMIC-T5 or SciFive+MIMIC-T5 show improvement over T5-Sup or FLAN-T5.

Finally, these evaluations further highlight the generalization ability of T5-Sup and FLAN-T5, which do better than clinical T5 models on both datasets. Even without clinical tasks in hand, the additional supervised training yields a model that does better on a novel clinical task across both data sources.

6. Do Clinical T5 Models Perform Well in Low-Resource Settings?

One of the possible reasons we have observed only a small performance gap between general-purpose and clinical models may be the presence of sufficient task training data. Indeed, previous work has found that tuning general-purpose BERT models for a clinical task using supervised data can nullify a clinical model's advantage (Harrigian et al., 2023a). In this regard, we may expect to see larger benefits to clinical models in a low-resource setting. Importantly, such settings are common in clinical tasks, where annotation typically requires advanced expertise, or is nigh impossible due to data privacy constraints. We therefore explore if the advantages of clinical models emerge more prominently in low-resource settings.

Methods We selected MedNLI and the two stigmatizing language datasets to test a low-resource setting. For MedNLI, we used the same cross validation splits as before, but downsampled the training subset to 1% (\sim 99 examples on average per cross validation fold). For the stigmatizing language datasets, which has less data than MedNLI to begin with, we were able to explore multiple downsampled training size settings – 1% (~ 2 examples on average per fold), 5%(\sim 12 examples on average per fold), and 25% (\sim 59 examples on average per fold) for every task. We note that 1% of training data represents an extreme scenario, albeit one that is not inconceivable given the complexities of annotating some clinical datasets (Spasic and Nenadic, 2020). Experimental details are described in the Appendix D.

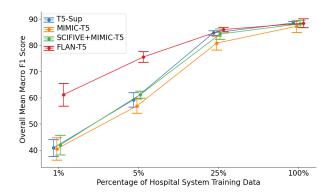
Results Figure 1 and Figure 2 show results for the Hospital System (anonymized) and MIMIC-IV stigmatizing language datasets, respectively, while Table 3 shows results for MedNLI. Additional task-specific results are reported in the Appendix G.

In the 1% MedNLI experiment, MIMIC-T5 is no longer leading, nor is it even competitive with, the other T5 models. For the two stigmatizing language

^{4.} The MIMIC-IV stigmatizing language dataset may have been seen verbatim during MIMIC-T5 pre-training, as they did not leverage the same evaluation splits.

		Hospital Syster	MIMIC-IV					
Model	Credibility & Obstinacy	Compliance	Descriptors	Average	Credibility & Obstinacy	Compliance	Descriptors	Average
T5-Sup	88.2(86.1,90.6)	86.7(85.5,88.2)	90.6(89.3,91.4)	88.5 _(87.8,89.0)	76.0(74.1,77.9)	92.9(91.7,94.2)	85.7(83.4,88.0)	84.9(84.2,85.8)
MIMIC-T5	88.1(83.1,94.1)		88.9(88.2,89.5)	87.2(85.1,89.7)	$75.0_{(72.1,77.8)}$	$91.6_{(90.4,92.7)}$		83.8(82.5,84.5)
SciFive+MIMIC-T5	$87.3_{(82.1,91.8)}$	87.2(86.8,87.6)	$90.0_{(89.2,90.7)}$	88.2(86.6,89.2)	$74.6_{(70.6,78.8)}$	$91.0_{(89.9,92.1)}$	$86.3_{(85.0,87.8)}$	$84.0_{(82.0,85.8)}$
FLAN-T5	90.4(88.3,92.3)	88.0(86.8,88.9)				92.9(91.3,94.5)	86.3 _(85.3,87.0)	85.6 _(85.0,86.1)

Table 2: Macro F1 scores on stigmatizing language datasets from the Hospital System (anonymized) and MIMIC-IV.



T5-Sup MIMIC-T5
SCIFIVE+MIMIC-T5
FLAN-T5

1/6

1/6

5/6

Percentage of MIMIC-IV Training Data

Figure 1: Mean macro-F1 scores across the three stigmatizing language classification tasks for the Hospital System (anonymized) dataset using a random 1%, 5%, and 25% sample of the training data.

Figure 2: Mean macro-F1 scores across the three stigmatizing language classification tasks for the MIMIC-IV dataset using a random 1%, 5%, and 25% sample of the training data.

Model	Accuracy
T5-Sup	$78.80_{(77.69,79.90)}$
MIMIC-T5	$63.87_{(60.35,66.95)}$
SciFive+MIMIC-T5	$78.92_{(78.35,79.84)}$
FLAN-T5	$79.49_{(78.97,80.06)}$

Table 3: Performance with 1% training data on MedNLI dataset

datasets, T5-Sup, MIMIC-T5, and SciFive+MIMIC-T5 all perform roughly equivalently to one another in the low-resource settings as we gradually increase the sample size. That said, MIMIC-T5 finds itself at the bottom of relative performance rankings in more cases than not. Across all three tasks, FLAN-T5 excels in the low-resource settings compared to the three non-FLAN-tuned T5 models. Altogether, our results suggest that strong general-purpose lan-

guage models are more appropriate for new distributions and those with limited amounts of task-specific training data.

7. Discussion

In this study, we independently replicated and extended prior work concerning specialized clinical T5 models (Lu et al., 2022; Lehman et al., 2023). By placing the models in an expanded context (i.e., additional non-clinical variants, evaluation datasets), we find ourselves in a position to make the following recommendations for clinical NLP practitioners.

Clinical models pre-trained from scratch using comparatively small datasets should not be used beyond their pre-training distributions.

Language models that are pre-trained on the target domain from scratch can achieve strong target-domain performance due to their in-domain pre-training (Gupta et al., 2023). For example, PubMed-BERT (Gu et al., 2021), which was pre-trained from scratch on 3.2 billion words from PubMed Abstracts only, achieved remarkable performance in biomedical tasks drawn also from PubMed Abstracts. Unfortunately, large amounts of diverse and relevant pre-training data are generally hard to obtain for most real-world clinical applications.

Currently, MIMIC provides the largest public clinical text corpora, coming in at approximately 1.2 billion words across both MIMIC-III and MIMIC-IV (Lehman et al., 2023). However, practitioners must remember that MIMIC only represents a small subset of the broader clinical text landscape. Not only are there sample biases related to the demographics of patients in the corpora, but also topic biases (e.g., intensive care unit patients) and syntactic biases (e.g., hospital-specific documentation practices). LLMs trained on this comparatively narrow sample of data provide optimistic estimates of performance when evaluated on data also drawn from the MIMIC corpora (Lehman et al., 2023). As shown in our experiments, they struggle to adapt to the diverse needs of the clinical and biomedical space. So-called "clinical" language models may perhaps be better described as "MIMIC" language models.

Clinical models should only be used in downstream tasks if they have access to sufficient supervised training data.

In low-data settings, specificalized clinical models are more likely to underperform their generic counterparts that have been pre-trained on more data and/or using supervised fine-tuning. For example, in §6, we saw that T5-Sup, SciFive+MIMIC-T5, and FLAN-T5 were all more qualified to operate in at the 1% downsample regime than MIMIC-T5. Even larger generic LLMs such as ChatGPT and PaLM are likely to provide even more significant advantages in low-resource settings (Agrawal et al., 2022; Singhal et al., 2023). This leads us to our next recommendation.

The combination of task fine-tuning and FLAN instruction fine-tuning is hard to beat.

Recent research has shown that models pre-trained in a supervised manner are explicitly guided towards flat loss regions. Consequently, they are more robust to sequential fine-tuning in new domains (Mehta et al., 2023). It has also been shown that heterogeneity within pre-training data is important for promoting generalization in downstream tasks across diverse domains (Longpre et al., 2023). FLAN-T5's exceptional performance across all tasks in our study, especially in low-resource settings, affirms these prior findings.

Adapting general-purpose models rather than training from scratch, or leveraging a mix of both training strategies, may be the best option for clinical language models moving forward.

General-domain pre-training is beneficial given that clinical text has limited availability, and domainadaptive or task-adaptive training can further improve performance on downstream tasks (Gururangan et al., 2020). For example, continued pre-training on small amounts of annotated clinical data (i.e., task-adaptive pre-training) has improved transfer to out-of-domain clinical data in a note-section classification task (Zhou et al., 2023), while adapting a generic language model, such as BERT, has been shown to be more efficient than pre-training a new model from scratch in unseen target domains (Lamproudis et al., 2022). Various contemporary works have presented ways to further optimize continued pre-training for adaptation. For example, by reverse engineering the effects of instruction-tuning (Fleshman and Durme, 2024), or by finding more efficient warm-up strategies (Gupta et al., 2023).

8. Conclusion

Our study aimed to determine whether existing clinical T5 models offer performance improvements over non-clinical T5 variants in clinical tasks. Our findings suggest that these clinical language models may outperform their general counterparts, but only under specific conditions: a) the task is on clinical EHR data sourced from MIMIC; b) sufficient annotated training data is available. Meanwhile, general-purpose language models, especially those leveraging supervised multi-task instructing tuning (e.g., FLAN-T5), excel when these criteria are not met.

More broadly, our results do provide support to previous claims that truly "domain-specific" language models, trained for specific data distributions and tasks, are ideal (Lehman et al., 2023). However, they also suggest that domain-specific models are not always practical. Data distributions may change over time (e.g., suddenly due to a pandemic, gradually as clinical practice evolves) (Khanday et al., 2020; Jeong et al., 2024) or across target populations (Harrigian et al., 2023a). Likewise, not all domains are data rich and diverse enough to support training a language model from scratch that endows broader linguistic abilities. In the clinical domain, non-clinical language models will almost certainly continue to have access to pre-training corpora that are orders of magnitude larger than clinical pre-training corpora, and that data scale is greatly advantageous for training high-performing LLMs (Kaplan et al., 2020).

Lastly, we would be remiss not to comment on the cost and sustainability implications of training domain-specific language models. While it is true that given the same inference budget, a clinical model trained from scratch is typically able to achieve better performance than a generic model (Lehman et al., 2023), such a perspective regarding model efficiency is inherently narrow. Using floating point operations (FLOPs) at inference time as a proxy for efficiency does not accurately reflect the additional cost complexities of training a language model from scratch - e.g., training compute, human labor (development and annotation), data access, and dealing with future distribution shift. Put another way, specialized models may achieve improvements in performance that are statistically significant, but not necessarily practically significant.

Limitations

T5 Model Family Our study focused on clinically pretrained T5 models, made possible by the released models from Lu et al. (2022) and Lehman et al. (2023). Comparing them allowed us to study the effect of pretraining strategies while controlling for the architecture. Future work could extend these investigations to other pretrained model architectures.

Data Availability Due to the scarcity of clinical text datasets, our evaluation was limited to a total of seven datasets. This constraint might not provide a comprehensive assessment for a wide range of clinical

tasks. We have cited relevant work where applicable to supplement our findings.

Hyperparameter Search Due to computational costs associated with training each T5 model over multiple cross validation folds, we were only able to perform limited amounts of hyperparameter tuning at the onset of the study. Upon identifying a reasonable set of hyperparameters that worked well across the different datasets and models, we opted to fix them across all the cross validation runs. It is certainly possible that different hyperparameters may have yielded different outcomes. That said, the computational limitations faced here reflect what many practitioners face.

Statistical Measures We are aware that traditional statistical methods in NLP may not be sufficient to describe the divergence of performances in the clinical domain. While these statistical methods are effective in establishing the statistical significance of performance differences between models, they may fall short of providing a comprehensive understanding of how these differences manifest in real-world clinical settings. This limitation underscores the need for specialized evaluation frameworks in healthcare applications, thereby ensuring that the models not only perform well statistically but also deliver practical, reliable, and clinically relevant outcomes.

Future Directions The clinical domain presents unique challenges and opportunities. Future work could consider extending model architectures and training strategies (Gema et al., 2024; Labrak et al., 2024; Toma et al., 2023; Wu et al., 2023a), evaluation (Tang et al., 2023; Hager et al., 2024; Ness et al., 2024)), and techniques for addressing data scarcity and privacy issues (Lin et al., 2022).

Ethics Statement

Our clinical datasets were obtained from the PhysioNet website, accessed under the PhysioNet Credentialed Health Data Use Agreement, and from Hospital System (anonymized) after IRB approval. All data from MIMIC-III or MIMIC-IV has been de-identified to ensure privacy and confidentiality. Clinical-T5 models were pre-trained on MIMIC; we access and use them under the PhysioNet Credentialed Health Data license and data use agreement. All data processing and model training was done on remote servers secured using OS-level security protocols.

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Appendix A. Dataset Details

We adapted the data processing methods mostly from the original MIMIC-T5, SciFive+MIMIC-T5 and Sci-Five's paper and codebases.

- MedNLI MedNLI (Romanov and Shivade, 2018) is a natural language inference dataset derived from MIMIC-III (Johnson et al., 2016). Given two sentences premise and hypothesis it labels the relation between these two sentences as "entailment", "natural", and "contradiction."
- RadQA RadQA (Soni et al., 2022) is a questionanswering dataset sourced from the radiology report in MIMIC-III (Johnson et al., 2016). It annotates the answer span to the questions; if no answers for the question, the answer text is empty.
- CLIP CLIP (Mullenbach et al., 2021) is a multilabel classification dataset of clinical action items, such as "patient instructions" and "appointment." Due to the long context of clinical records of these two tasks, we referred to the methodology utilized by MIMIC-T5, which involves segmenting long records

and appropriately mapping labels to the segmented sequences.

BC5CDR-disease BC5CDR-disease (Li et al., 2016b) is a Disease Named Entity Recognition corpus, which annotates 1500 PubMed articles with disease entities labels. Suppose (x, y) is the inputoutput pair: y is the same as x except that disease entities are enclosed with "disease" and "*disease".

NCBI-disease NCBI-disease (Doğan et al., 2014) is also a corpus for Disease Named Entity Recognition from PubMed abtracts. Both BC5CDR-disease and NCBI-disease are preprocessed by SciFive and available in their repo.

HOC HOC dataset (Baker et al., 2015) is accessible on Huggingface. It is developed from PubMed publication abstracts. We use the sentence-level dataset, where each sentence is annotated for different hallmarks of cancer.

Clinical Stigmatizing Language Datasets Clinical Stigmatizing Language Datasets (Harrigian et al., 2023b) characterizes stigmatizing languages in three classification tasks:

- 1. Credibility & Obstinacy (Disbelief, Difficult, Exclude): expressions of doubt or resistance.
- 2. Compliance (Negative, Neutral, Positive): adherence to medical advice.
- 3. Descriptors (Negative, Neutral, Positive, Exclude): characterization of patient behavior and demeanor.

Appendix B. Prefix and Instructions

T5 Prefix We used the same prefix as MIMIC-T5 on MedNLI, RadQA, and CLIP datasets. We referred to SCFIVE-T5's github repository for NCBI, BC5CDR, NCBI's prefix. On the stigmatizing language datasets, we referred to its keyword category as prefix before inputs: "adamant" for the Credibility & Obstinacy task; "compliance" for the Compliance task; "other" for the Descriptors task.

FLAN-T5 Instructions We didn't exhaustively experiment with different instructions for FLAN-T5. We referred to some instructions from MIMIC-T5 and most instructions are straight-forward.

 MedNLI Answer entailment, contradiction or neutral. Premise: {premise}
 Hypothesis: {hypothesis}

- RadQA Context: {context} Question: {question} If no answer is found in the context, do not reply; otherwise, give an answer from the context:
- CLIP Context: {context}. Label the above sentence as an empty string or as one or more of the following options, delimited by comma: Options: {labels}
- HOC Sentence: {input} Assign the above sentence as zero or more of the following class labels: {labels}
- BC5CR-disease & NCBI-disease Sentence: {input} Identify and label disease terms in the sentence:
- Stigmatizing Language Dataset
 - Credibility & Obstinacy Classify this sentence as difficult, disbelief, or exclude, regarding the credibility and obstinacy of the patient: {input
 - Compliance Classify this sentence as negative, neutral, or positive, regarding the patient's compliance with medical advice: {input}
 - Descriptors Classify this sentence as exclude, negative, neutral, or positive, regarding the patient's behavior and demeanor: {input}

Appendix C. Model Details

The details of the pre-training corpus for each model are listed in Table 4.

Appendix D. Training Details

D.1. Training settings

Training on Full Datasets We use code provided by prior work (Phan et al., 2021; Lehman et al., 2023) to fine-tune the various T5 variants with pytorch==2.1.2. Throughout our experiments, we used the adafactor optimizer (Shazeer and Stern, 2018) with constant lr = 1e - 4, batch size of 64, and set max sequence lengths that depend on the dataset.

	Weight Initialization	pre-training Data
T5-Sup (Raffel et al., 2023)	Random	Colossal Clean Crawled Corpus (C4) + Supervised Tasks (Raffel et al., 2023)
T5-Den (Raffel et al., 2023)	Random	Colossal Clean Crawled Corpus (C4) (Raffel et al., 2023)
Clinical-T5 (Lehman et al., 2023)	Random	MIMIC-III (Johnson et al., 2016) & MIMIC-IV (Johnson et al., 2023)
Clinical-T5 (Lu et al., 2022)	SciFive-PubMed-PMC (Phan et al., 2021)	MIMIC-III (Johnson et al., 2016)

Table 4: pre-training corpus for the models we compared. All models compared have 770 million parameters.

For all fine-tuning runs, we train for a fixed amount of epochs (30) and pick the best checkpoint based on validation set performance. We chose accuracy for MedNLI, and F-1 for other datasets to pick the best checkpoint during the validation and then evaluated on the complete test split per model and dataset.

Training on Downsampled Datasets Due to variations in the downsampled training data, we conducted a learning rate search for all models in the range of [1e - 3, 5e - 4, 1e - 4, 1e - 5] using a batch size of 16 with 5% and 25% downsampling training data and using a batch size of 4 with 1% downsampling training data on stigmatizing language datasets and ran the learning rate search in the range of [5e - 3, 1e - 3, 5e - 4, 1e - 4] with a batch size of 64 on MedNLI dataset. We trained each model for 30 epochs. We selected the best checkpoint for each model based on their average validation score across folds (i.e. macro-F1 for the stigmatizing task and accuracy for the MedNLI task) and subsequently evaluated on the test dataset. The best-found learning rates are reported in Table 5, Table 6, and Table 7.

D.2. Differences from Prior Work

We summarize a list of changes we made that possibly contribute to the difference in results compared to prior Clinical-T5s works (Lu et al., 2022; Lehman et al., 2023):

- We trained each model on each supervised dataset from scratch.
- We implemented the 5-fold cross-validation experiment, focusing on variance in data. This is a different evaluation structure from prior works.

1% Training data		
Model	Learning Rate	Batch Size
$T\bar{5}$ - \bar{Sup}	1e-4	4
FLAN-T5	1e-4	4
MIMIC-T5	5e-4	4
SCIFIVE+MIMIC-T5	1e-4	4
5% Training data		
Model	Learning Rate	Batch Size
T5-Sup	1e-4	<u>16</u>
FLAN-T5	1e-4	16
MIMIC-T5	5e-4	16
SCIFIVE+MIMIC-T5	5e-4	16
25% Training data		
Model	Learning Rate	Batch Size
T5-Sup	1e-4	<u>16</u>
FLAN-T5	1e-4	16
MIMIC-T5	1e-4	16
SCIFIVE+MIMIC-T5	1e-4	16

Table 5: Best-found hyperparameter settings for down-sampling experiments on stigmatizing language dataset - MIMIC-IV

1% Training data		
Model	Learning Rate	Batch Size
T5-Sup	1e-3	4
FLAN-T5	1e-4	4
MIMIC-T5	1e-3	4
SCIFIVE+MIMIC-T5	5e-4	4
5% Training data		
Model	Learning Rate	Batch Size
T5-Sup	1e-4	_ <u>_ 16 </u>
FLAN-T5	1e-4	16
MIMIC-T5	1e-3	16
SCIFIVE+MIMIC-T5	5e-4	16
25 % Training data		
Model	Learning Rate	Batch Size
T5-Sup	1e-4	16
TT 137 TT		

Table 6: Best-found hyperparameter settings for down-sampling experiments on stigmatizing language dataset - Hospital System

1e-4

5e-4

1e-4

16

16

16

FLAN-T5

MIMIC-T5

SCIFIVE+MIMIC-T5

1 % Training Data		
Model	Learning Rate	Batch Size
T5-Sup	1e-4	64
FLAN-T5	1e-4	64
MIMIC-T5	1e-3	64
SCIFIVE+MIMIC-T5	1e-4	64

Table 7: Best-found hyperparameter settings for down-sampling experiments on MedNLI dataset with 1% training data

• We added T5-Den and FLAN-T5 in addition to widely used T5-Sup into the comparison.

D.3. Computing Resources

We conducted our experiments on multiple 40G/80G A100 GPUs and Tesla M60 GPUs. Each run of the fine-tuning takes a few hours, except for CLIP dataset which could take approximately 10 hours for 30 epochs with multiple GPUs. However, it should be noted that these models are approaching a size that is not necessarily easy to accommodate in clinical settings.

Appendix E. Evaluation Details

We use the Python 3.9 packages sklearn==1.1.2, and evaluate==0.3.0 for evaluations.

Classification For the CLIP dataset, we followed the same evaluation strategy as Lehman2023DoWS by transforming predicted values into binary matrices, whose dimensions indicate the presence of class labels. And then used the package sklearn.metrics to calculate macro-F1 and micro-F1. On HOC dataset, we evaluate the outputs in the same manner as Phan et al., 2021 and Yasunaga et al., 2022. For stigmatizing datasets, we also used sklearn.metrics for macro-F1 scores.

Natural Language Inference For the MedNLI dataset, without additionally post-processing the outputs, we compute the F1 score and the accuracy directly between the labels and predicted values.

Question Answering We followed the same processing method as Lehman et al., 2023 utilizing SQuAD 2.0 (Rajpurkar et al., 2018).

Named Entity Recognition As instructed by Clinical-T5 (Lu et al., 2022), we referred to the evaluation of SciFive (Phan et al., 2021). BC5CDR-disease and NCBI-disease datasets label the entity by inserting "disease*" and "*disease" around disease names in the input. We convert generated sentence outputs into sequences of "B-disease", "I-disease", "O", and padding and then evaluate using the metrics sequent (Ramshaw and Marcus, 1995; Nakayama, 2018).

Appendix F. Reproducibility Experiments

We evaluated T5, MIMIC-T5, and SciFive+MIMIC-T5 on the six datasets, following the original methodologies (i.e. a single train/dev/test split) proposed in each respective paper. Results of three clinical datasets are shown in Table 9 and results of three biomedical datasets are shown in Table 10. And these results are single-run.

Appendix G. Cross-Validation Experiments

We did 5-fold cross-validation in the following procedures: we merged the training and validation datasets and then shuffled them. Within each fold, 4 subsets are for training and one subset is for validation. For down-sampling experiments, we down-sampled the training data only while keeping the validation the same per fold. In the end, all the models were evaluated on the original test dataset. The detailed performances cross models are listed in Appendix G with paired T-test results.

Specifically, 11 shows 5-fold cross validation results on MedNLI Dataset. Table 12 shows 5-fold cross validation results on RadQA Dataset. Table 13 shows 5-fold cross validation results on CLIP dataset. Table 14 shows 5-fold cross validation results on HOC dataset. Table 15 shows 5-fold cross validation results on BC5CDR-disease dataset. Table 16 shows 5-fold cross validation results on NCBI-disease dataset. Table 17 and Table 18 are results for the stigmatizing language datasets from MIMIC-IV and Hospital System separately. Table 19, Table 20, and Table 21 illustrate the outcomes of down-sampling experiments on the MIMIC-IV dataset; Table 22, Table 23, and Table 24 present the outcomes of down-sampling experiments on the Hospital System dataset.

Task	Dataset	Source	Evaluated By	Metrics
NLI	MedNLI	MIMIC	Both	Accuracy
Classif.	CLIP HOC	MIMIC Biomedical	MIMIC-T5 SciFive+MIMIC-T5	F1(Macro, Micro) F1, P, R
QA	RadQA	MIMIC	MIMIC-T5	EM, F1
NER	BC5CDR NCBI	Biomedical	SciFive+MIMIC-T5	F1, P, R

Table 8: Statistics of datasets used in MIMIC-T5 and SciFive+MIMIC-T5

	MedNLI	Rac	lQA	CLIP		
	Accuracy	EM	F1	Macro-F1	Micro-F1	
T5-Den	85.16	52.44	71.42	64.68	79.11	
T5-Sup	84.95	$\bar{5}4.07^{-}$	71.14	64.20	78.93	
MIMIC-T5						
— Ours	86.64	54.72	73.38	65.76	80.11	
— Reported	87.2	55.0	74.5	66.3	80.0	
SciFive+MIMIC-T5						
— Ours	85.79	52.28	70.74	63.79	78.24	
— Reported	85.86	N/A	N/A	N/A	N/A	

Table 9: We reproduced the results of MIMIC-T5 on three clinical datasets as reported and complemented evaluations of SciFive+MIMIC-T5 on the clinical datasets that were not used previously.

		HOC		I	BC5CDI	?	NCBI			
	F1	Р	R	F1	Р	R	F1	Р	R	
T5-Den	85.57	85.81	85.34	80.43	81.74	79.17	85.29	82.86	87.88	
T5-Sup	$-85.\overline{91}$	86.01	85.80	81.70	81.46	81.94	-82.53	82.49	82.58	
MIMIC-T5										
— Ours	81.93	82.00	81.85	78.39	77.52	79.29	77.97	77.80	78.14	
— Reported	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	
SciFive+MIMIC-T5										
— Ours	85.42	85.52	85.32	82.44	81.22	83.70	85.67	83.18	88.31s	
—Reported	84.78	85.37	84.79	80.35	79.24	81.49	86.73	86.37	87.09	

Table 10: Evaluations on three biomedical datasets in SciFive+MIMIC-T5 were also reproduced. MIMIC-T5 and T5's performances are also shown in the table.

MedNLI										
Metrics		T5-Den	T5-S	Sup	FLAN-	$\Gamma 5$	MIM	IC-T5	SciI	Five+MIMIC-T5
Accuracy Mean		85.89 ♠	84.	94	86.03 ♦		86.75 ↑ ♠0.86, ↑ ♣1.81, ↑ ♦0.7		85.64 ↓ ♠0.25, ↑ ♣0.70, ↓ ♦0	
	95% CI	[85.34, 86.24]	[84.38,	85.49]	[85.72, 86]	[5.43]	[86.27	,87.29]		[85.13, 86.37]
MedNLI-Acc	uracy-T-test									
Mod	lels	T5-Den		T5-	-Sup	FI	AN-T5	MIMIC-T5		${\bf SciFive+MIMIC-T5}$
T5-Den	-Large	-			-					-
T5-Sup	-Large	t(df=4)=-2.63,	P=.06		-		-	-		-
FLAN-T	5-Large	t(df=4)=0.31, I	P=.77 t((df=4)=3	3.17, P=.03		-	-		-
MIMIC-T	Γ5-Large	t(df=4)=1.97, I	P=.12 t((df=4)=4	1.14, P=.01	t(df=4)	=1.75, P=.15	-		-
SciFive+MIM	IIC-T5-Large	t(df=4)=-0.45,	P=.68 t((df=4)=1	.18, P=.30	t(df=4)=	-1.45, P=.22	t(df=4)=-2.18, H	P=.10	-

Table 11: 5-fold cross-validation results across models on MedNLI dataset and the t-test results among T5, MIMIC-T5, and SciFive+MIMIC-T5

RadQA										
Metrics		T5-Den	T	5-Sup	FLAN-	Т5	MIM	C-T5	Scil	Five+MIMIC-T5
EM	Mean	52.57 ♠	5	1.95 ♣	53.0; ♦	3	54.50 ↑ ♠1.92, ↑ ♣2.54, ↑ ♦1.47		53.42 ↑ ♠0.85, ↑ ♣1.47, ↑ ♦0.39	
	95% CI	[51.9, 53.27]	[51.0	4,52.95]	[51.79, 5]	4.01]	[54.04,	55.05]		[52.69, 54.22]
F1	Mean	68.90 ♠	6	8.68 ♣	70.65 •	2		.40 4.72, ↑ ♦ 2.78	↑ ♠ 1.4	70.37 47, ↑ ♣1.7, ↓ ♦0.25
	95% CI	[68.04, 69.79]	[67.8	2,69.77]	[69.76, 7]	1.34]	[72.75,	74.00]		[69.44, 71.49]
RadQA-I	EM-T-test									
Mo	dels	T5-Den		Т5	-Sup		FLAN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Der	n-Large	-			-		-	-	-	
T5-Suj	p-Large	t(df=4)=-0.91,	P = .41		-					-
FLAN-	Γ5-Large	t(df=4)=0.47, I	P = .67	t(df=4)=	1.4, P=.23		-	-		-
MIMIC-	T5-Large	t(df=4)=4.3, F	P = .01	t(df=4)=3	2.92, P=.04	t(df=	=4)=1.84, P=.14	-		-
SciFive+MIN	MIC-T5-Large	t(df=4)=1.19, I	P = .30	t(df=4)=	1.95, P=.12	t(df=	=4)=0.54, P=.62	t(df=4)=-1.97, P	P = .12	-
RadQA-l	F1-T-test									
Mo	dels	T5-Den		T5-	Sup		FLAN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Dei	n-Large	-			-		-	-		-
T5-Sup	T5-Sup-Large $t(df=4)=-0.55, P=.61$ -		-		-	-		-		
FLAN-T	AN-T5-Large $t(df=4)=2.53, P=.06$ $t(df=4)=3.65,$		8.65, P=.02		-	-		-		
MIMIC-	T5-Large	t(df=4)=5.95, P	=.004	t(df=4)=5	5.2, P=.007	t(df=	=4)=3.5, P=.02	-		-
SciFive+MIM	IIC-T5-Large	t(df=4)=2.19, F	P=.09	t(df=4)=2	2.15, P=.10	t(df=	4)=-0.32, P=.76	t(df=4)=-5.47, P	=.005	-

Table 12: 5-fold cross-validation results across models on RadQA dataset and the t-test results among T5, MIMIC-T5, and SciFive+MIMIC-T5.

CLIP									
Metrics		T5-Den	T5-Sup	FLAN-	Т5	MIMI	C-T5	SciFi	ve+MIMIC-T5
Macro-F1	Mean	63.86 ♠	62.42 ♣	64.33 ♦	1	66.0 ↑ ♠ 2.23, ↑ ♣ 3	~ ~	↓ ♠0.4,	63.46 ↑ ♣1.04, ↓ ♦0.88
	95% CI	[62.18, 65.55]	[61.37, 63.4]	[62.94, 65	5.72]	[65.84,	66.26]	[62.2, 64.76]
Micro-F1	Mean	78.62 ♠	78.38 ♣	79.40 ♦	1	79.5 ↑ ♠0.67, ↑ ♣		↓ ♠0.38	78.23 , ↓ ♣0.15, ↓ ♦1.17
	95% CI	[77.48, 79.79]	[78.03, 78.77]	[78.85, 79	0.94]	[78.51,	80.01]	[7	7.93, 78.51]
CLIP-Macro	o-F1-T-test								
Mod	lels	T5-Den	Т5	-Sup		FLAN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Den	-Large	-		-		-	-		-
T5-Sup	-Large	t(df=4)=-1.39, F	P=.24	-		-	-		-
FLAN-T	5-Large	t(df=4)=0.38, P	=.72 t(df=4)=	1.83, P=.14		-	-		-
MIMIC-T	75-Large	t(df=4)=2.15, P	'=.10 t(df=4)=5	5.86, P=.004	t(df=	=4)=2.02, P=.11	-		-
SciFive+MIM	IC-T5-Large	t(df=4)=-0.3, P	=.78 t(df=4)=	0.95, P=.39	t(df=	-4)=-0.73, P=.51	t(df=4)=-4.33, F	'= .01	-
CLIP-Micro	-F1-T-test								
Mod	lels	T5-Den	T5	-Sup		FLAN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Den	-Large	-		-		-	-		-
T5-Sup	-Large	t(df=4)=-0.34, H	P=.75	-		-	-		-
FLAN-T	5-Large	t(df=4)=1.83, P	e.14 t(df=4)=	2.55, P=.06		-	-		-
MIMIC-T	75-Large	t(df=4)=0.64, P	?=.56 t(df=4)=	1.89, P=.13	t(df=	(4)=-0.18, P=.87	-		-
SciFive+MIM	IC-T5-Large	t(df=4)=-0.67, I	P=.54 t(df=4)=-	0.49, P = .65	t(df=	(4)=-4.19, P=.01	t(df=4)=-2.19, P	·=.09	-

 $\begin{array}{ll} \text{Table 13: 5-fold cross-validation results across models on CLIP dataset and the t-test results among T5,} \\ \text{MIMIC-T5, and SciFive+MIMIC-T5} \end{array}$

HOC										
Metrics		T5-Den	-	Γ5-Sup	FLAN-	·T5		MIMI	C-T5	SciFive+MIMIC-T5
F1	Mean	84.81 •		84.76 ♣	84.79 •	9	↓ ♠2.07	82. 7, ↓ ♣ 2		85.1 ♦ 0.29, ↑ ♣ 0.34, ↑ ♦ 0.3
	95% CI	[84.26, 85.49]	[84.	.38, 85.11]	[84.22, 8	5.26]	[82.53,	82.98]	[84.81, 85.4]
P	Mean	84.72 •		84.77	84.60 ♦)	↓ ♠1.8	82. 2, ↓ ♣		85.04 ♦ 0.32, ↑ ♣ 0.27, ↑ ♦ 0.4
	95% CI	[84.04, 85.47]	[84.	.43, 85.11]	[83.94, 8	5.18]	[82.77,	83.04]	[84.67, 85.39]
R	Mean	84.91 •		84.76 •	84.99 ♦	9	↓ ♠2.3	82. 1, ↓ ♣	~ ~	85.17 ♦ 0.26, ↑ ♣ 0.41, ↑ ♦ 0.1
	95% CI	[84.49, 85.52]	[84.	.28, 85.21]	[84.47, 8]	[5.4]	[82.26,	83.01]	[84.85, 85.52]
HOC-F1	-T-test									
Mod	lels	T5-Den		T5-3	Sup		FLAN-T5		MIMIC-T5	SciFive+MIMIC-T5
T5-Den	-Large	-		-			-		-	-
T5-Sup	-Large	t(df=4)=-0.09, P	=.93	-			-		-	-
FLAN-T	5-Large	t(df=4)=-0.05, P	=.96	t(df=4)=0.	07, P=.95		-		-	-
MIMIC-7	Γ5-Large	t(df=4)=-5.75, P=	=.005	t(df=4)=-6.	86, $P=.002$	t(df=	4)=-7.02, I	P = .002	-	-
SciFive+MIM	IIC-T5-Large	t(df=4)=1.05, P=	=.35	t(df=4)=1	31, P=.26	t(df=	=4)=1.08, <i>i</i>	P = .34	t(df=4)=10.06, P=i	.001 -
HOC-P										
Mod		T5-Den		Т5-	Sup		FLAN-T	5	MIMIC-T5	SciFive+MIMIC-T5
T5-Der		-		-			-		-	-
T5-Sup		t(df=4)=0.1, P=		-			-		-	-
FLAN-T	75-Large	t(df=4)=-0.29, P		t(df=4)=-0			-		-	-
MIMIC-		t(df=4)=-4.3, P=	=.01	t(df=4)=-6.	62, P=.003	t(df=	(4) = -5.46,	P = .005	-	-
SciFive+MIN		t(df=4)=0.86, P	=.44	t(df=4)=1	19, P=.30	t(df	=4)=1.41,	P = .23	t(df=4)=7.87, P=.0	001 -
HOC-R										
Mod		T5-Den		T5-3	Sup		FLAN-T5		MIMIC-T5	SciFive+MIMIC-T5
T5-Den				-			-		-	-
T5-Sup		t(df=4)=-0.28, P		- (10 1) -			-		-	-
FLAN-T		t(df=4)=0.22, P=		t(df=4)=0.		. (10	-	2 00:	-	-
MIMIC-7		t(df=4)=-6.84, P=		t(df=4)=-6.			4)=-8.09, <i>I</i>		- (10 1) 1-1	-
SciFive+MIM	HC-T5-Large	t(df=4)=1.26, P=	=.27	t(df=4)=1.	11, P=.33	t(df=	=4)=0.57, I	P=.60	t(df=4)=10.45, P=i	.001 -

Table 14: 5-fold cross-validation results across models on HOC dataset and the t-test results among T5, MIMIC-T5, and SciFive+MIMIC-T5

BC5CDR										
Metrics			T5-Den	T5-Sup	FLA	N-T5	MI	MIC-T5	Scil	Five+MIMIC-T5
F1	Mea	n	82.90 •	82.94 ♣	83	3.30 ♦		31.17 ♣1.77, ↓ ♦2.13	↑ ♠0.7	83.61 1, ↑ ♣0.68, ↑ ♦0.31
	95%	CI [82.33, 83.38]	[82.51, 83.45]	[82.89	0,83.82]	[80.9	02, 81.53]		[83.38, 83.82]
P	Mea	n	81.67 ♠	81.62 ♣	82	2.14 ♦		80.62 ♣1.01, ↓ ♦1.52	↑ ♠1.4	83.07 1, ↑ \$ 1.44, ↑ ♦ 0.93
	95%	CI [80.77, 82.51]	[81.30, 81.96]	[81.53	3, 82.72]	[80.2	20, 80.94]		[82.51, 83.6]
R	Mea	n	84.19 •	84.29 •	84	4.50 ♦		31.73 ♣2.56, ↓ ♦2.76	↓ ♠0.0	84.17 2, ↓ ♣0.12, ↓ ♦0.32
	95%	CI [83.64, 84.63]	[83.55, 85.05]	[84.23	3,84.93]	[81.2	23, 82.31]		[83.60, 84.82]
BC5CDR-F1-	T-test									
Models			T5-Den	T5-Sup		FI	AN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Den			-	-			-	-		-
T5-Sup		t(df=	4)=0.06, P=.96	-			-	-		-
FLAN-T	5	t(df=	4)=0.82, P=.46	t(df=4)=2.16,	P = .10		-	-		-
MIMIC-T	75	t(df=4	P=-9.49, P=1.001	t(df=4)=-4.01,	P = .02	t(df=4)=	-5.29, P=.006	-		-
SciFive+MIM	IC-T5	t(df=	(4)=2.38, P=.08	t(df=4)=1.7,	P = .16	t(df=4)	=0.76, P=.49	t(df=4)=14.45, P=	j.001	-
BC5CDR-P-	T-test									
Models	3		T5-Den	T5-Sup)	Fl	LAN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Der	1		-	-			-	-		-
T5-Sup)	t(df=	4)=-0.08, P=.94	-			-	-		-
FLAN-T	75	t(df=	=4)=0.71, P=.52	t(df=4)=2.37,	P = .08		-	-		-
MIMIC-	Γ5	t(df=	4)=-3.07, P=.04	t(df=4)=-3.23	P = .03	t(df=4)=	=-3.09, P=.04	-		-
SciFive+MIN	IIC-T5	t(df=	4)=2.08, P=.11	t(df=4)=3.41,	P = .03	t(df=4)	=1.73, P=.16	t(df=4)=6.04, P=	.004	-
BC5CDR-R-	Γ-test									
Models			T5-Den	T5-Sup		FI	AN-T5	MIMIC-T5		SciFive+MIMIC-T5
T5-Den			-	-			-	-		-
T5-Sup		t(df=	4)=0.19, P=.86	-			-	-		-
FLAN-T	5	t(df=	=4)=0.8, P=.47	t(df=4)=0.78,	P = .48		-	-		-
MIMIC-T	75	t(df=4	e)=-6.28, P=.003	t(df=4)=-3.88,	P = .02	t(df=4)=	-6.14, P=.004	-		-
SciFive+MIM	IC-T5	t(df=	4)=-0.07, P=.95	t(df=4)=-0.17,	P = .88	t(df=4)=	-0.61, P=.57	t(df=4)=10.35, P=	j.001	-

Table 15: 5-fold cross-validation results across models on BC5CDR-disease dataset and the t-test results among T5, MIMIC-T5, and SciFive+MIMIC-T5

NCBI									
Metrics		T5-Den	T5-Sup	FLAN	N-T5	MIM	IC-T5	Scil	Five+MIMIC-T5
F1	Mean	85.08 ♠	84.58 •	84.4	44		.10 5.49, ↓ ♦ 5.35	↑ ♠0.3	85.43 6, ↑ ♣0.85, ↑ ♦0.99
	95% CI	[84.61, 85.46]	[84.22, 85.02]	[83.94,	85.03]	[78.28,	79.94]		[84.89, 86.06]
P	Mean	83.49 ♠	83.58 ♣	82.: ♦	20		.35 5.24, ↓ ♦ 3.85	↑ ♠0.3	83.80 1, ↑ ♣0.21, ↑ ♦1.60
	95% CI	[82.78, 84.16]	[83.13, 84.12]	[81.59,	83.04]	[77.57,	78.87]		[82.95, 84.68]
R	Mean	86.73 ♠	85.61 ♣	86.8	82		.87 5.74, ↓ ♦ 6.95	↑ ♠0.4	87.14 1, ↑ ♣1.54, ↑ ♦0.32
	95% CI	[86.31, 87.22]	[85.16, 86.15]	[86.31,	87.30]	[78.70,	81.30]		[86.86, 87.50]
NCBI-F1-T-	test								
Models		T5-Den	T5-Sup)]	FLAN-T5	MIMIC-T	5	SciFive+MIMIC-T5
T5-Den		-	-			-	-		-
T5-Sup	t	(df=4)=-0.95, P=.40	-			-	-		-
FLAN-T	5 t	(df=4)=-1.73, P=.16	t(df=4)=-0.33	, P=.76		-	-		-
MIMIC-T	5 t(c	df=4)=-12.52, P=j.00	t(df=4)=-10.1,	P=j.001	t(df=4)	=-11.66, P=j.001	-		-
SciFive+MIM	IC-T5	t(df=4)=0.94, P=.40	t(df=4)=1.71,	P = .16	t(df=4	1)=1.62, P=.18	t(df=4)=8.65, P	'=j.001	-
NCBI-P-T-	test								
Models		T5-Den	T5-Sup)		FLAN-T5	MIMIC-T5	j	SciFive+MIMIC-T5
T5-Den		-	-			-	-		-
T5-Sup	t	(df=4)=0.14, P=.90	-			-	-		-
FLAN-T	5 t	(df=4)=-1.88, P=.13	t(df=4)=-3.52	, P=.02		-	-		-
MIMIC-T	75 t(df=4)=-7.08, P=.002	t(df=4)=-14.76	, P=j.001	t(df=4)=-8.28, P=.001	-		-
SciFive+MIM	IC-T5 t	(df=4)=0.74, P=.50	t(df=4)=0.33,	P=.76	t(df=4	1)=1.93, P=.13	t(df=4)=6.65, P	=.003	-
NCBI-R-T-t	test								
Models		T5-Den	T5-Sup)	I	FLAN-T5	MIMIC-T	5	SciFive+MIMIC-T5
T5-Den		-	-			-	-		-
T5-Sup		t(df=4)=-2.0, P=.12	-			-	-		-
FLAN-T	5	t(df=4)=0.38, P=.72	t(df=4)=2.71,	P = .05		-	-		-
MIMIC-T	5 t(df=4)=-10.43, P=j.00	1 t(df=4)=-7.16,	P = .002	t(df=4)	=-12.53, P=j.001	-		-
SciFive+MIM	IC-T5	t(df=4)=1.07, P=.34	t(df=4)=3.25,	P = .03	t(df=4)=0.72, P=.51	t(df=4)=8.63, P	=j.001	-

Table 16: 5-fold cross-validation results across models on NCBI-disease dataset and the t-test results among T5, MIMIC-T5, and SciFive+MIMIC-T5

		•					
Metrics		T5-Sup	FLAN-T	5 MIM	IC-T5	SciFive	+MIMIC-T5
F1	Mean	76.04 ♣	77.44 ♦		98 , ↓ ♦ 2.46		74.65 $39, \downarrow \spadesuit 2.79$
	95% CI	[74.14, 77.94]	[74.96, 79.5	34] [72.11	,77.75]	[70.	59, 78.82]
MIMIC-I	V-Complian	ce					
Metrics		T5-Sup	FLAN-T	5 MIM	IC-T5	SciFive	+MIMIC-T5
F1	Mean	92.94 ♣	92.93 ♦		.57 , ↓ ♦ 1.36		90.98 96, ↓ ♦ 1.95
	95% CI	[91.74, 94.18]	[91.27, 94.4	48] [90.38	, 92.70]	[89.	90, 92.12]
MIMIC-I	V-Descripto	rs					
Metrics		T5-Sup	FLAN-T	5 MIM	IC-T5	SciFive	+MIMIC-T5
F1	Mean	85.73 ♣	86.31 ♦		76 , ↓ ♦ 1.55		86.28 54, ↓ ♦ 0.03
	95% CI	[83.45, 87.99]	[85.32, 87.0	03] [81.90	,88.15]	[85.0	00, 87.79]
IMIC-IV-Credibil	itv & Obstinacy-	F1-T-test					
	Models		5-Sup	TT 137 TT			
,)-Sup	FLAN-T5	MI	MIC-T5	SciFive+MIMIC-7
E	T5-Sup		- -	FLAN-T5	MI	MIC-T5	SciFive+MIMIC-7
Г	T5-Sup LAN-T5	t(df=4)=	- - -:0.64, P=.56	FLAN-T5 - -	MII	MIC-T5 - -	
M	LAN-T5 IMIC-T5		- -0.64, P=.56	FLAN-T5		MIC-T5 - - -	-
M SciFive	LAN-T5 IMIC-T5 e+MIMIC-T5	t(df=4)= $t(df=4)=$	-0.64, P=.56 -0.52, P=.63	-	÷.35	MIC-T5 0.12, P=.91	- - -
M SciFive IIMIC-IV-Compl	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test	t(df=4)= $t(df=4)=$	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t	- c(df=4)=-1.05, P= c(df=4)=-1.08, P=	=.35 =.34 t(df=4)=	- - 0.12, P=.91	
M SciFive HMIC-IV-Compl Mod	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test	t(df=4)= $t(df=4)=$	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t	- c(df=4)=-1.05, P=	÷.35	- - 0.12, P=.91	
M SciFive HMIC-IV-Compl Mod T5-S	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test els up	t(df=4)= t(df=4)= T5-Sup	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t	- c(df=4)=-1.05, P= c(df=4)=-1.08, P=	=.35 =.34 t(df=4)=	- - 0.12, P=.91	- - -
M SciFive HMIC-IV-Compl Mod T5-S FLAN	LAN-T5 IMIC-T5 p+MIMIC-T5 liance-F1-T-test els up	t(df=4)= $t(df=4)=$ $t(df=4)=$ $t(df=4)=-0.0, P=$	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t	- c(df=4)=-1.05, P= c(df=4)=-1.08, P= FLAN-T5	=.35 =.34 t(df=4)=	- - 0.12, P=.91	-
M SciFive HMIC-IV-Compl Mod T5-S	LAN-T5 IMIC-T5 =+MIMIC-T5 liance-F1-T-test els up I-T5 C-T5	t(df=4)= t(df=4)= T5-Sup	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t -1.00 =.30 t(df=4	- c(df=4)=-1.05, P= c(df=4)=-1.08, P=	=.35 =.34 t(df=4)=	- - 0.12, <i>P</i> =.91	
M SciFive SciFive MIMIC-IV-Compl Mod T5-S FLAN MIMIC SciFive+M	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test els up I-T5 C-T5 IMIC-T5	t(df=4)= t(df=4)= T5-Sup - t(df=4)=-0.0, P= t(df=4)=-1.2, P t(df=4)=-1.5, P	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t -1.00 =.30 t(df=4	- c(df=4)=-1.05, P= c(df=4)=-1.08, P= FLAN-T5 - - - 4)=-1.2, P=.30	:.35 :.34 t(df=4)= MIMIC	- - 0.12, <i>P</i> =.91	- - - SciFive+MIMIC-T - -
M SciFive HMIC-IV-Compl Mod T5-S FLAN MIMIC	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test els up f-T5 C-T5 IMIC-T5 ptors-F1-T-test	t(df=4)= t(df=4)= T5-Sup - t(df=4)=-0.0, P= t(df=4)=-1.2, P t(df=4)=-1.5, P	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t -1.00 -1.00 -1.00 t -1.00 t -1.00 t -1.00 t	- c(df=4)=-1.05, P= c(df=4)=-1.08, P= FLAN-T5 - - - 4)=-1.2, P=.30	:.35 :.34 t(df=4)= MIMIC		- - - SciFive+MIMIC-T - - -
M SciFive IIMIC-IV-Compl Mod T5-S FLAN MIMIC SciFive+M IIMIC-IV-Descri	LAN-T5 IMIC-T5 p+MIMIC-T5 liance-F1-T-test els up I-T5 C-T5 IMIC-T5 ptors-F1-T-test	t(df=4) = t(df=4) = t(df=4) = t(df=4) = -0.0, P = t(df=4) = -1.2, P = t(df=4) = -1.5, P = t(df=4) = -1.5	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t -1.00 -1.00 -1.00 t -1.00 t -1.00 t -1.00 t		MIMIO - - - - - - - - - - - - -		- - - SciFive+MIMIC-T - - -
M SciFive IIMIC-IV-Compl Mod T5-S FLAN MIMIC SciFive+M IIMIC-IV-Descri	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test els up I-T5 C-T5 IMIC-T5 ptors-F1-T-test els up	t(df=4) = t(df=4) = t(df=4) = t(df=4) = -0.0, P = t(df=4) = -1.2, P = t(df=4) = -1.5, P = t(df=4) = -1.5	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t -1.00 =.30 t(df=4) =.21 t(df=4)		MIMIO - - - - - - - - - - - - -		SciFive+MIMIC-T SciFive+MIMIC-T
M SciFive HMIC-IV-Compl Mod T5-S FLAN MIMIC SciFive+M HMIC-IV-Descri Mod T5-S	LAN-T5 IMIC-T5 e+MIMIC-T5 liance-F1-T-test els up f-T5 C-T5 IMIC-T5 ptors-F1-T-test els up	t(df=4) = t(df=4) = t(df=4) = 0.0, P = t(df=4) = -1.2, P = t(df=4) = -1.5, P = t(df=4) = -1.5, P = t(df=4) = -1.5, P = -1.5	-0.64, P=.56 -0.52, P=.63 t -0.81, P=.46 t =1.00 =.30 t(df=4 =.21 t(df=4		MIMIC - t(df=4)=-0.8		SciFive+MIMIC-T SciFive+MIMIC-T

Table 17: 5-fold cross-validation results across models on stigmatizing dataset from MIMIC-IV and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

Metrics		T5-Sup	FLAN-T5	MIMIC-	-T5 SciFiv	e+MIMIC-T5
F1	Mean	88.21 ♣	90.38 ◆	88.07 ↓ ♣ 0.15, ↓		87.31 0.9, ↓ ♦ 3.07
	95% CI	[86.12, 90.59]	[88.26, 92.33]	[83.11, 94	[82	2.11, 91.76]
Hospital	System-Con	npliance				
Metrics		T5-Sup	FLAN-T5	MIMIC-	-T5 SciFive	e+MIMIC-T5
F1	Mean	86.71 •	88.03 ♦	84.72 ↓ ♣ 2.0, ↓ •		87.21 .49, ↓ ♦ 0.82
	95% CI	[85.52, 88.19]	[86.84, 88.92]	[80.99, 88	[86]	.82, 87.63]
Hospital S	System-Desc	riptors				
Metrics		T5-Sup	FLAN-T5	MIMIC-	-T5 SciFiv	e+MIMIC-T5
F1	Mean	90.58 ♣	86.82 ♦	88.85 ↓ ♣ 1.73, ↑		89.98 0.6, ↑ ♦ 3.17
	95% CI	[89.25, 91.37]	[79.94, 90.69]	[88.24, 89	0.46] [89	0.21, 90.67]
ital System-Cred	libility & Obstin	acy-F1-T-test				
	Models		T5-Sup	FLAN-T5	MIMIC-T5	SciFive+MIMIC
	THE Comm					
	T5-Sup			-	-	-
	FLAN-T5		4)=1.02, P=.37	- -	-	-
N	FLAN-T5 MIMIC-T5	t(df=4	1)=-0.07, P=.95 t(- df=4)=-0.63, P=.5		-
M SciFiv	FLAN-T5 MIMIC-T5 ve+MIMIC-T5	t(df=4	1)=-0.07, P=.95 t(- df=4)=-0.63, P=.5 df=4)=-1.48, P=.2		- - - .83 -
M SciFiv	FLAN-T5 MIMIC-T5 re+MIMIC-T5 ompliance-F1-T	t(df=4	1)=-0.07, P=.95 t(1)=-0.34, P=.75 t(
N SciFiv	FLAN-T5 IIMIC-T5 re+MIMIC-T5 ompliance-F1-T dels	t(df=4) $t(df=4)$	1)=-0.07, P=.95 t(1)=-0.34, P=.75 t(df=4)=-1.48, P=.2	t(df=4)=-0.23, P=	
Moo SciFiv ital System-Co Moo	FLAN-T5 IIMIC-T5 re+MIMIC-T5 ompliance-F1-T dels Sup	t(df=4) $t(df=4)$	1)=-0.07, P=.95 t(1)=-0.34, P=.75 t(pp F	df=4)=-1.48, P=.2	t(df=4)=-0.23, P=	SciFive+MIMIC-
Moo SciFiv ital System-Co Moo T5-	FLAN-T5 MIMIC-T5 re+MIMIC-T5 ompliance-F1-T dels Sup N-T5	t(df=4 t(df=4 -test T5-Su	$(4)=-0.07, P=.95 ext{ t}$ $(4)=-0.34, P=.75 ext{ t}$ $(4)=-0.34, P=.75 ext{ t}$ $(5) ext{ } P=.16$	df=4)=-1.48, P=.2	t(df=4)=-0.23, P=	SciFive+MIMIC-
N SciFiv ital System-Co Moo T5- FLA	FLAN-T5 MIMIC-T5 re+MIMIC-T5 ompliance-F1-T- dels Sup N-T5	t(df=4) $t(df=4)$ $t(df=4)$	4)=-0.07, P=.95 t(4)=-0.34, P=.75 t(pp F 5, P=.16 4, P=.36 t(df=4)	LAN-T51.52, P=.20	t(df=4)=-0.23, P=	SciFive+MIMIC- - - -
SciFive+M	FLAN-T5 MIMIC-T5 re+MIMIC-T5 ompliance-F1-T- dels Sup N-T5	t(df=4) = t(df=4) = t(df=4) = 1.75 $t(df=4) = 1.75$ $t(df=4) = 1.0$ $t(df=4) = 0.55$	4)=-0.07, P=.95 t(4)=-0.34, P=.75 t(pp F 5, P=.16 4, P=.36 t(df=4)	LAN-T51.52, P=.20	MIMIC-T5	SciFive+MIMIC- - - -
SciFive+Asital System-De	FLAN-T5 IIMIC-T5 re+MIMIC-T5 ompliance-F1-Todels Sup N-T5 IC-T5 IIMIC-T5	t(df=4) = t(df=4) = t(df=4) = 1.75 $t(df=4) = 1.75$ $t(df=4) = 1.0$ $t(df=4) = 0.55$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	LAN-T51.52, P=.20	MIMIC-T5	SciFive+MIMIC
SciFive+Asital System-De	FLAN-T5 MIMIC-T5 re+MIMIC-T5 ompliance-F1-T dels Sup N-T5 IC-T5 MIMIC-T5 escriptors-F1-T dels	t(df=4) $t(df=4)$ -test $T5-Su$ $t(df=4)=1.75$ $t(df=4)=-1.0$ $t(df=4)=0.52$ -test	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	LAN-T5	MIMIC-T5 t(df=4)=1.06, P=.35	SciFive+MIMIC
SciFive SciFive Mod T5- FLAI MIMI SciFive+Mod Mod Mod Mod Mod Mod Mod SciFive Mod Mod Mod SciFive Mod Mod Mod SciFive Mod Mod Mod SciFive Mod Mod Mod SciFive Mod Mod Mod SciFive Mod Mod Mod Mod SciFive Mod Mod Mod Mod Mod Mod SciFive Mod Mod Mod Mod Mod Mod Mod Mo	FLAN-T5 IIMIC-T5 re+MIMIC-T5 mpliance-F1-T dels Sup N-T5 IC-T5 IIMIC-T5 escriptors-F1-T dels Sup	t(df=4) $t(df=4)$ -test $T5-Su$ $t(df=4)=1.75$ $t(df=4)=-1.0$ $t(df=4)=0.52$ -test	4)=-0.07, P=.95 t(4)=-0.34, P=.75 t(4)=-0.34, P=.75 t(5, P=.16 4, P=.36 t(df=4) 2, P=.63 t(df=4)	LAN-T5	MIMIC-T5 t(df=4)=1.06, P=.35	SciFive+MIMIC
SciFive+Notal System-De Moo	FLAN-T5 IMMIC-T5 re+MIMIC-T5 ompliance-F1-T dels Sup N-T5 IC-T5 IMMIC-T5 escriptors-F1-T dels Sup	t(df=4) t(df=4) t(df=4) -test T5-Su t(df=4)=1.75 t(df=4)=-1.0 t(df=4)=0.52 -test T5-Su	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	LAN-T5	MIMIC-T5 t(df=4)=1.06, P=.35	SciFive+MIMIC- - - -

Table 18: 5-fold cross-validation results across models on stigmatizing dataset from Hospital System and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

MIMIC-IV-Credibility & Obstinacy - 25% downsampling data

	WIIIVII O I V	Creatonity	& Obbillacy	2070 downsan	npmig date	U		
	Metrics		T5-Sup	FLAN-T5	MIMI	C-T5	SciFive	+MIMIC-T5
_	F1	Mean	63.81 ♣	71.92 ♦	64. ↑ ♣ 0.63,			57.67 $15, \downarrow \spadesuit 14.25$
		95% CI	[62.15, 65.77]	[69.6, 74.16]	[61.69,	66.63]	[54.	39,61.06]
_	MIMIC-IV	7-Complian	ce- 25% downs	ampling data				
_	Metrics		T5-Sup	FLAN-T5	MIMI	C-T5	SciFive-	+MIMIC-T5
-	F1	Mean	89.6 ♣	91.44 ◆	86.0 ↓ ♣ 3.58,			89.58 02, ↓ ♦ 1.86
		95% CI	[86.8, 92.19]	[89.15, 93.09]	[84.63,	87.5]	[87.1	19,92.12]
-	MIMIC-I	V-Descripto	ors- 25% downs	sampling data				
	Metrics		T5-Sup	FLAN-T5	MIMI	C-T5	SciFive+	-MIMIC-T5
	F1	Mean	76.83 ♣	78.21 ♦	75.5 ↓ ♣ 1.52,			3.46 $3, \uparrow •4.25$
		95% CI	[72.49, 79.4]	[77.29, 79.19]	[72.32,	77.52]	[82.0	[2,82.85]
иіміс	C-IV-Credibility	& Obstinacy-F	'1-T-test					
	Me	odels	Т	5-Sup	FLAN-T5	M	IMIC-T5	SciFive+MIMIC-T5
	Т5	-Sup		-	-		-	-
	FLA	AN-T5	t(df=4)=	=3.37, P=.03	-		-	-
	MIN	IIC-T5	t(df=4)=	=0.42, P=.69 t(df=	=4)=-2.66, P=.	.06	-	-
		MIMIC-T5	t(df=4)=	-3.08, P=.04 t(df=	=4)=-4.9, P=.0	008 t(df=4)	=-2.42, P=.07	7 -
MIMI	IC-IV-Complia							
	Models		T5-Sup	FL.	AN-T5	MIMI	IC-T5	SciFive+MIMIC-T5
	T5-Suj		-		-	-	-	-
	FLAN-T		t(df=4)=1.2, H		-	-	-	-
	MIMIC-		t(df=4)=-4.16,		-4.54, P=.01	-	-	-
	SciFive+MIN		t(df=4)=-0.01,	P = .99 t(df = 4) =	-1.47, P=.21	t(df=4)=1	1.9, $P=.13$	-
MIMI	IC-IV-Descript							
	Models		T5-Sup	FLA	N-T5	MIMI	C-T5	SciFive+MIMIC-T5
	T5-Sup		-		-	-		-
	FLAN-T		t(df=4)=0.72, P		-	-		-
	MIMIC-7		t(df=4)=-0.82, P		1.67, P=.17	<u>-</u>		-
	SciFive+MIM	HC TE	t(df=4)=2.66, P	=.06 $t(df=4)=5$.	01 D 00F	t(df=4)=4.6	24 D 010	_

Table 19: 5-fold cross-validation results across models on stigmatizing dataset from MIMIC-IV with 25% downsampled training data and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

MIMIC-I	· Creammi					~	3 573 57 67 575
Metrics		T5-Sup	FLAN-T5	MIMI	IC-T5	SciFive	+MIMIC-T5
F1	Mean	51.62 ♣	66.33 ♦	58. ↑ ♣ 6.96,	.58 ↓ ♦ 7.76		53.14 $52, \downarrow \spadesuit 13.2$
	95% CI	[42.57, 58.72]	[64.37, 67.61]	[53.25,	64.66]	[49.	77, 56.59]
MIMIC-I	V-Complian	ce- 5% downsai	mpling data				
Metrics		T5-Sup	FLAN-T5	MIMI	[C-T5	SciFive	+MIMIC-T5
F1	Mean	70.2 ♣	82.33 ♦	72. ↑ ♣ 2.78,	.98 ↓ ♦ 9.35		69.36 34, ↓ ♦ 12.97
	95% CI	[67.48, 72.75]	[74.22, 87.16]	[70.25,	76.61]	[55.	76, 79.52]
MIMIC-I	V-Descripto	rs- 5% downsa	mpling data				<u> </u>
Metrics	-	T5-Sup	FLAN-T5	MIMI	C-T5	SciFive-	+MIMIC-T5
F1	Mean	61.5 ♣	63.16 ♦	62. ↑ ♣ 0.99,			67.3 8, ↑ ♦ 4.14
	95% CI	[59.2, 63.54]	[49.6, 71.45]	[60.26,	64.6]	[64.9	92,69.49]
MIC-IV-Credibili	tv & Obstinacy-l	F1-T-test					
	Models		5-Sup	FLAN-T5	MI	MIC-T5	SciFive+MIMIC-
Г	75-Sup		-	-		-	-
FI	LAN-T5	` '	2.78, P=.05	-		-	-
MI	MIC-T5	t(df=4)=	1.35, P = .25 $t(df =$	(-4)=-2.55, P=	06	-	-
	+MIMIC-T5	t(df=4)=	0.27, P=.80 t(df=	4)=-5.36, P=.0	006 t(df=4)=	=-1.25, P=.28	-
MIC-IV-Compli		TET C	Tel A	N Dr	MIMI	O mr	C.:E: MIMIC !
Mode		T5-Sup	FLA	N-T5	MIMI	U-15	SciFive+MIMIC-
T5-Sı FLAN-		t(df=4)=2.4, P	= 07	- -	-		
MIMIC	'-T5	t(df=4)=0.93, H		1.83, P=.14	-		_
MIMIC SciFive+MI		t(df=4)=0.93, H t(df=4)=-0.11, H	P=.40 t(df=4)=-1	1.83, <i>P</i> =.14 1.38, <i>P</i> =.24	t(df=4)=-0.	36, P=.73	-
SciFive+MI	MIC-T5		P=.40 t(df=4)=-1		t(df=4)=-0.	36, P=.73	-
	MIC-T5 etors-F1-T-test		P=.40 t(df=4)=-1 P=.92 t(df=4)=-1		t(df=4)=-0.	,	- SciFive+MIMIC-7
SciFive+MI MIC-IV-Descrip	MIC-T5 etors-F1-T-test	t(df=4)=-0.11, I	P=.40 t(df=4)=-1 P=.92 t(df=4)=-1	1.38, P=.24	,	,	- SciFive+MIMIC-
SciFive+MI MIC-IV-Descrip Mode	MIC-T5 otors-F1-T-test ls	t(df=4)=-0.11, I	P=.40 t(df=4)=-1 P=.92 t(df=4)=-1 FLA	1.38, P=.24	,	,	SciFive+MIMIC-
SciFive+MI MIC-IV-Descrip Mode T5-Su	MIC-T5 otors-F1-T-test ls up	t(df=4)=-0.11, I	P=.40 t(df=4)=-1 P=.92 t(df=4)=-1 FLA	1.38, P=.24	,	,	- SciFive+MIMIC- - -

Table 20: 5-fold cross-validation results across models on stigmatizing dataset from MIMIC-IV with 5% downsampled training data and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

Metrics		T5-Sup	FLAN-T5	MIMI	C-T5	SciFive	+MIMIC-T5
F1	Mean	27.7 ♣	51.02 ◆	37. ↑ ♣ 9.38,		↑ ♣ 5.′	33.49 78, ↓ ♦ 17.54
	95% CI	[22.94, 32.17]	[45.89, 55.52	2] [29.74,	42.96]	[27.	39, 40.18]
MIMIC-IV	V-Compliano	ce- 1% downsar	npling data				
Metrics		T5-Sup	FLAN-T5	MIMI	C-T5	SciFive	e+MIMIC-T5
F1	Mean	54.32 ♣	66.18 ♦	42 ↓ ♣ 12.22,		↓ ♣ 3.	50.44 88, ↓ ♦ 15.75
	95% CI	[49.27, 59.91]	[65.28, 66.94	[38.57,	46.91]	[43	5.57, 56.06]
MIMIC-I	V-Descripto	rs- 1% downsar	npling data				
Metrics		T5-Sup	FLAN-T5	MIMIO	C-T5	SciFive	+MIMIC-T5
F1	Mean	43.41 ♣	47.85 ♦	29.0 \$\pm\$14.32,		↓ ♣8.	35.14 $27, \downarrow \spadesuit 12.7$
	95% CI	[40.3, 46.32]	[42.16, 55.4]	[25.76,	33.03]	[29.	11, 41.98]
IIC-IV-Credibil	lity & Obstinacy	-F1-T-test					
	Models		Г5-Ѕир	FLAN-T5	MIM	IIC-T5	SciFive+MIMIC
	T5-Sup		-	-		-	-
F	LAN-T5	t(df=4)	=5.92, P=.004	-		-	-
	IMIC-T5	t(df=4)	=3.43, P=.03 t	(df=4)=-2.43, P=.	07	-	-
	e+MIMIC-T5	, ,	=1.77, P=.15 to	(df=4)=-3.58, P=.	02 t(df=4)=-	1.76, P=.15	-
MIC-IV-Comp	Hance- F. I Ites	t					0 IFI 3 IFI IF O
Mod			- F	T. A N_T5	MIMIC	_T5	Sci Five + MIMIC
Mod	els	T5-Sup	F	LAN-T5	MIMIC	-T5	SciFive+MIMIC
Mod T5-S FLAN	els up			TLAN-T5 - -	MIMIC -	-T5	SciFive+MIMIC
T5-S	els Jup J-T5	T5-Sup	P=.03	LAN-T5	MIMIC - -	-T5	SciFive+MIMIC
T5-S FLAN	els lup I-T5 C-T5	T5-Sup - t(df=4)=3.46, A	P=.03 P=.001 t(df=4):	-	MIMIC t(df=4)=2.98		SciFive+MIMIC
T5-S FLAN MIMIO SciFive+M	els lup I-T5 C-T5	T5-Sup t(df=4)=3.46, A t(df=4)=-8.5, F t(df=4)=-1.14,	P=.03 P=.001 t(df=4):	- 9.03, P=i.001	- - -		SciFive+MIMIC
T5-S FLAN MIMIO SciFive+M	els lup I-T5 C-T5 IMIC-T5 ptors-F1-T-tes	T5-Sup t(df=4)=3.46, A t(df=4)=-8.5, F t(df=4)=-1.14,	P=.03 P=.001 t(df=4): P=.32 t(df=4)	- 9.03, P=i.001	- - -	8, <i>P</i> =.04	SciFive+MIMIC SciFive+MIMIC
T5-S FLAN MIMIO SciFive+M MIC-IV-Descri	els fup I-T5 C-T5 IMIC-T5 ptors-F1-T-tes els	T5-Sup t(df=4)=3.46, A t(df=4)=-8.5, F t(df=4)=-1.14,	P=.03 P=.001 t(df=4): P=.32 t(df=4)	- 9.03, P=i.001)=-4.2, P=.01	t(df=4)=2.9	8, <i>P</i> =.04	- - - -
T5-S FLAN MIMIO SciFive+M MIC-IV-Descri Mod	els fup I-T5 C-T5 IMIC-T5 ptors-F1-T-tes els	T5-Sup t(df=4)=3.46, A t(df=4)=-8.5, F t(df=4)=-1.14,	P=.03 P=.001 t(df=4): P=.32 t(df=4) F	- 9.03, P=i.001)=-4.2, P=.01	t(df=4)=2.90	8, <i>P</i> =.04	- - - -
T5-S FLAN MIMIO SciFive+M MIC-IV-Descri Mod T5-S	els fup I-T5 C-T5 IMIC-T5 ptors-F1-T-tes els up I-T5	T5-Sup t(df=4)=3.46, A t(df=4)=-8.5, F t(df=4)=-1.14, t T5-Sup	P=.03 P=.001 t(df=4): P=.32 t(df=4) F	- 9.03, P=i.001)=-4.2, P=.01	t(df=4)=2.90	8, <i>P</i> =.04	- - -

Table 21: 5-fold cross-validation results across models on stigmatizing dataset from MIMIC-IV with 1% downsampled training data and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

Metrics		T5-Sup	FLAN-T	5 MIMIC	C-T5	SciFive-	+MIMIC-T5
F1	Mean	85.1 ♣	86.59 ♦	75.9 ↓ ♣ 9.19, ↓			84.7 4, ↓ ♦ 1.89
	95% CI	[81.26, 87.88]	[83.57, 88.9	[71.71, 8	80.43]	[80.5	58, 88.32]
Hospital	System-Con	npliance- 25% d	lownsamplin	ng data			
Metrics		T5-Sup	FLAN-T	C5 MIMIC	C-T5	SciFive+	MIMIC-T5
F1	Mean	82.52 ♣	85.3 ♦	79.3 ↓ ♣ 3.2, ↓			1.53 9, ↓ ♦ 3.77
	95% CI	[78.41, 84.92]	[82.98, 87.	65] [76.29, 8	33.15]	[75.26	5, 86.29]
Hospital	System-Desc	criptors- 25% do	ownsampling	g data			
Metrics		T5-Sup	FLAN-T	5 MIMIC	C-T5	SciFive+	-MIMIC-T5
F1	Mean	86.45 ♣	85.98 ♦	87.1 ↑ ♣ 0.75, 1	-		6.51 6, ↑ ♦ 0.53
	95% CI	[85.24, 88.16]	[85.14, 86.0	[85.32, 8	0 41	[07.0	0 00 14]
	00,00=	[00.21,00.10]	[00.14,00.0	Jaj [65.5∠, c	9.41	[85.0	9,88.14]
oital System-Cre		. , ,	[00.14,00.0	[65.52,6	9.41]	[85.0	9,88.14]
bital System-Cre	dibility & Obstin Models	. , ,	T5-Sup	[65.52, e	,	[85.0 MIC-T5	
pital System-Cre	dibility & Obstin	. , ,		J L /	,		
-	dibility & Obstin Models	acy-F1-T-test		J L /	,		
-	dibility & Obstin Models T5-Sup	acy-F1-T-test t(df=	T5-Sup	FLAN-T5	MI	MIC-T5	SciFive+MIMIC
SciFi	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5	acy-F1-T-test $t(\mathrm{df} = \\ t(\mathrm{df} = \\ t($	T5-Sup - - -(4)=0.54, P=.62	FLAN-T5	MI	MIC-T5	SciFive+MIMIC
SciFi pital System-C	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T	acy-F1-T-test $t(\mathrm{df}=$ $t(\mathrm{df}=$ $t(\mathrm{df}=$ -test	T5-Sup4)=0.54, P=.62 4)=-2.36, P=.08 4)=-0.15, P=.89	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P=	MI :.04 :.62 t(df=4):	MIC-T5	SciFive+MIMIC
SciFi pital System-C	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5	acy-F1-T-test $t(\mathrm{df} = \\ t(\mathrm{df} = \\ t($	T5-Sup4)=0.54, P=.62 4)=-2.36, P=.08 4)=-0.15, P=.89	FLAN-T5 - t(df=4)=-2.94, P=	MI	MIC-T5	SciFive+MIMIC
SciFi pital System-C	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T	acy-F1-T-test $t(\mathrm{df}=$ $t(\mathrm{df}=$ $t(\mathrm{df}=$ -test	T5-Sup4)=0.54, P=.62 4)=-2.36, P=.08 4)=-0.15, P=.89	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P=	MI :.04 :.62 t(df=4):	MIC-T5	SciFive+MIMIC
SciFi pital System-C Mc T5-	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T	acy-F1-T-test $t(\mathrm{df}=$ $t(\mathrm{df}=$ $t(\mathrm{df}=$ -test	T5-Sup	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P=	MI :.04 :.62 t(df=4):	MIC-T5	SciFive+MIMIC
SciFi pital System-C Mc T5-	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-Todels -Sup	t(df= t(df= t(df= -test	T5-Sup	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P=	MI :.04 :.62 t(df=4):	MIC-T5	SciFive+MIMIC
SciFi SciFi Mc T5- FLA MIM	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-Todels -Sup	acy-F1-T-test $t(df=\\t(df=\\t(df=\\t(df=-test)$ $T5-Su$ $t(df=4)=1.79$	T5-Sup	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P= FLAN-T5 -	MI :.04 :.62 t(df=4):	MIC-T54.8, P=.009	SciFive+MIMIC SciFive+MIMIC
SciFi SciFi Mc T5 FLA MIM SciFive+1	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-Todels -Sup N-T5	acy-F1-T-test $t(df = t(df = $	T5-Sup -4)=0.54, P =.62 4)=-2.36, P =.08 4)=-0.15, P =.89 Tp T7, P =.19 T (df 8, P =.48 T (df	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P= FLAN-T5	MI = .04 = .62 t(df=4): MIMIO	MIC-T54.8, P=.009	SciFive+MIMIC SciFive+MIMIC
SciFive+I	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T odels -Sup .N-T5 IC-T5 MIMIC-T5	acy-F1-T-test $t(df = t(df =$	T5-Sup -4)=0.54, P =.62 4)=-2.36, P =.08 4)=-0.15, P =.89 Tp T7, P =.19 T (df 8, P =.48 T (df	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P= FLAN-T5	MI = .04 = .62 t(df=4): MIMIO	MIC-T54.8, P=.009 C-T5	SciFive+MIMIC SciFive+MIMIC
SciFi pital System-C Mc T5 FLA MIM SciFive+I pital System-D	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 we+MIMIC-T5 ompliance-F1-T odels -Sup N-T5 IC-T5 MIMIC-T5 escriptors-F1-T	acy-F1-T-test $t(df = t(df = $	T5-Sup -4)=0.54, P =.62 4)=-2.36, P =.08 4)=-0.15, P =.89 Tp T7, P =.19 T (df 8, P =.48 T (df	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P= FLAN-T5	MIMIO MIMIO t(df=4)=0.	MIC-T54.8, P=.009 C-T5	SciFive+MIMIC SciFive+MIMIC
SciFi pital System-C T5 FLA MIM SciFive+ pital System-D Mc T5	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ove+MIMIC-T5 odels -Sup N-T5 IC-T5 MIMIC-T5 dels escriptors-F1-Todels	acy-F1-T-test $t(df = t(df = $	T5-Sup	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P= FLAN-T5	MIMIO MIMIO t(df=4)=0.	MIC-T54.8, P=.009 C-T5	SciFive+MIMIC SciFive+MIMIC
SciFipital System-Complete System-Complete System-Double System-S	dibility & Obstin Models T5-Sup FLAN-T5 MIMIC-T5 ompliance-F1-T odels -Sup N-T5 IC-T5 MIMIC-T5 escriptors-F1-T odels -Sup	t(df= t(df= t(df= t(df= -test T5-Su t(df=4)=-1.5 t(df=4)=-0.7 -test T5-Su	T5-Sup4)=0.54, P=.62 4)=-2.36, P=.08 4)=-0.15, P=.89	FLAN-T5 - t(df=4)=-2.94, P= t(df=4)=-0.54, P= FLAN-T5	MIMIO MIMIO t(df=4)=0.	MIC-T54.8, P=.009 C-T5	SciFive+MIMIC SciFive+MIMIC

Table 22: 5-fold cross-validation results across models on stigmatizing dataset from Hospital System with 25% downsampled training data and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

Metrics		T5-Sup	FLAN-T5	MIMIC-	-T5	SciFive-	+MIMIC-T5
F1	Mean	48.94 ♣	75.66 ♦	48.88 ↓ ♣ 0.06, ↓			50.2 $6, \downarrow \spadesuit 25.46$
	95% CI	[43.03, 55.34]	[67.77, 82.11	[46.72, 5]	0.4]	[48.	32, 52.7]
Hospital	System - Co	ompliance - 5%	downsamplin	g data			
Metrics		T5-Sup	FLAN-T5	MIMIC-	-T5	SciFive+	-MIMIC-T5
F1	Mean	58.62 ♣	71.09 ♦	51.82 ↓ ♣ 6.8, ↓ ∢			66.84 8, ↓ ♦ 14.25
	95% CI	[55.48, 60.66]	[63.96, 76.1	4] [49.19, 54	1.12]	[52.6	5,60.79]
Hospital S	System - Des	scriptors - 5% d	lownsampling	g data			
Metrics		T5-Sup	FLAN-T5	MIMIC-	-T5	SciFive-	+MIMIC-T5
F1	Mean	69.95	79.82	69.71 ↓ ♣0.24, ↓			76.41 46, ↓ ♦ 3.41
	95% CI	[68.13, 71.33]	[77.17, 82.3]	[63.77, 73	3.81]		0,77.68]
ital System-Cre	dibility & Obstin	acv-F1-T-test					
ital System-Cre	dibility & Obstin Models	acy-F1-T-test	T5-Sup	FLAN-T5	MIN	IIC-T5	SciFive+MIMIC
ital System-Cre	-	acy-F1-T-test	T5-Sup	FLAN-T5	MIN	MIC-T5	SciFive+MIMIC
-	Models		T5-Sup - - 	FLAN-T5	MIN	/IC-T5 - -	SciFive+MIMIC
	Models T5-Sup	t(df=	-	FLAN-T5 - t(df=4)=-5.36, P=.0		-	-
1	Models T5-Sup FLAN-T5	t(df=	-4)=3.89, P=.02	-	06	-	- - -
] SciFi	Models T5-Sup FLAN-T5 MIMIC-T5	t(df= t(df= t(df=	- (-4)=3.89, P=.02 (4)=-0.02, P=.99	- t(df=4)=-5.36, P=.0	06	-	- - -
l SciFi vital System-C	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5	t(df= t(df= t(df=	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70	- t(df=4)=-5.36, P=.0	06	- - 0.93, P=.40	
SciFi Stal System-C Mo	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T	t(df= t(df= t(df=	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70	t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0	06 01 t(df=4)=	- - 0.93, P=.40	-
SciFi SciFal System-C Mc	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T	t(df= t(df= t(df=	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70	t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0	06 01 t(df=4)=	- - 0.93, P=.40	SciFive+MIMIC-
SciFi SciFi Dital System-C Mo T5 FLA	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T dels -Sup	t(df= t(df= t(df= T-test	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70 up 2, P=.07	t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0	06 01 t(df=4)=	- - 0.93, P=.40	SciFive+MIMIC-
SciFi SciFi Mc T5 FLA MIM	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T odels -Sup N-T5	t(df= t(df= t(df= T-test T5-Si - t(df=4)=2.5	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70 -4)=0.42, P=.70 -2, P=.07 -3, P=.009 t(df=	- t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0 FLAN-T5	06 01 t(df=4)=	- - - 0.93, P=.40	SciFive+MIMIC
SciFive+	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T odels -Sup N-T5	t(df= t(df= t(df= t(df= T-test T5-St - t(df=4)=2.5 t(df=4)=-4.8 t(df=4)=-1.1	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70 -4)=0.42, P=.70 -2, P=.07 -3, P=.009 t(df=	- t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0 FLAN-T5 	06 01 t(df=4)= MIMIC - -	- - - 0.93, P=.40	SciFive+MIMIC
SciFive+Doital System-D	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T odelsSup N-T5 IIC-T5 MIMIC-T5	t(df= t(df= t(df= t(df= T-test T5-St - t(df=4)=2.5 t(df=4)=-4.8 t(df=4)=-1.1	24)=3.89, P=.02 4)=-0.02, P=.99 4)=0.42, P=.70 up 22, P=.07 3, P=.009 t(df=17, P=.31 t(df=17, P=.31)	- t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0 FLAN-T5 	06 01 t(df=4)= MIMIC - -		SciFive+MIMIC
SciFi SciFive+ SciFive+ Mo Mo Mo Mo Mo Mo Mo Mo Mo M	Models T5-Sup FLAN-T5 MIMIC-T5 ve+MIMIC-T5 ompliance-F1-T odels -Sup N-T5 IIC-T5 MIMIC-T5 escriptors-F1-T	$t(df=\\ t(df=\\ t(df=\\ -test)$ $T5-St$ $-\\ t(df=4)=-2.5$ $t(df=4)=-4.8$ $t(df=4)=-1.1$ $-test$	24)=3.89, P=.02 4)=-0.02, P=.99 4)=0.42, P=.70 up 22, P=.07 3, P=.009 t(df=17, P=.31 t(df=17, P=.31)	- t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0 FLAN-T5 - - -4)=-5.18, P=.007 =4)=-3.12, P=.04	06 01 t(df=4)= MIMIC t(df=4)=2.8		SciFive+MIMIC-
SciFive+: bital System-C T5 FLA MIM SciFive+: bital System-D Mo	Models T5-Sup FLAN-T5 MIMIC-T5 ove+MIMIC-T5 ompliance-F1-T odels -Sup N-T5 IIC-T5 MIMIC-T5 escriptors-F1-T odels	$t(df=\\ t(df=\\ t(df=\\ -test)$ $T5-St$ $-\\ t(df=4)=-2.5$ $t(df=4)=-4.8$ $t(df=4)=-1.1$ $-test$	-4)=3.89, P=.02 4)=-0.02, P=.99 -4)=0.42, P=.70 up 2, P=.07 3, P=.009 t(df=17, P=.31 t(df=119)	- t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0 FLAN-T5 - - -4)=-5.18, P=.007 =4)=-3.12, P=.04	06 01 t(df=4)= MIMIC t(df=4)=2.8		SciFive+MIMIC-
SciFi SciFi Stal System-C Mc T5 FLA MIM SciFive+: Sital System-D Mc T5- FLA	Models T5-Sup FLAN-T5 MIMIC-T5 ove+MIMIC-T5 ompliance-F1-T odels -Sup N-T5 IIC-T5 MIMIC-T5 escriptors-F1-T odels -Sup	t(df= t(df= t(df= T-test T5-St t(df=4)=-2.5 t(df=4)=-1.1 -test	24)=3.89, P=.02 4)=-0.02, P=.99 4)=0.42, P=.70 24)=0.42, P=.70 25, P=.07 26, P=.07 27, P=.31 t(df=1) 28, P=.004	- t(df=4)=-5.36, P=.0 t(df=4)=-4.42, P=.0 FLAN-T5 - - -4)=-5.18, P=.007 =4)=-3.12, P=.04	06 01 t(df=4)= MIMIC t(df=4)=2.8		SciFive+MIMIC-

Table 23: 5-fold cross-validation results across models on stigmatizing dataset from Hospital System with 5% downsampled training data and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5

Hospital System-Credibility & Obstinacy - 1% downsampling data

Metrics		T5-Sup	FLAN-T5	MIMIC-	T5 SciFive	e+MIMIC-T5
F1	Mean	23.29 ♣	54.45 ♦	34.95 ↑ ♣ 11.66, ↓		31.79 $5, \downarrow \spadesuit 22.66$
	95% CI	[17.55, 27.46]	[45.76, 61.9	1] [26.12, 46	[25]	45, 38.43]
Hospital S	System-Com	pliance- 1% do	wnsampling	data		
Metrics		T5-Sup	FLAN-T5	MIMIC-	T5 SciFive	+MIMIC-T5
F1	Mean	45.44 ♣	70.6 ♦	41.92 ↓ ♣ 3.52, ↓ ♦		38.52 92, ↓ ♦ 32.08
	95% CI	[40.34, 52.88]	[68.81, 72.0	5] [36.33, 46	[33	42, 42.52]
Hospital S	System-Desc	riptors- 1% dov	vnsampling o	lata		
Metrics		T5-Sup	FLAN-T5	MIMIC-	T5 SciFive	+MIMIC-T5
F1	Mean	53.78 ♣	58.36 ♦	44.52 ↓ ♣ 9.26, ↓ ♦		55.52 $74, \downarrow \spadesuit 2.83$
	95% CI	[45.07, 60.86]	[52.25, 64.7	5] [32.8, 54.	.06] [53	29, 58.91]
oital System-Cre	dibility & Obstin	acy-F1-T-test				
	Models	·	T5-Sup	FLAN-T5	MIMIC-T5	SciFive+MIMIC-
	T5-Sup		-	-	-	-
	FLAN-T5	`	4)=4.77, P=.009	-	-	-
	MIMIC-T5	`	(4)=1.39, P=.24	t(df=4)=-5.22, P=.00		-
	ve+MIMIC-T5 ompliance-F1-T-		(4)=1.49, P=.21	t(df=4)=-6.22, P=.00	03 $t(df=4)=-0.56, P=.6$	1 -
<u> </u>	dels	T5-Suj	p	FLAN-T5	MIMIC-T5	SciFive+MIMIC-
Т5-	·Sup	-		-	-	-
FLA	N-T5	t(df=4)=6.2,	P=.003	-	-	-
MIM	IC-T5	t(df=4)=-0.85	5, P=.44 t(df=	4)=-9.06, P=j.001	-	-
SciFive+l	MIMIC-T5	t(df=4)=-1.63	3, P=.18 t(df=-	4)=-14.93, P=j.001	t(df=4)=-1.02, P=.36	-
pital System-D	escriptors-F1-T	-test				
Mo	odels	T5-S1	up	FLAN-T5	MIMIC-T5	SciFive+MIMIC-7
	-Sup	-		-	-	-
FLA	N-T5	t(df=4)=0.6	3, P=.56		-	
MIM	IC-T5	t(df=4)=-2.7	76, P=.05 t(df=	=4)=-1.67, P=.17	-	-

Table 24: 5-fold cross-validation results across models on stigmatizing dataset from Hospital System with 1% downsampled training data and the t-test results among T5-Sup, FLAN-T5, MIMIC-T5, and SciFive+MIMIC-T5