BioGPT: Generative Pre-trained Transformer for Biomedical Text Generation and Mining

Renqian Luo, Liai Sun, Yingce Xia, Tao Qin, Sheng Zhang, Hoifung Poon, and Tie-Yan Liu *Microsoft Research, October - 2022*

Bhautik Lukhi Advanced AI in Biomedicine(Graded) 21st November, 2024

Agenda

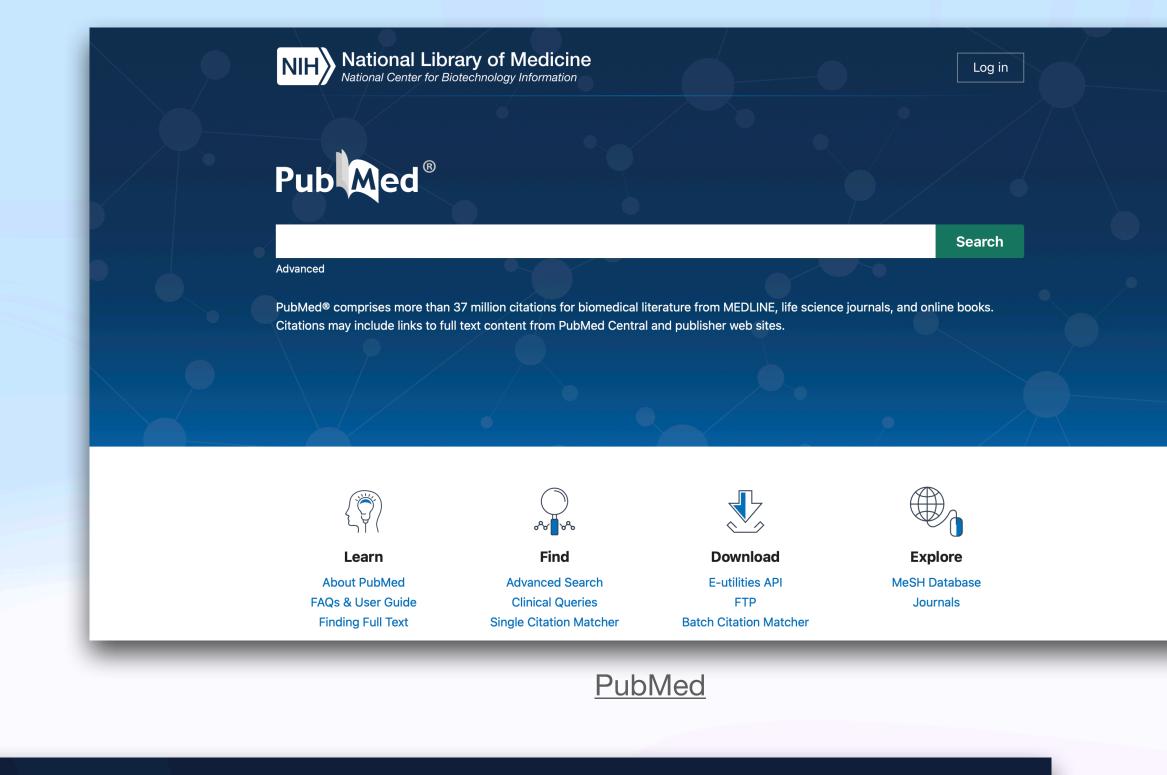
- Motivation
- Biomedical NLP Tasks
- BERT and GPT
- Introduction to BioGPT
- Architecture, Training and Fine-tuning
- Results on Biomedical Tasks
- BioGPT in Action

Why

- Millions of Research Articles
- PubMed
- Semantic Scholar
- PMC
- Arxiv

.





SEMANTIC SCHOLAR

A free, Al-powered research tool for scientific literature

Search 222,191,906 papers from all fields of science

Try: Sylvia T. Ceyer • Influenza • Metaphysics

SEMANTIC SCHOLAR



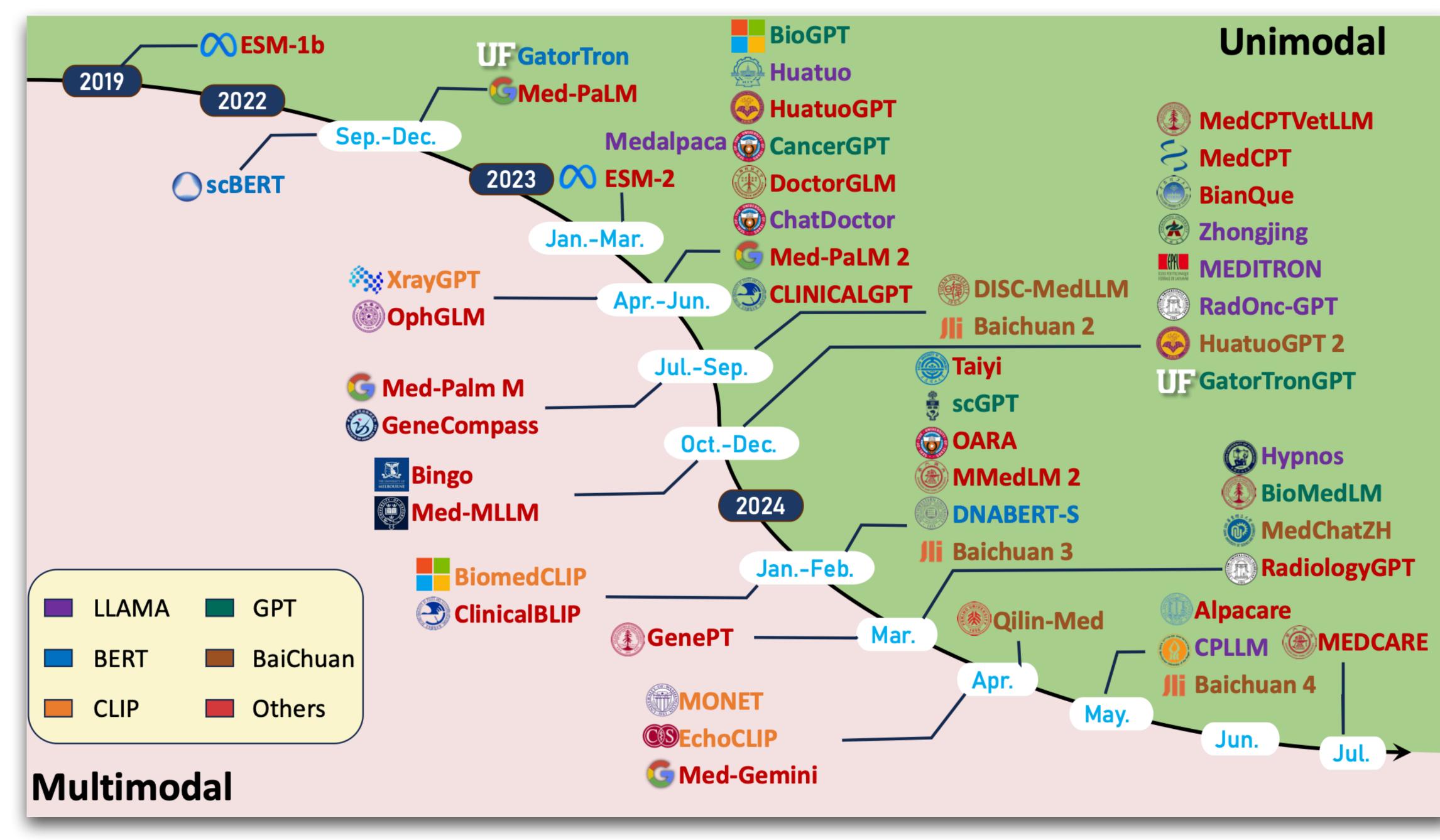
NLP in Biomedicine

- Great at extracting insights from structured/unstructured biomedical text.
- Exceptional capabilities in complex language understanding and generation tasks.
- Applications:
 - Drug discovery
 - Clinical therapy enhancement
 - Pathology research

Biomedical NLP Tasks

- **Relation Extraction:** Identifying relationships between entities in text.
- **Question Answering:** Providing answers based on biomedical literature.
- **Document Classification:** Categorizing documents into predefined classes.
- Text Generation: Generating relevant biomedical text based on prompts.
- Named Entity Recognition: Identifying and classifying key entities.
- Text Summarization: Condensing lengthy articles into concise summaries.





A Survey for Language Models in Biomedicine (Wang et al. 2024)



Pre-trained models for Bio-medicine

Model	Pros	Cons
BERT	Pretrained on massive data	General Domain
BioBERT	Continue pre-trained on bio domain Shared vocab with general dor	
BlueBERT	Continue pre-trained on bio domain	Shared vocab with general domain
SciBERT	Pretrained on Science domain Out-domain knowledge	
PubMedBERT	Pretrained on bio domain Encoder only architecture	
ELECTRAMed	Pretrained on bio domain Encoder only architecture	

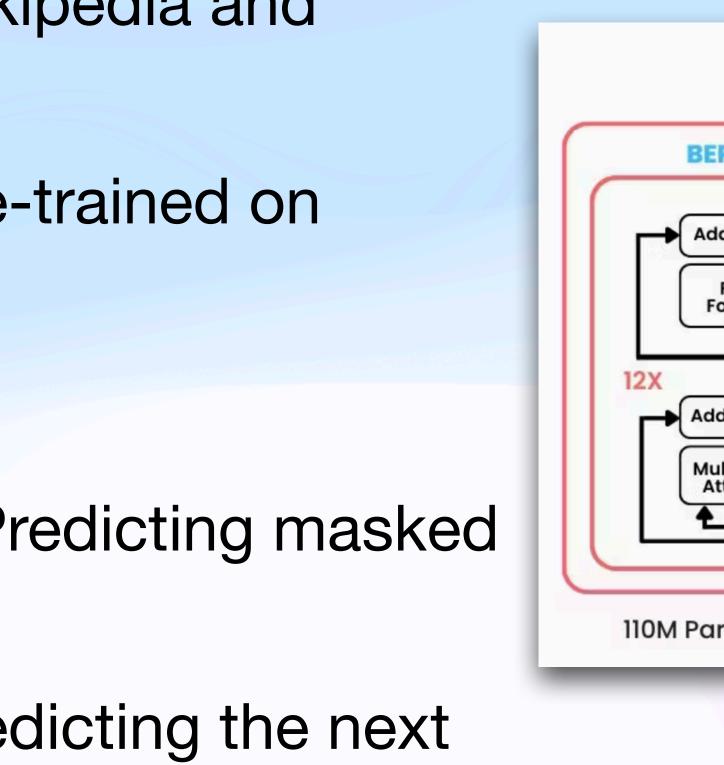
*until November, 2022

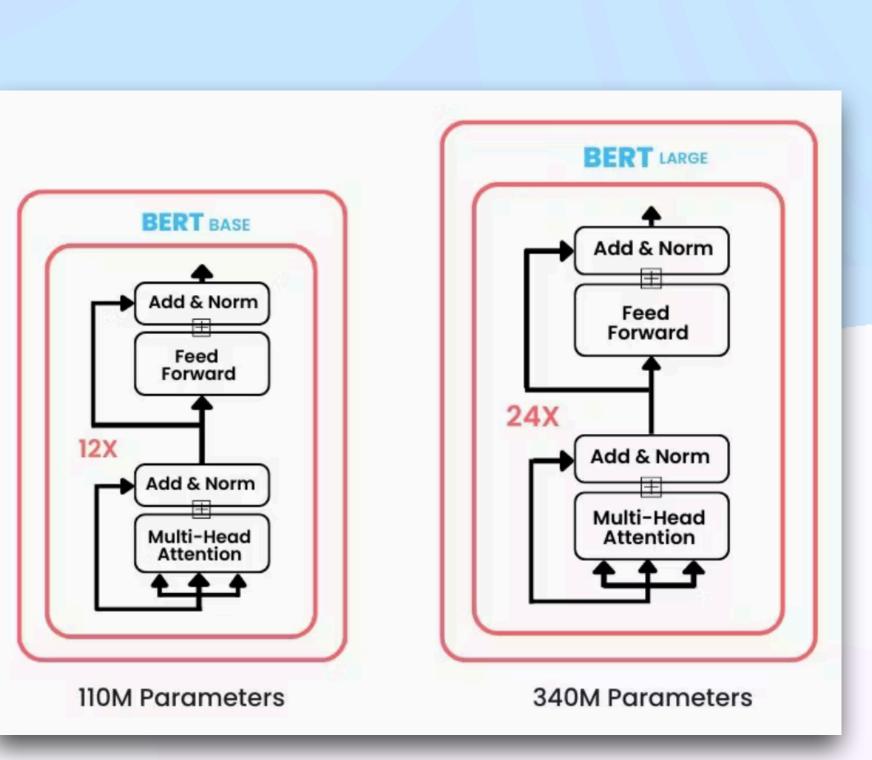
BERT and GPT



BERT and GPT

- **BERT**: Pre-trained on English Wikipedia and BooksCorpus.
- **GPT**: Pre-trained Transformer pre-trained on BookCorpus.
- Self-Supervision:
 - Masked Language Modeling: Predicting masked words based on **full** context.
 - Causal language modeling: Predicting the next word of the sentence based only on the past.





BERT (Base) and BERT (Large).

Where BERT Lacks?

- **BERT's Constraints:**
 - Better at understanding rather than generating text.
- GPT's Generative Edge:
 - Better at language generation through a causal language modeling.

 - Multi-task and also Few-shot learner.

GPT-2 and GPT-3 enhance performance on multiple downstream tasks.

Why not use GPT Directly in Biomedicine?

- Even GPT-3 struggles with biomedical tasks due to the Domain Shift.
- Previous Adaption Attempt:
 - DARE: Pre-trained on limited data (0.5M abstracts) for data augmentation in relation extraction.
 - But results were limited.
 - Task-Specific Adaptation(InstructGPT): GPT model adapted for unconventional downstream clinical tasks.

Introduction to BioGPT

What is **BioGPT**?

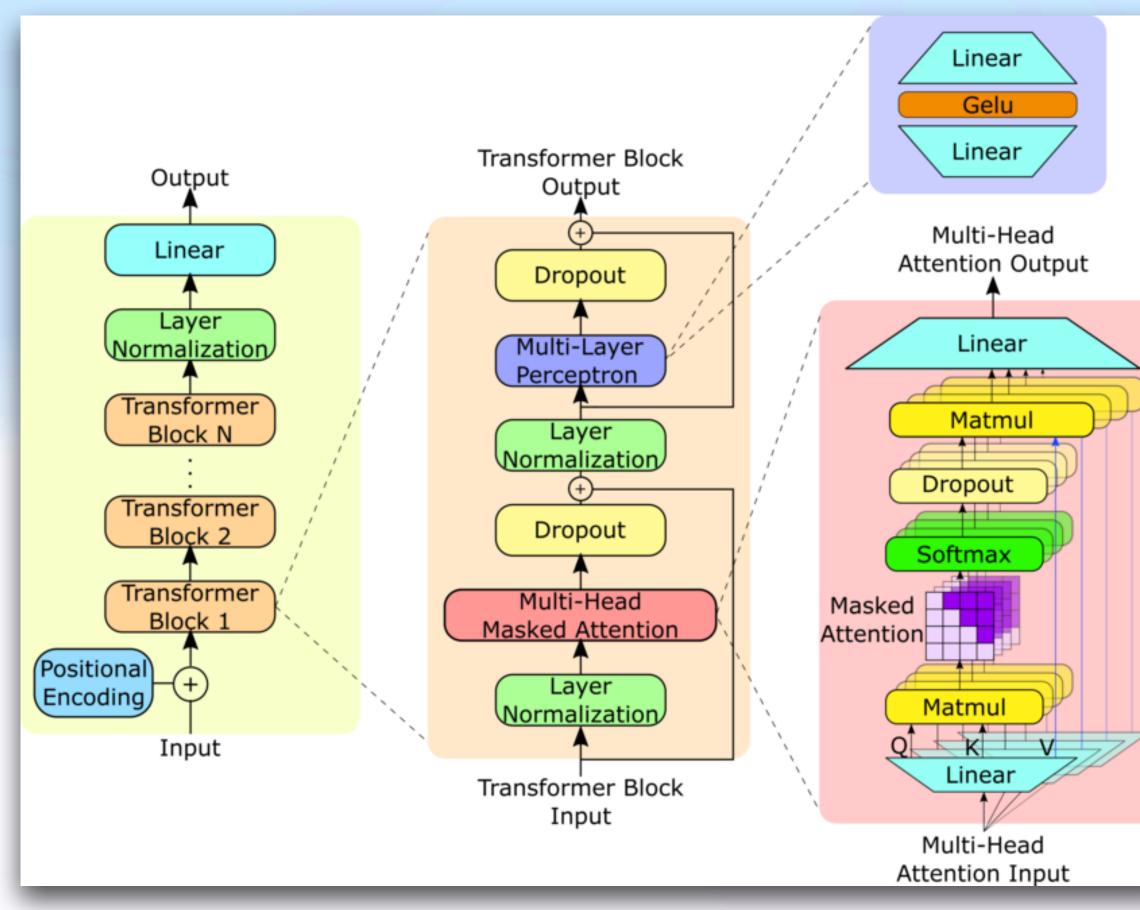
- **PubMed** abstracts.
- Key Features:
 - Combines generative modeling with biomedical relevance.
 - classification, and Text generation.

• A generative pre-trained Transformer for biomedical text, trained on **15M**

Can be used for Relation extraction, Question Answering, Document

Architecture

- Based on the GPT-2_{medium} architecture.
- Features a Transformer decoder structure.
- Key Components:
 - Multi-head attention mechanism.
 - Trained end-to-end for optimized performance across various tasks.



GPT-2 Architecture (Yang et al., 2023)



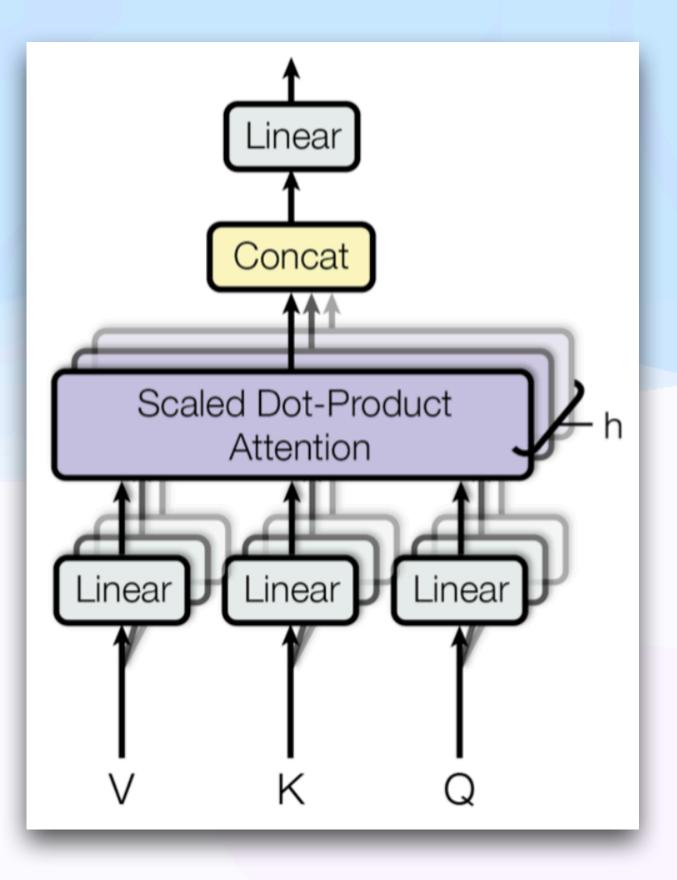
Multi-Head Attention Mechanism

- Runs multiple attention mechanisms in parallel.
- Outputs are concatenated and linearly transformed to match dimensions.

 $Multihead(Q, K, V) = Concat(head_1, head_2, ..., head_h)W,$

$$head_{i} = softmax\left(\frac{Q_{i}K_{i}^{\top}}{\sqrt{d}}\right)V_{i}$$

- Q, K, V: Queries, Keys, Values.
- W: Learnable parameter matrices. •
- *d*: Dimensionality scaling factor.



Multi-Head Attention (Vaswani et al., 2017)

Training and Fine-tuning

Training Dataset

- Pre-trained on 15M PubMed abstracts.
- Enables training effective language models with domain-specific knowledge.
- Bridge between general language understanding and biomedical text generation.
- **Key Characteristics:**
 - Diverse range of biomedical topics.
 - Continuously updated with newly published research.

Training

- Pre-trained using standard language modeling tasks.
- **Criteria:**
 - Minimize negative log-likelihood.

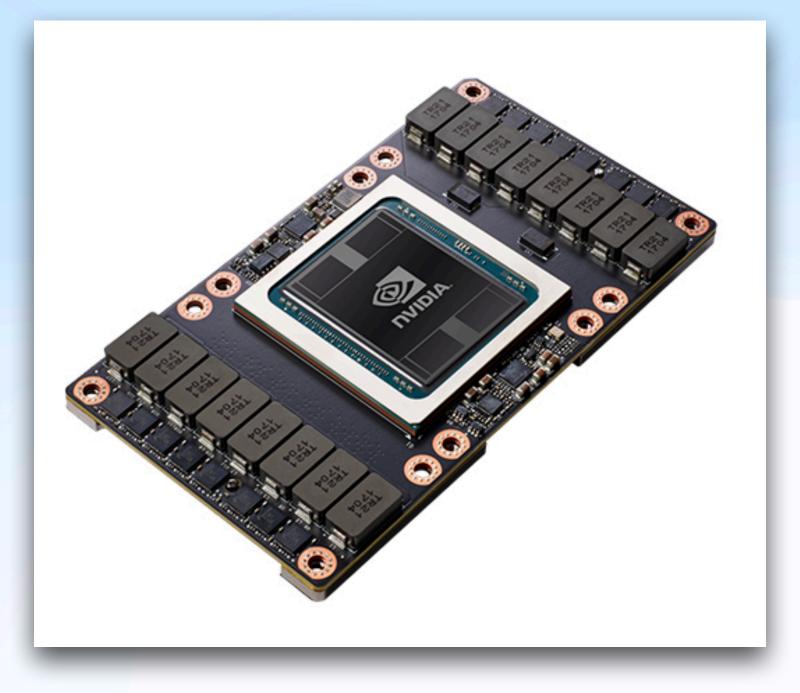
$$\min - \frac{1}{|D|} \sum_{i=1}^{|D|} \sum_{j=1}^{n_i} \log P(s_j \mid x_j)$$

- D: Dataset of sequences and s_i : token
- Effective batch size of **524,288 tokens**. •
- Adam optimizer with a learning rate schedule.
- Employed a warm-up phase(20,000 steps) to stabilize the training.

 $S_{j-1}, S_{j-2}, \dots, S_1$

Training Hardware

• Pre-trained on 8 NVIDIA V100 GPUs for 200,000 steps.



NVIDIA V100 FOR NVLINK, from NVIDIA website

Vocabulary Development

- Why Domain-Specific Vocabulary?
 - Language model's performance hinges on its vocabulary quality.
 - General vocabularies can complicate specialized biomedical terms.
- Vocabulary Creation Process

 - Byte Pair Encoding (BPE): Derives vocabulary directly from biomedical datasets. Final Vocabulary Size: 42,384 tokens (50,257 for GPT)
- **Advantages:**
 - Better understanding of biomedical terminology.
 - More precise and contextually relevant text generation.

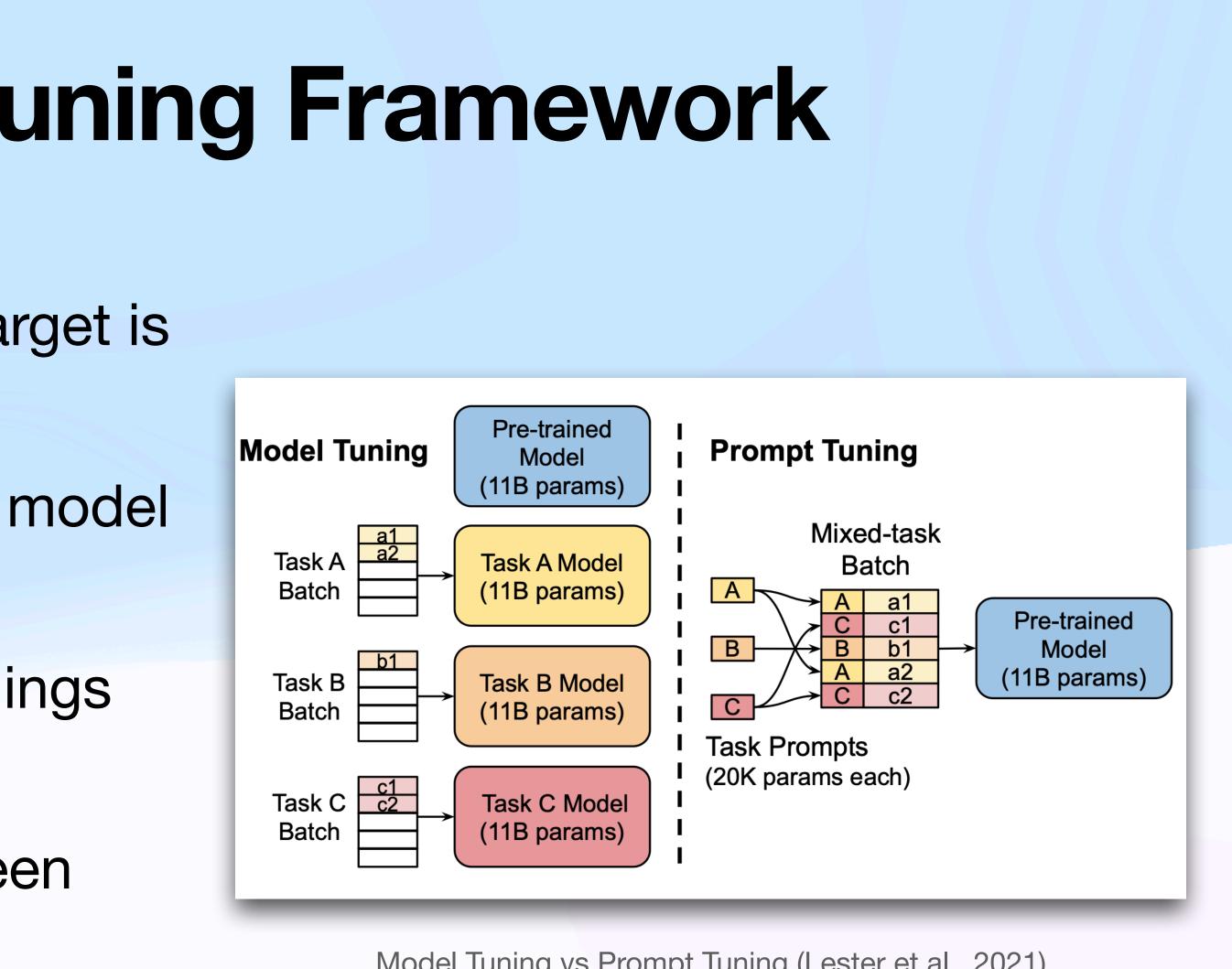
Fine-tuning Method Overview

- To adapt BioGPT for downstream tasks.
- **Key Adaptation:**
 - Convert labels into natural language sequences.
 - Maintains consistency with pre-training task format.
 - Avoids structured formats or special tokens for smoother semantics.



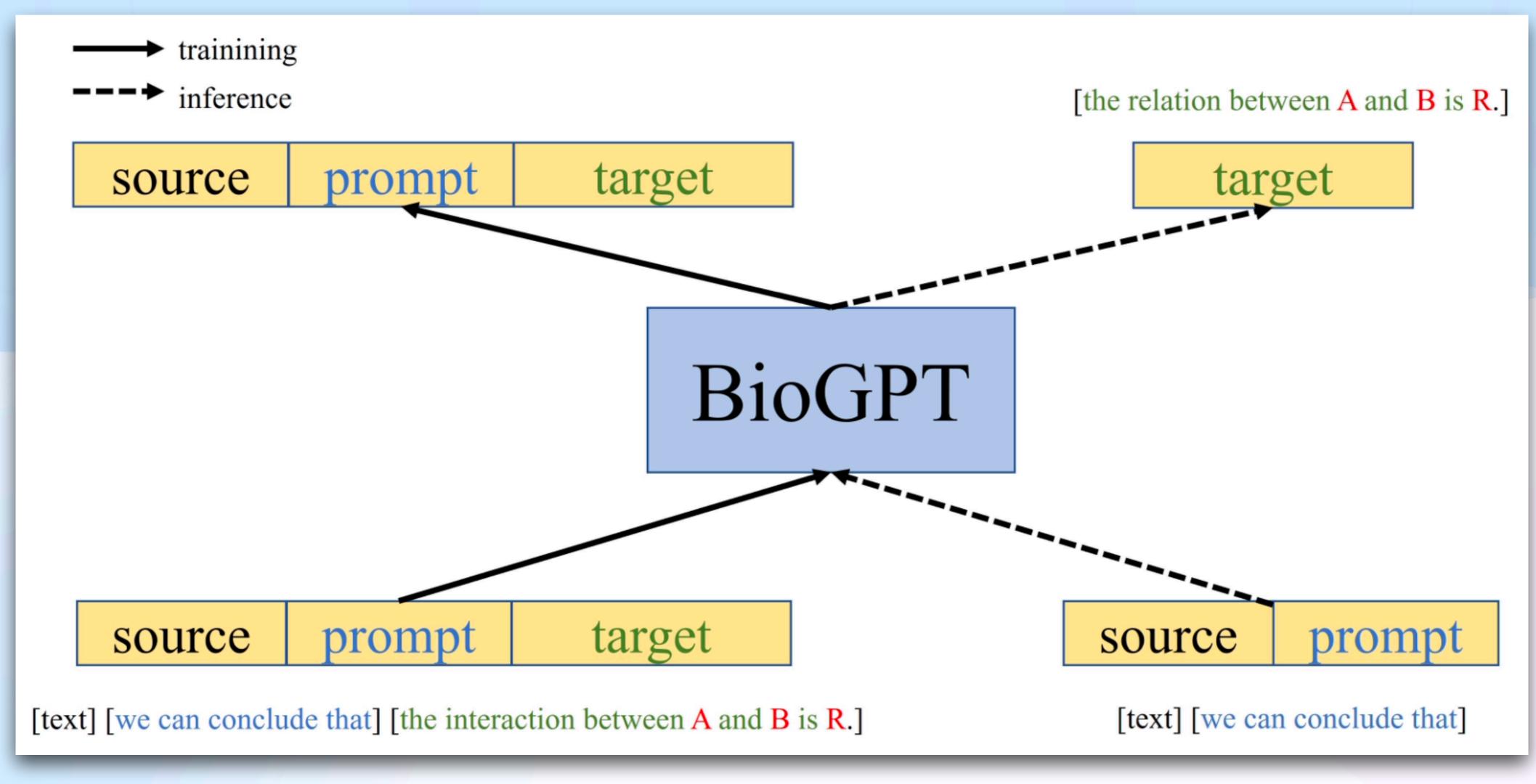
Prompt Based Fine-tuning Framework

- Simply concatenating source and target is ineffective.
- Workaround: prompts to guide the model in generating task-specific output.
- Soft Prompts: Continuous embeddings (virtual tokens) learned end-to-end.
 - Virtual tokens are placed between source and target sequences.



Model Tuning vs Prompt Tuning (Lester et al., 2021)

Prompt Based Fine-tuning Framework



Source: Paper

Downstream Tasks

- Relation Extraction:
 - Key for biomedicine.
- **Question Answering (QA):** •
 - Answering questions based on context.
 - Produce answer sequences, improving upon span prediction methods.
- **Document Classification:**
 - Classifying documents into predefined categories.
 - Leverages large pre-trained model for enhanced understanding and prediction.

Focuses on generating relational triplets directly from text without intermediate annotations.

Downstream Tasks

Task	Method	Dataset
Relation Extraction	GLRE, REBEL, seq2rel	KD-DTI, BC5CDR, DDI
Question Answering	QA-Net, LUKE, BERT, PubMedBERT, BioELECTRA, LinkBERT	PubMedQA, BioASQ
Document Classification	BERT, BlueBERT, SciBERT, SPECTER, PubMedBERT, BioELECTRA, LinkBERT	HoC, SciDocs

Summary of the downstream tasks for evaluation

BioGPT Variants

BioGPT

- 24-layer Transformer
- 347M parameters
- 15M PubMed abstracts
- Approx. 4B tokens

BioGPT_{large}

- 48- layer Transformer
- 1.57B parameters
- 15M PubMed abstracts + 6M PMC full paper
- Approx. 8B tokens

Results



1. Relation Extraction Task

needing intermediate steps.

Model	Precision	Recall	F1
GLRE(gt+pred)	34.82	18.29	23.99
GLRE(pred+pred)	23.00	4.88	8.05
GPT-2	43.92	32.55	37.39
REBEL	34.28	39.49	36.70
REBEL _{pt}	40.94	21.20	27.94
Seq2rel	43.5	37.5	40.2
BioGPT	49.44	41.28	44.98
BioGPT ⁺	49.52	43.25	46.17

Model	Precision	Recall	F1
Transformer + PubMedBERT-attn	25.35	24.14	24.19
GPT-2 _{medium}	30.53	27.87	28.45
REBEL	32.36	29.58	30.39
REBEL _{pt}	35.73	32.61	33.32
BioGPT	40.00	39.72	38.42

Results on **BC5CDR** chemical-disease-relation task

Results on **KD-DTI** drug-target-interaction task

Extracting relationships between entities(triplets) in a single pass, without

Model	Precision	Recall	F1
GPT-2 _{medium}	23.39	31.93	24.68
REBEL	35.36	28.64	28.27
REBEL _{pt}	46.59	39.60	40.56
BioGPT	41.70	44.75	40.76

Results on **DDI** drug-drug-interaction task



2. Question Answering

- Goal: Answer questions using reference context.
- Labels: Yes, No, Maybe.

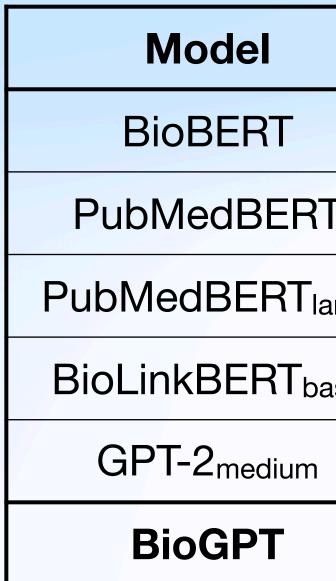
Model
PubMedBER
BioELECTRa
BioLinkBERT
BioLinkBERT
BioGPT

Results on **PubMedQA** question answering task

	Accuracy
٦T	55.8
a	64.2
T _{base}	70.2
T large	72.2
	78.2

3. Document Classification

- Goal: Classify document type based on text.
- Target Sequence Format: The type of this document is 'label'.



Results on HoC document classification task

	F1
	81.54
Т	82.32
arge	82.70
ase	84.35
	81.84
	85.12

Results Summary

Relation Extraction:

- Drug-Target Interaction (KD-DTI)
- Chemical-Disease Interaction (BC5CDR)
- Drug-Drug Interaction (DDI)
- Up to 4% improvement over all methods

Question Answering:

- PubMedQA
- 6.0% improvement over previous best

Document Classification:

- HoC
- 3.28% improvement over previous ones



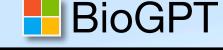
BioGPT in Action





Drug-Target Interaction Extraction

The Janus family kinases (Jaks), Jak1, Jak2, Jak3, and Tyk2, form one subgroup of the non-receptor protein tyrosine kinases. They are involved in cell growth, survival, development, and differentiation of a variety of cells but are critically important for immune cells and hematopoietic cells. Data from experimental mice and clinical observations have unraveled multiple signaling events mediated by Jak in innate and adaptive immunity. Deficiency of Jak3 or Tyk2 results in defined clinical disorders, which are also evident in mouse models. A striking phenotype associated with inactivating Jak3 mutations is severe combined immunodeficiency syndrome, whereas mutation of Tyk2 results in another primary immunodeficiency termed autosomal recessive hyperimmunoglobulin E syndrome. In contrast, complete deletion of Jak1 or Jak2 in the mouse are not compatible with life and, unsurprisingly, do not have counterparts in human disease. However, activating mutations of each of the Jaks are found in association with malignant transformation, the most common being gain-of- function mutations of Jak2 in polycythemia vera and other myeloproliferative disorders. Our existing knowledge on Jak signaling pathways and fundamental work on their biochemical structure and intracellular interactions allow us to develop new strategies for controlling autoimmune diseases or malignancies by developing selective Jak inhibitors, which are now coming into clinical use. Despite the fact that Jaks were discovered only a little more than a decade ago, at the time of writing there are 20 clinical trials underway testing the safety and efficacy of Jak inhibitors.



• **Output:** the interaction between pnus156804 and janus kinase 3 (jak 3) is inhibitor.

 Structured Output: cpnus156804, janus kinase 3(jak 3), inhibator >

Task: Find <drug, target, interaction> triplet given the document



Drug-Drug Interaction Extraction

BioGPT

An inhibitor of CYP2C8 (such as gemfibrozil) may increase the AUC of rosiglitazone and an inducer of CYP2C8 (such as rifampin) may decrease the AUC of rosiglitazone. Therefore, if an inhibitor or an inducer of CYP2C8 is started or stopped during treatment with rosiglitazone, changes in diabetes treatment may be needed based upon clinical response.

- **Output:** the interaction between gemfibrozil and rosiglitazone is mechanism; the interaction between rifampin and rosiglitazone is mechanism.
- Structured output: (gemfibrozil, rosiglitazone, mechanism), (rifampin, rosiglitazone, mechanism)

Task : Find <drug, drug, interaction> triplet given the document

Question Answering

Question: Do some US states have higher / lower injury mortality rates than others?

Context: this article examines the hypothesis that the six u.s. states with the highest rates of road traffic deaths (group 1 states) also had above-average rates of other forms of injury such as falling, poisoning, drowning, fire, suffocation, homicide, and suicide, and also for the retail trade and construction industries. the converse, second hypothesis, for the six states with the lowest rates of road traffic deaths (group 2 states) is also examined. data for these 12 states for the period 1983 to 1995 included nine categories of unintentional and four categories of intentional injury, seventy-four percent of the group 1 states conformed to the first hypothesis, and 85% of the group 2 states conformed to the second hypothesis, answer: group 1 states are likely to exhibit above-average rates for most other categories of injury death, whereas group 2 states are even more likely to exhibit below-average rates for most other categories of injury death.

Ground truth: Yes

BioGPT: the answer to the question given the context is **Yes.**



Zero-shot QA

Context: [tl;dr: Some background introduction] The aim of this work is to assess a previously described slit-lamp biomicroscopy-based method (SLBM) for measuring pupil diameter and compare it to Colvard infrared pupillometry (CIP). Two examiners performed three repeated measurements with each instrument in 40 healthy eyes. We determined the agreement of SLBM and CIP, intraobserver and interobserver repeatabilities, and interobserver concordance (kappa) and SLBM ability for detecting pupil sizes over 6.0 mm. The mean (±standard deviation [SD]) pupil diameter was 5.81 ± 0.70 mm with SLBM and 6.26 ± 0.68 mm with CIP (p=0.01) averaging both examiner's results. Mean differences between the SLBM and CIP were -0.60 mm and -0.30 mm for each examiner using the average of the three readings (p=0.02), and they were very similar using the first reading. Intraobserver reproducibility: the width of the 95% LoA ranged from 1.79 to 2.30 mm. The ICCs were 0.97 and 0.92 for SLBM, and 0.96 and 0.90 for CIP. Interobserver reproducibility, the width of the LOA ranged from 1.82 to 2.09 mm. Kappa statistics were 0.39 and 0.49 for the first and mean SLBM readings, respectively, and 0.45 for both the first and mean CIP readings. Sensitivity and specificity of SLBM for detection of pupils larger than 6 mm ranged from 55.56% to 73.68% and from 76.19% to 95.45%, respectively. The best tradeoff between sensitivity and specificity ranged from 5.4 mm to 6.2 mm.

Ground truth: No

BioGPT: The answer is that SLBM is not a good alternative to CIP for measuring pupil size.

Question: Can we measure mesopic pupil size with the cobalt blue light slit-lamp biomicroscopy method?



Text Generation Common Words

Input	Model	Text
Bicalutamide	GPT-2	Bicalutamide, a synthetic bicalutar elegans cells. This effect was not o line with our previous work which s but not in the presence of synthetic
	BioGPT	Bicalutamide (Casodex) is an andro castration-resistant prostate cancer docetaxel chemotherapy, as well as progressed on or are ineligible for castration-sensitive prostate cancer docetaxel chemotherapy.

GPT-2 vs BioGPT when given relatively common names as input

amide derivative, was found to increase the cellular proliferation of C. observed with the other synthetic bicalutamide derivatives. This is in showed that bicalutamide increases the proliferation of C. elegans cells ic bicalutamide.

ogen receptor (AR) antagonist approved for the treatment of metastatic er (mCRPC) in patients who have progressed on or are ineligible for s for the treatment of early-stage prostate cancer in men who have not c docetaxel chemotherapy, as well as for the treatment of metastatic er (mCSPC) in men who have not progressed on or are ineligible for



Text Generation Uncommon Words I

Xylazine	GPT-2	Xylazine, the "active ingredient" in those of former University of Califor having a bath filled with it.
	BioGPT	Xylazine is an alpha 2-adrenoceptor medicine for many years, but its effects the dog, and its effects on the centr despite the fact that xylazine has b more than 30 years.
Psoralen	GPT-2	Psoralen, the city of the spiritualis way of salvation, and to draw him b his fellow-man.
	BioGPT	Psoralen photochemotherapy (PUV by the risk of skin cancer, particul which is the most common site of s

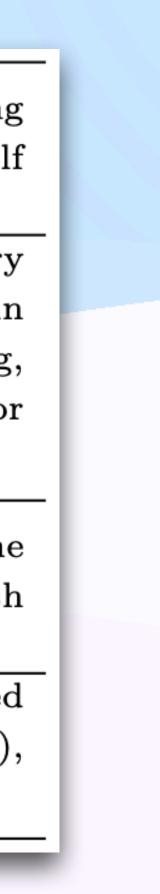
GPT-2 vs BioGPT when given somewhat uncommon names as input

" "bath salts" (see above) has been linked to numerous deaths, including ornia, Berkeley, professor Andrew Anglin, who posted a video of himself

or agonist that has been used as a sedative and analgesic in veterinary fects on the cardiovascular system have not been extensively studied in ral nervous system (CNS) have not been well characterized in the dog, been widely used as a sedative and analgesic in veterinary medicine for

st, and had been given to him by God, who wished to show him the back to the true life of Christ, and to bring him into communion with

VA) is a well-established treatment for psoriasis, but its use is limited larly squamous cell carcinoma (SCC) of the head and neck (H & N), skin cancer in the United States (US)



Text Generation Uncommon Words II

GPT-2	CP-673451) and 4-5-6-3-1-2-3-4-6-7
	31-32-33-34-35-36-37-38-39-40-41-42-
BioGPT	CP-673451 is a potent, selective, a human cathepsin G (CatG) with in models of inflammation and in a m instillation of lipopolysaccharide (Li lung injury (ALI) in which neutroph
GPT-2	BIIB-021A-2-1.pdf. Wright, C. (200 L.E.R. Wright, S.R. Warrick, and J. T. Klare (New York: Oxford University)
BioGPT	BIIB-021 is a novel, orally active, n lasting anti-inflammatory activity in of adjuvant-induced arthritis (AIA), collagen-induced arthritis (CIA), an is induced by immunization with bo
	GPT-2

7-8-9-10-11-12-13-14-15-16-17-18-19-20-21-22-23-24-25-26-27-28-29-30- $2 - 43 - 44 - 45 - 46 - 47 - 48 - 49 - 50 \dots$

and orally active inhibitor of human neutrophil elastase (HNE) and n vitro and in vivo anti-inflammatory activity in a variety of animal nodel of acute lung injury (ALI) in the rat induced by intratracheal LPS) and tumor necrosis factor-alpha (TNF-alpha), a model of acute hils play an important role.

02). "The War on Terror: A Legal Theory of Terrorist Attacks," in .H. Rothman (Eds.), The War on Terror: Essays in Honor of Michael rsity Press), 143-57.

non-peptide bradykinin B2 receptor antagonist with potent and longn animal models of acute and chronic inflammation and in a rat model , an animal model of rheumatoid arthritis (RA), and in a rat model of n animal model of collagen-induced arthritis (CIA), in which arthritis ovine type II collagen (CII).



Text Generation Keywords of interest

Input	Model	Text
COVID-19	GPT-2	COVID-19-0067-7 http://www.pinterest.com/pin/7117 http://www.pinterest.com/pin/7117
	BioGPT	COVID-19 is a global pandemic cau 2), the causative agent of coronavi- countries and territories, including Kingdom (UK), and the United Sta- confirmed cases and more than 800,
SARS-CoV-2	GPT-2	SARS-CoV-2-CKM-T1-1/pdf/T1/S. topics/programs/tox/tox-programs.
	BioGPT	SARS-CoV-2 is the causative agent infected more than 390,000 people v

GPT-2 vs BioGPT when manually given keywords of interest (COVID-19 related terms)

Hip-hop-romance-detective-franchise-unveiled-in-coco-houston 78278499149422/ Black Widow: World War II's Greatest Widow? 77842668650112/

used by severe acute respiratory syndrome coronavirus 2 (SARS-CoVvirus disease 2019 (COVID-19), which has spread to more than 200 the United States (US), Canada, Australia, New Zealand, the United ates of America (USA), as of March 11, 2020, with more than 800,000 0,000 deaths.

SARS-CoV-2-CKM-T1-1.pdf Cancer http://www.cancer.gov/cancer .html Other Research http://www.cdc.gov/cancer/cancer/index.html t of COVID-19, a severe acute respiratory syndrome (SARS) that has worldwide and killed more than 250,000 people.



Scaling to Larger Size **BioGPT**_{large}

- Developed on GPT-2 XL architecture, 1.5B parameters.

Task	
BC5CDR	
KD-DTI	
DDI	
PubMedQA	Α
НоС	

BioGPT-Large fine-tuned on downstream tasks



Fine-tuned and evaluated for enhanced performance on downstream tasks.

Performance	
50.12	
38.39	
44.89	
81.0	
84.40	

Conclusion

- Built on the GPT-2 backbone, pre-trained on 15M PubMed abstracts.
- **Pioneer** to adapt GPT effectively in Biomedicine domain.
- **Outperforms** GPT-2 in biomedical text generation.
- State-of-the-Art results on:
 - S relation extraction tasks.
 - I question answering task.
- Larger-scale BioGPT model on expanded biomedical datasets.

References

- https://arxiv.org/pdf/2210.10341
- https://pubmed.ncbi.nlm.nih.gov/
- https://www.semanticscholar.org/
- https://arxiv.org/pdf/2409.00133
- https://vitalflux.com/bert-vs-gpt-differences-real-life-examples/
- <u>shown fig1 373352176</u>
- https://arxiv.org/pdf/1706.03762
- https://www.nvidia.com/de-de/data-center/

https://www.researchgate.net/figure/GPT-2-model-architecture-The-GPT-2-model-contains-N-Transformer-decoder-blocks-as-

Any Questions? Thank you for listening!