



Introduction to LLMs with a Focus on Biomedical Data Science

Presenter: Shubo Tian



National Library of Medicine
National Center for Biotechnology Information

Learning Objective

- Understanding what is language model, type of language models, and existing major language models and their applications in general and in biomedicine.

What is Language Model

Given a sequence of words/tokens:

$$w_1, w_2, \dots, w_n$$

a model that computes either of the following probabilities:

$$P(w_1, w_2, \dots, w_n) \text{ or}$$

$$P(w_n | w_1, w_2, \dots, w_{n-1})$$

is called a language model.

$$P(w_1, w_2, \dots, w_n) = \prod_{i=1}^n P(w_i | w_1, \dots, w_{i-1})$$

$P(\text{"P53 is a tumor suppressor gene."})$

$= P(\text{"P53"}) \cdot$

$P(\text{"is"} | \text{"P53"}) \cdot$

$P(\text{"a"} | \text{"P53 is"}) \cdot$

$P(\text{"tumor"} | \text{"P53 is a"}) \cdot$

$P(\text{"suppressor"} | \text{"P53 is a tumor"}) \cdot$

$P(\text{"gene"} | \text{"P53 is a tumor suppressor"}) \cdot$

$P(\text{"."} | \text{"P53 is a tumor suppressor gene"})$

Language Modeling

Word count modeling (n-gram)

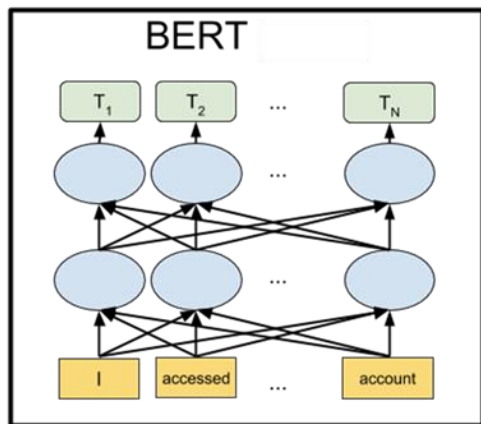
$$P(w_n | w_1, w_2, \dots, w_{n-1}) = \frac{\text{count}(w_1, w_2, \dots, w_n)}{\text{count}(w_1, w_2, \dots, w_{n-1})}$$

Neural network modeling

$$\begin{aligned} P(w_n | w_1, w_2, \dots, w_{n-1}) &= P(w_n | h) \\ &= \frac{\exp(h^T \text{emb}(w_n))}{\sum_{w' \in \text{Vocab}} \exp(h^T \text{emb}(w'))} \end{aligned}$$

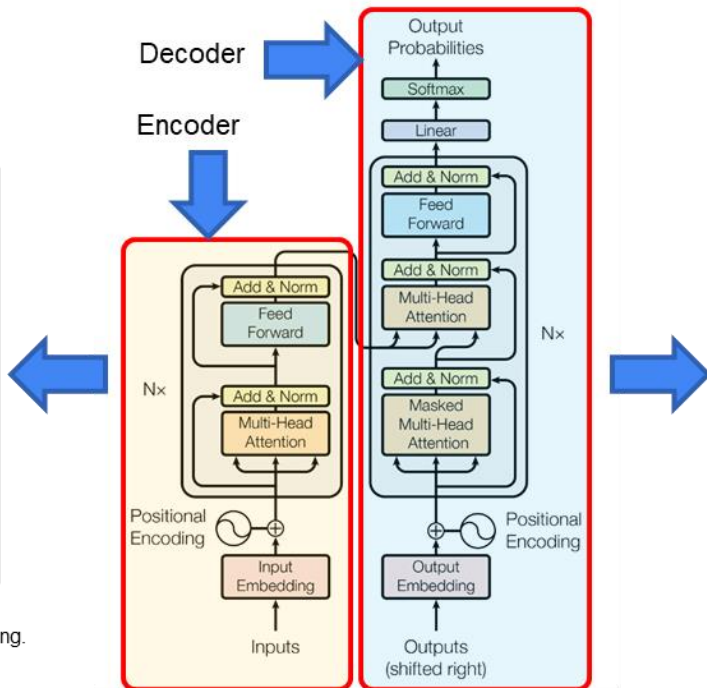
$$h = \text{Encoder}(w_1, w_2, \dots, w_{n-1})$$

Transformer and LLMs

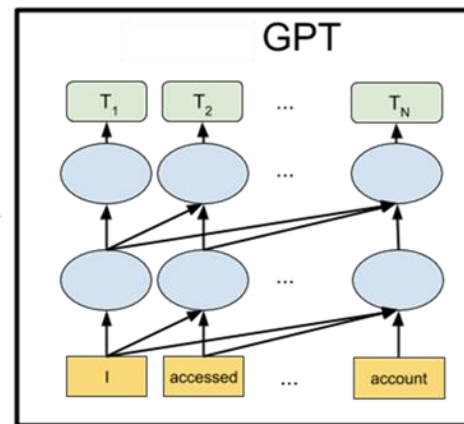


Devlin, J. *et al.* (2019) BERT: Pre-training of Deep Bidirectional Transformers for Language Understanding. *arXiv:1810.04805 [cs]*.

<https://ai.googleblog.com/2018/11/open-sourcing-bert-state-of-art-pre.html>



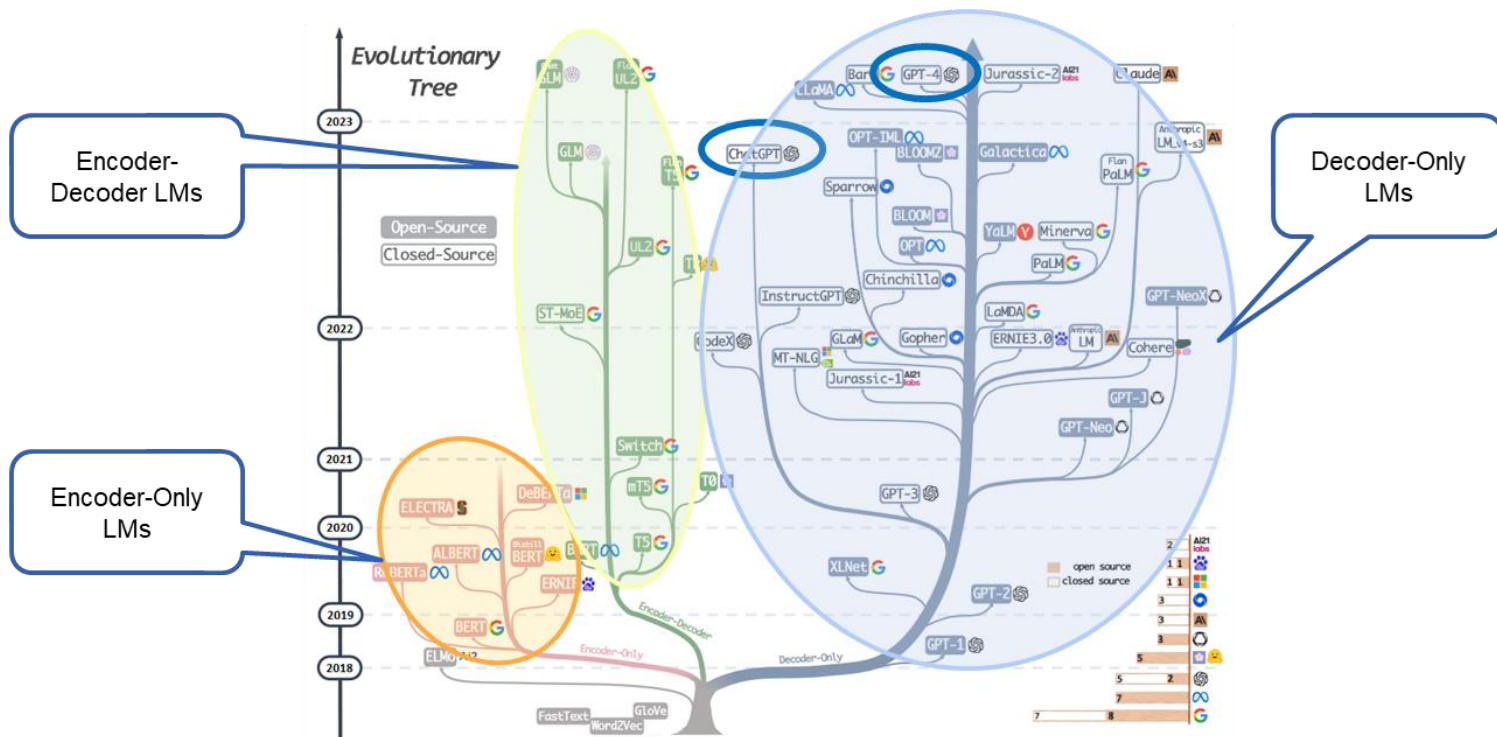
Vaswani, A. *et al.* (2017) Attention Is All You Need. *arXiv:1706.03762 [cs]*.



Radford, A. *et al.* Improving Language Understanding by Generative Pre-Training.

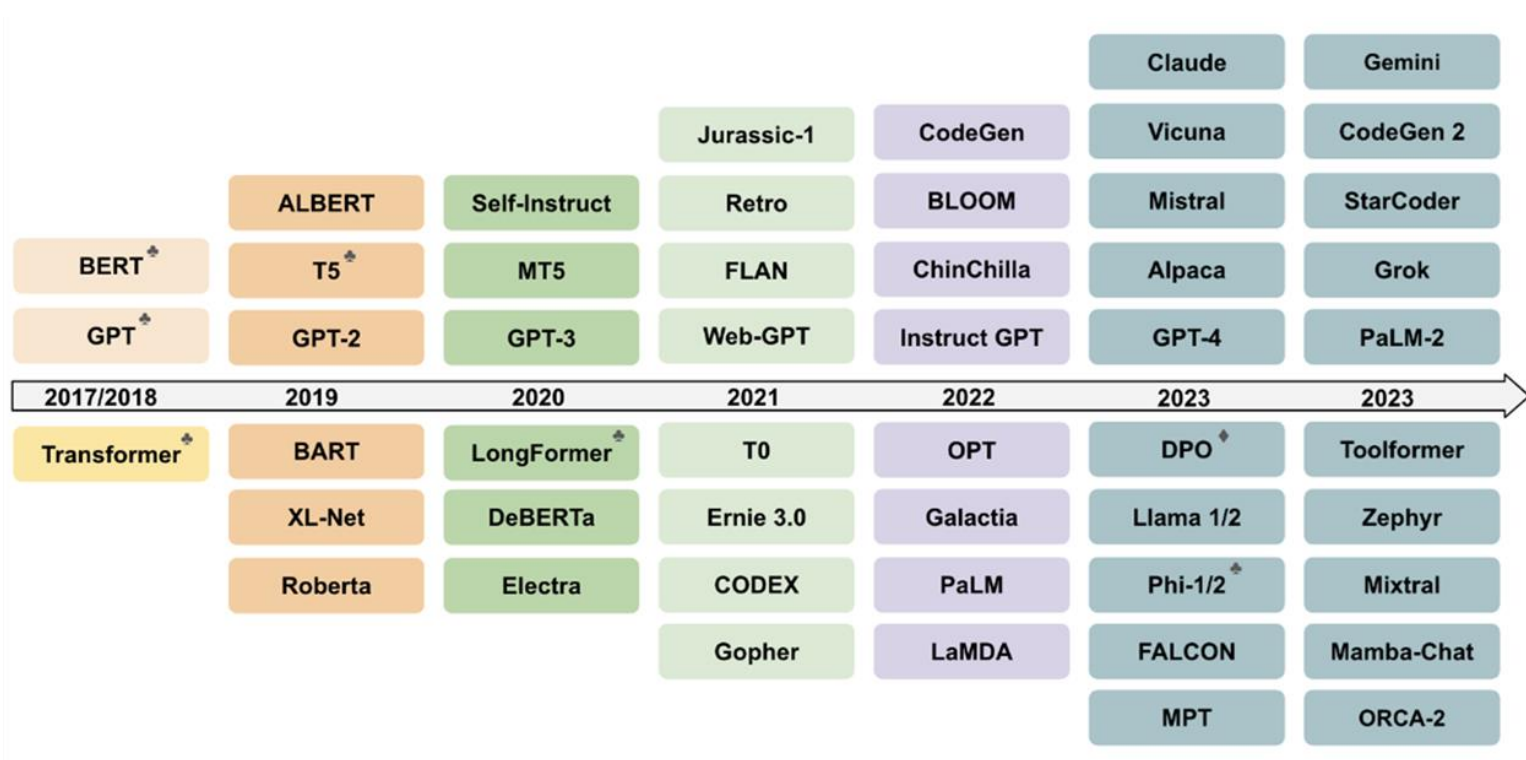
<https://ai.googleblog.com/2018/11/open-sourcing-bert-state-of-art-pre.html>

Type of LLMs

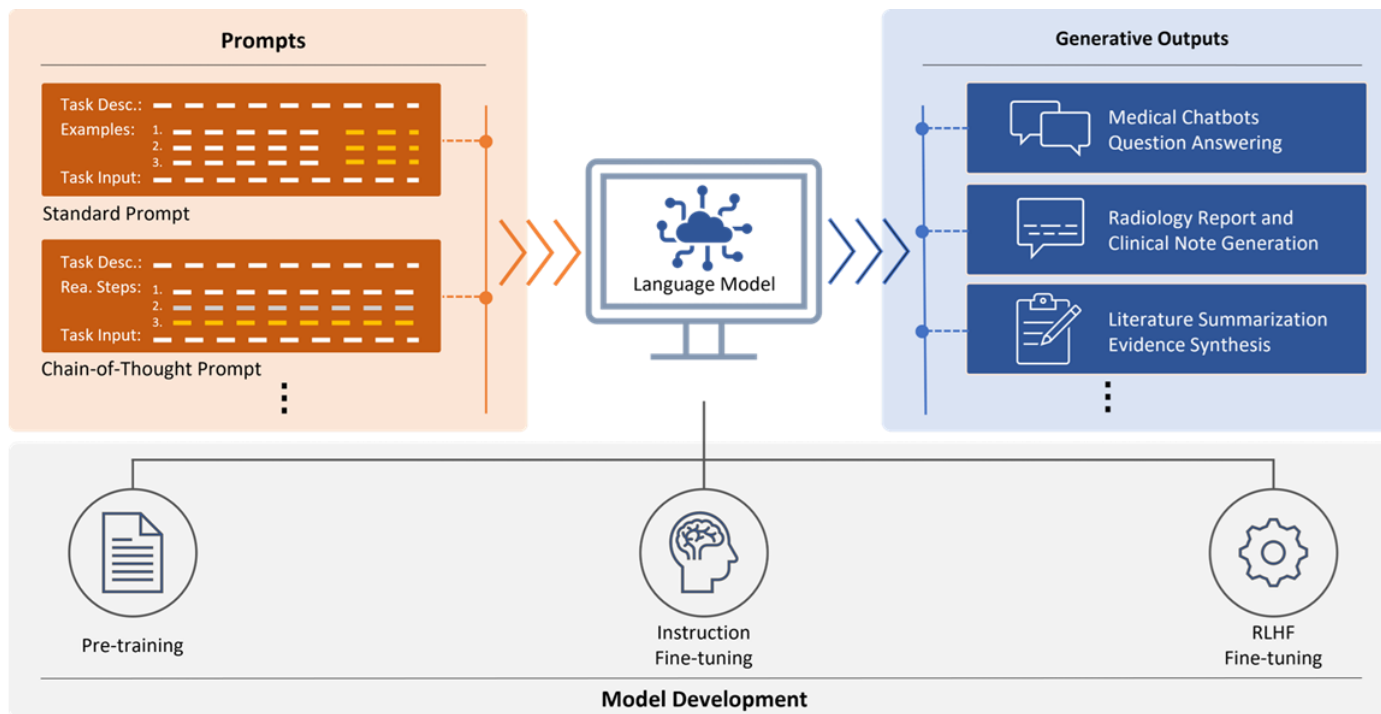


Yang, J., Jin, H., Tang, R., Han, X., Feng, Q., Jiang, H., Yin, B. and Hu, X. (2023) Harnessing the Power of LLMs in Practice: A Survey on ChatGPT and Beyond. 10.48550/arXiv.2304.13712.

Timeline of Representative LLMs

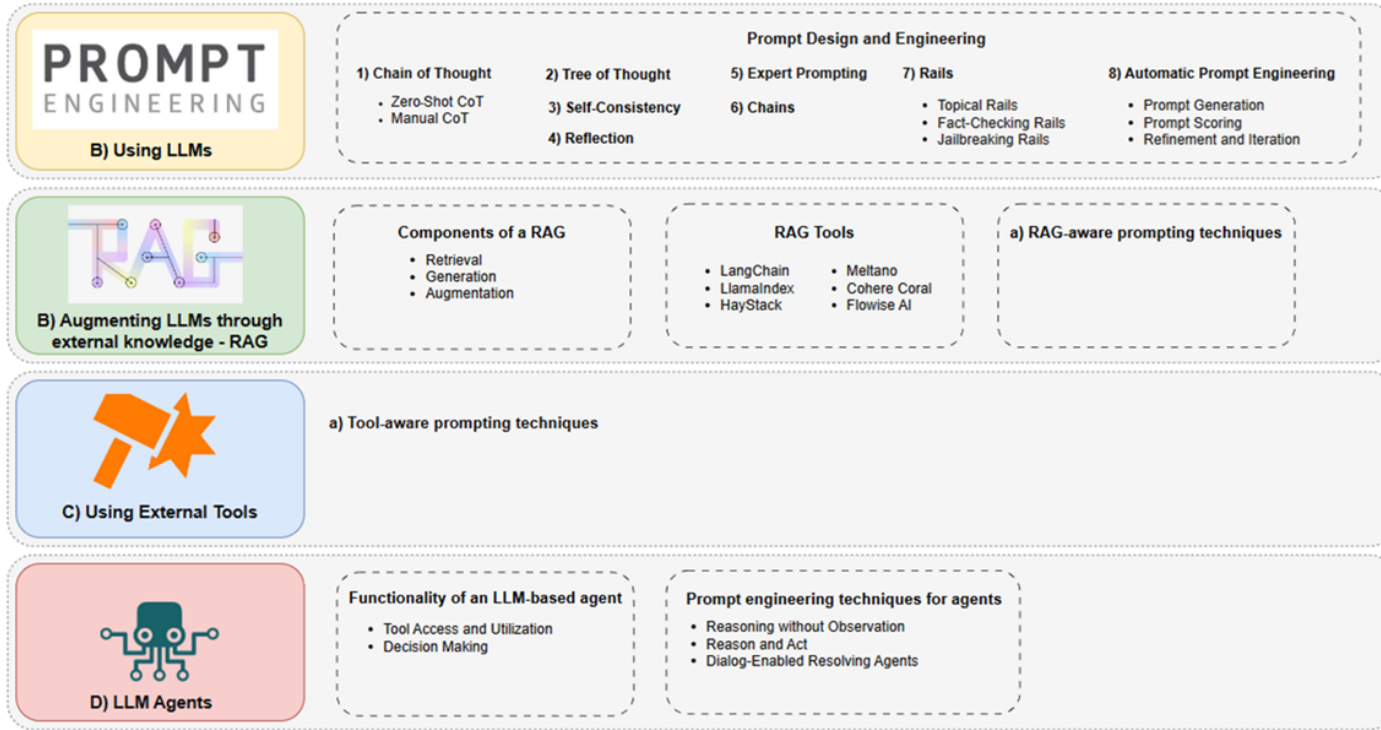


The Paradigm of LLMs



Tian, S., Jin, Q., Yeganova, L., Lai, P.-T., Zhu, Q., Chen, X., Yang, Y., Chen, Q., Kim, W., Comeau, D.C., et al. (2024) Opportunities and challenges for ChatGPT and large language models in biomedicine and health. *Briefings in Bioinformatics*, 25, bbad493.

How LLMs are Used

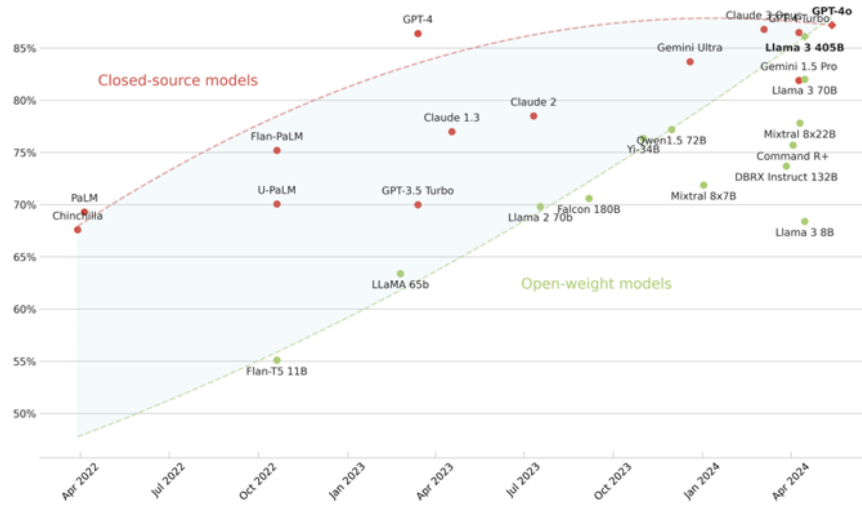


Major LLMs and Their Performance

Closed-source vs. open-weight models

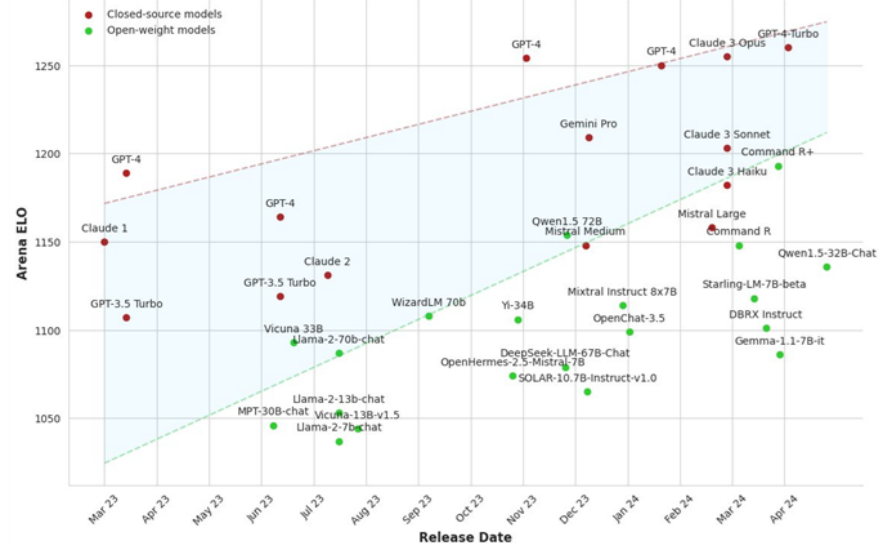
Llama 3 405B from Meta closes the gap between closed-source and open-weight models.

MMLU (5-shot)



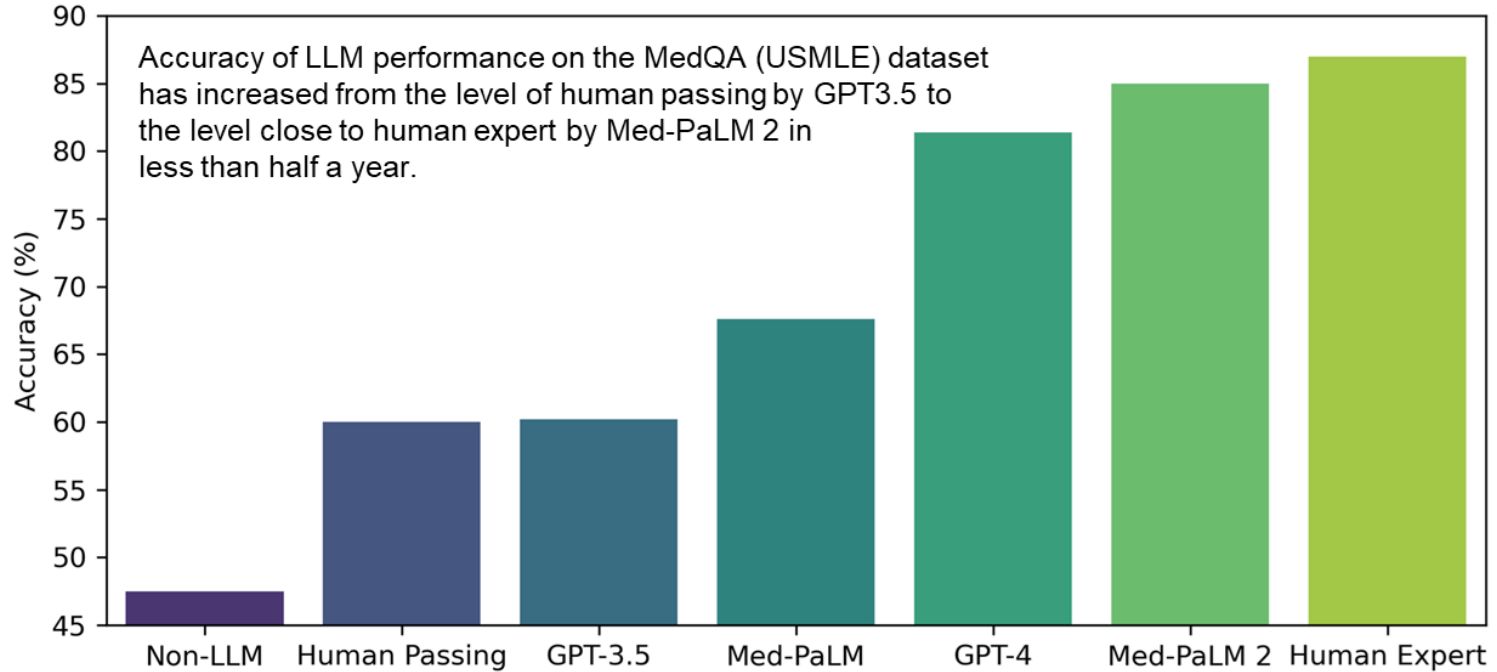
<https://twitter.com/maximelabonne/status/1793947985430057288>
<https://crfm.stanford.edu/helm/mmlu/latest/#/leaderboard>

Closed-source vs. Open-weight models (Arena ELO)



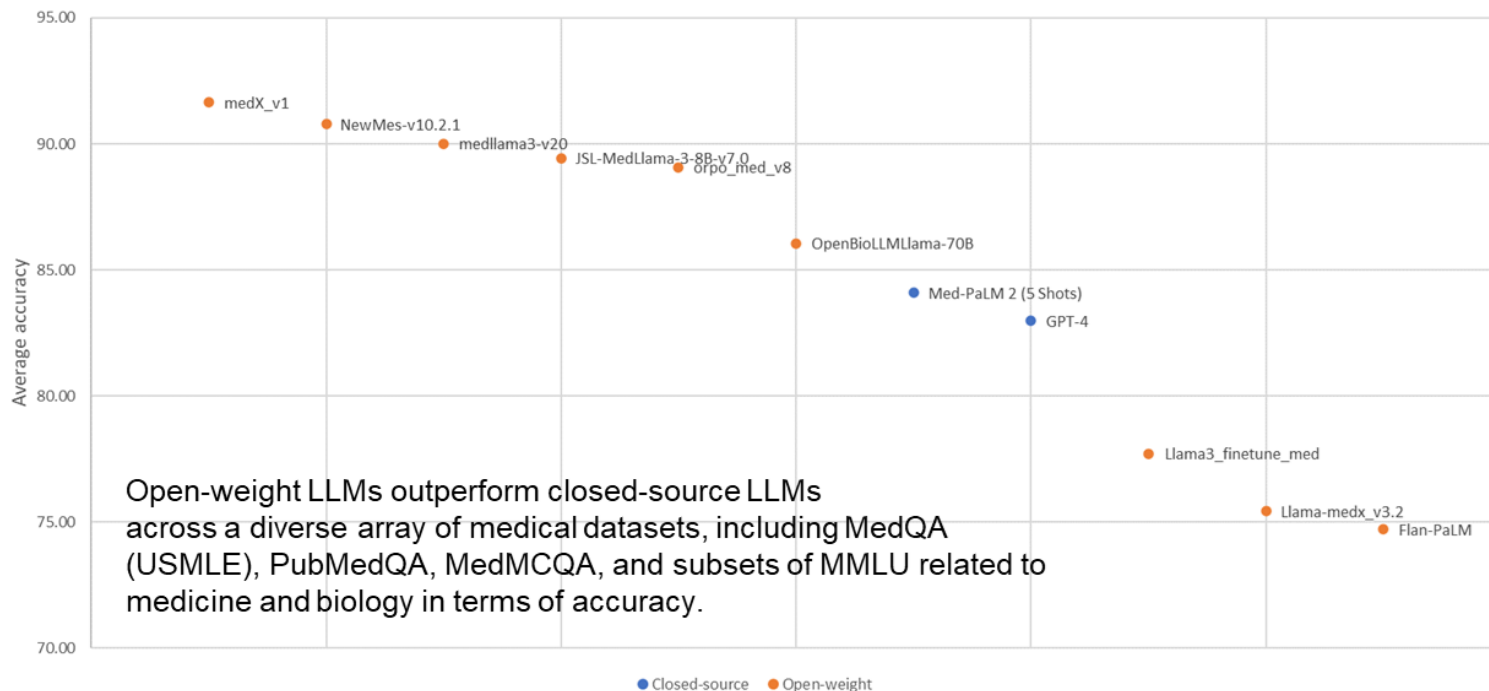
<https://twitter.com/maximelabonne/status/1779801605702836454>
<https://chat.lmsys.org/?leaderboard>

LLMs in Biomedicine



Tian, S., Jin, Q., Yeganova, L., Lai, P.-T., Zhu, Q., Chen, X., Yang, Y., Chen, Q., Kim, W., Comeau, D.C., et al. (2024) Opportunities and challenges for ChatGPT and large language models in biomedicine and health. *Briefings in Bioinformatics*, 25, bbad493.

Major LLMs in Biomedicine and Their Performance



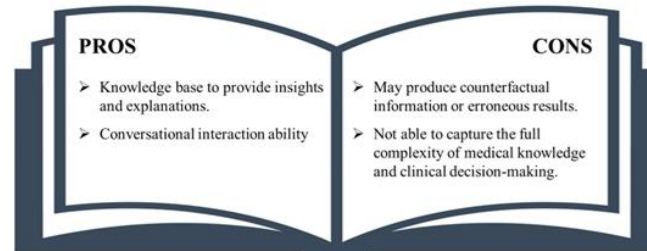
https://huggingface.co/spaces/openlifescienceai/open_medical_llm_leaderboard

Applications of LLMs in Biomedicine



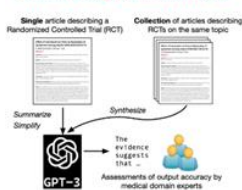
Macular corneal dystrophy in a Chinese family related with novel mutations of **CHST6**.

PURPOSE: To identify mutations in the **cathepsin B autoantigenase gene (CHST6)** for a Chinese family with **macular corneal dystrophy (MCD)** and to investigate the histopathological changes in the affected cornea. **METHODS:** A corneal button of the proband was obtained by penetrating keratoplasty. The half button and ultrathin sections from the other half button were examined with special stains under a light microscope (LM) and an electron microscope (EM) separately. Genomic DNA was extracted from peripheral blood of 11 family members, and the coding region of **CHST6** was amplified by the polymerase chain reaction (PCR) method. The PCR products were analyzed by **direct sequencing** and restriction enzyme digestion. **RESULTS:** The positive reaction to colloidal **iron** stain (intracellular blue accumulations in the stroma) was detected under light microscopy. Transmission electron microscopy revealed the enlargement of smooth endoplasmic reticulum and the presence of intracytoplasmic vacuoles. The compound heterozygous mutations, **C899C>G** and **C1027T>C**, were identified in exon 3 of **CHST6** in three patients. The two transmissions resulted in the substitution of a **high codon for glutamine at codon 294 (g1029900)** and a missense mutation at **codon 354, lysine to histidine (g29258)**. The six unaffected family individuals carried alternative heterozygous mutations. These two mutations were not detected in any of the 100 control subjects. **CONCLUSIONS:** Those novel compound heterozygous mutations were thought to contribute to the loss of function, which induced the abnormal metabolism of **keratan sulfate (KS)** that deposited in the corneal stroma. It could be proved by the observation of a positive stain reaction and the enlarged **keratan** fibers as well as hyperplastic fibroblasts under microscopes.

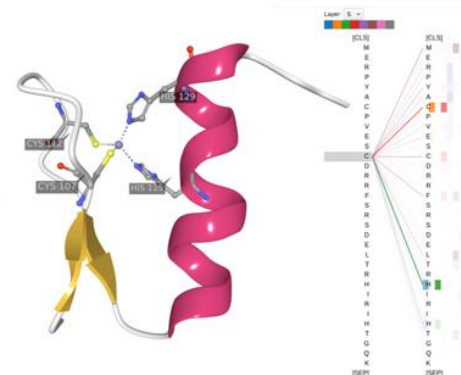
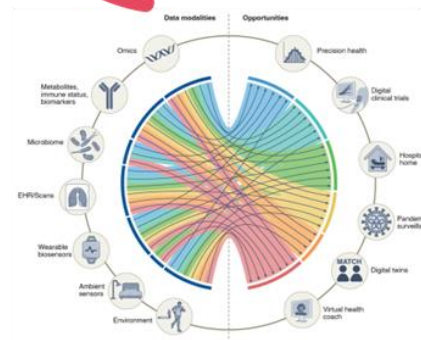
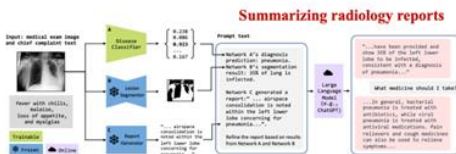


The ChatGPT/LLMs Learning Avenues

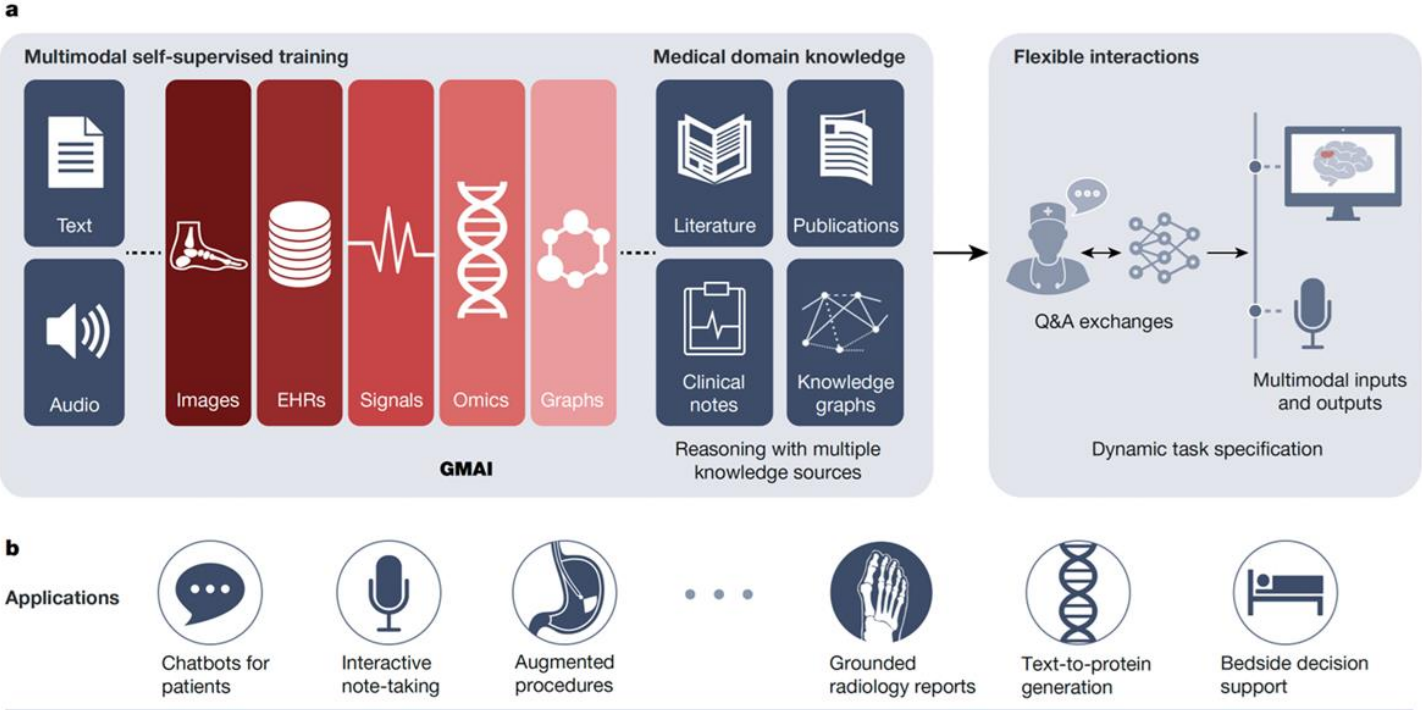
Summarizing medical evidence



ChatGPT: the future of discharge summaries?
 Sejan B Patel • Kyle Lam
 Open Access • Published: February 06, 2023 • DOI: [https://doi.org/10.1016/S2589-7500\(23\)00021-3](https://doi.org/10.1016/S2589-7500(23)00021-3)



LLMs in the Context of AI in Biomedicine



Moor, M. et al. (2023) Foundation models for generalist medical artificial intelligence. Nature, 616, 259–265.

Thank you!

Next talk in line: How to Use GPT-3.5 and GPT-4 with Python, Qiao Jin, M.D.