

Foundation Models in Medicine

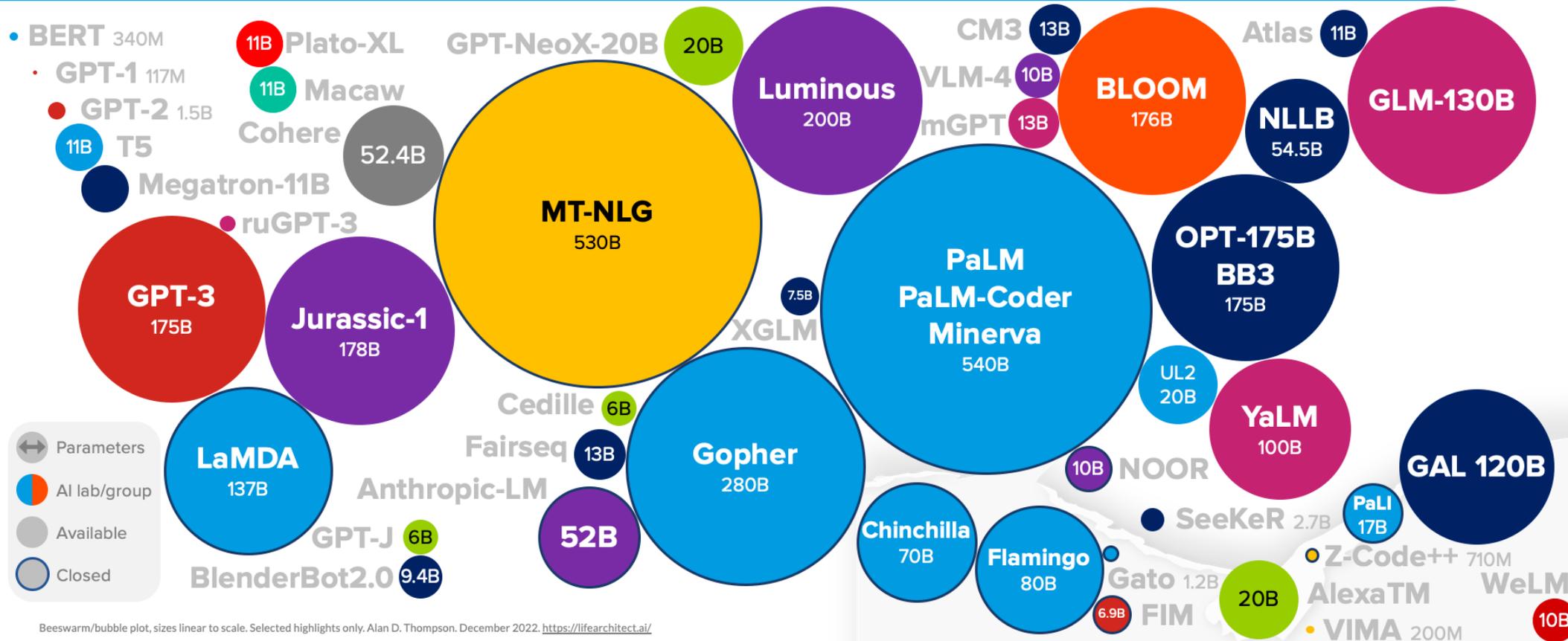
Generalist vs Specialist

Presented by Shaoting Zhang

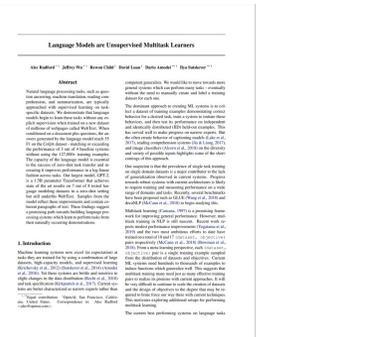
Date: 2023.12



LANGUAGE MODEL SIZES TO DEC/2022



Large-Language Models (LLM)



Transformer

- Proposed the Transformer architecture

2018/06

GPT

- Using the decoder of Transformer, a language model is trained on a large amount of unannotated text data, which is then fine-tuned for subtasks

2017/06

BERT

- Using the encoder of Transformer and pre-training on a larger dataset, the performance is much better than GPT

2019/02

GPT - 2

- Continuing to use the decoder of Transformer to explore the potential of language models, but the performance is not as impressive as previous models

GPT - 3

- Compared to GPT-2, both the data and the model are 100 times larger, with very impressive results

2020/05

2022/11

Instruct GPT

- Through instruction fine-tuning and RLHF (Reinforcement Learning from Human Feedback), the model's generalization ability is enhanced and aligned with human values, making the model controllable and safe

GPT - 4

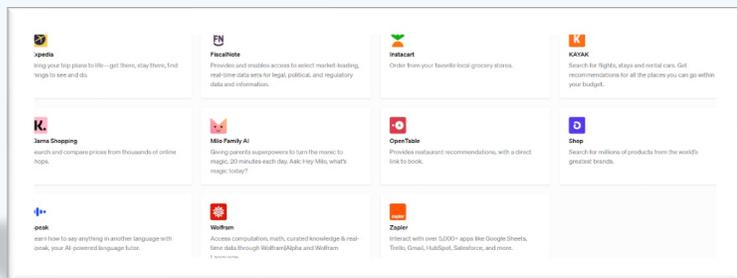
- Compared to GPT3.5, it supports image & text input, can analyze images, and provides text output

2023/03

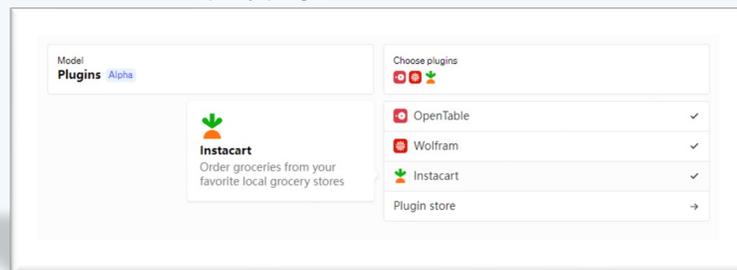


Initial support for plugins

- The first plugins have been created by [Expedia](#), [FiscalNote](#), [Instacart](#), [KAYAK](#), [Klarna](#), [Milo](#), [OpenTable](#), [Shopify](#), [Slack](#), [Speak](#), [Wolfram](#), and [Zapier](#).



- Hosting two plugins, a web browser and code interpreter (Alpha)
- Open-sourced the code for a knowledge base retrieval plugin
 - Browsing
 - Code Interpreter
 - Retrieval
 - Third-party plugins

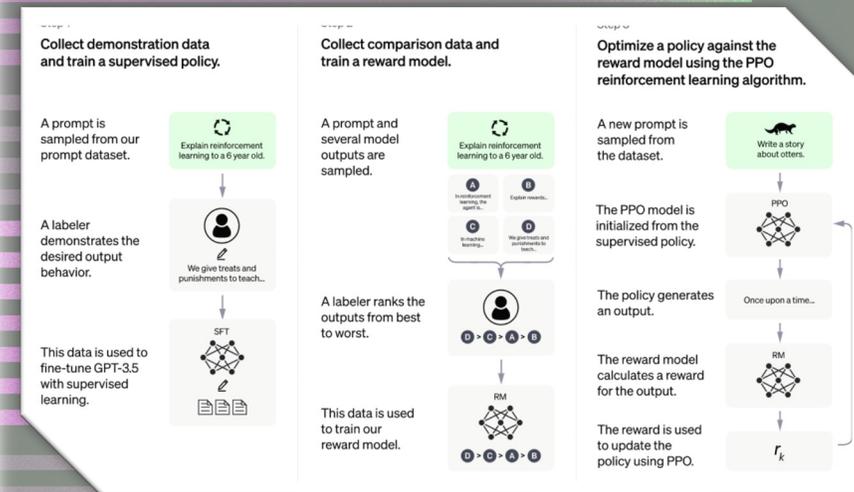


ChatGPT

We've trained a model called ChatGPT which interacts in a conversational way. The dialogue format makes it possible for ChatGPT to answer follow-up questions, admit its mistakes, challenge incorrect premises, and reject inappropriate requests.

The AI model acts as an intelligent API caller

- OpenAI plugins connect ChatGPT to third-party applications (Beta)
- Plugins enable ChatGPT to interact with APIs defined by developers
- Plugins enable ChatGPT to do things like:
 - Retrieve real-time information; e.g., sports scores, stock prices, the latest news, etc.
 - Retrieve knowledge-base information; e.g., company docs, personal notes, etc.
 - Assist users with actions; e.g., booking a flight, ordering food, etc.



Open-source Large-Language Models



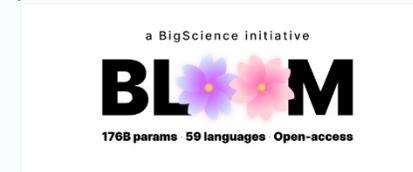
- **First unified full-stack open-source large-language model system — INTERN.**
- The InternLM (书生·浦语) 200 billion parameter version, InternLM-20B, has been officially open-sourced. The entire toolchain for the development and application of large models has been upgraded, encompassing five major stages: data, pre-training, fine-tuning, deployment, and evaluation.
- InternLM-20B was pre-trained on over 2.3T Tokens containing high-quality English, Chinese, and code data. It exhibits significant improvements in understanding, reasoning, mathematical, and programming abilities
 - ✓ Outstanding overall performance
 - ✓ Strong utility invocation capability
 - ✓ Supports a 16k context length (Through inference extrapolation)
 - ✓ Better value alignment.



- Parallel training based on RNN.
- Comparable performance with other transformer-based LLMs.



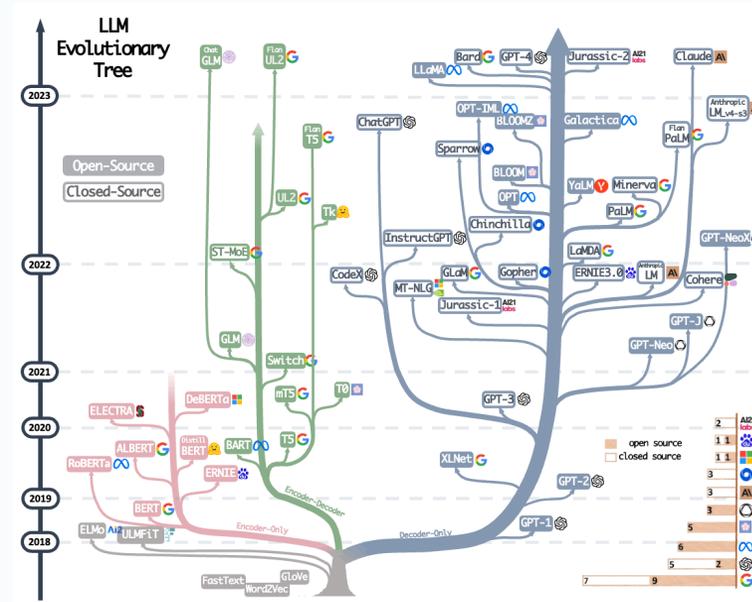
- A series of LLMs scaling from 7B to 65B.
- One of the most advanced large-scale foundational language models currently available, and the most famous open-source model.



- 176B LLM which could generate 46 natural languages and 13 programming language texts.
- Using ALiBi position embedding to have more smooth training process and better downstream performance.
- Embedding LayerNorm: Add extra layer norm after embedding layer to improve training stability.



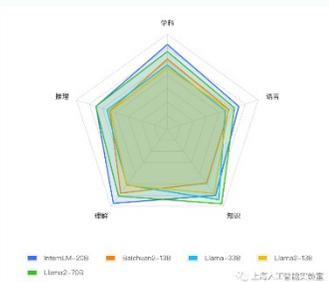
- GLM: A generative LLM, pretraining with autoregressive blank padding objects and could be fine-tuned in various natural language understanding tasks.
- New ChatGLM2-6B model: a GLM-based framework, aiming at chinese Q&A and dialogue model, users can deploy in consumer GPUs.



Baichuan

- A series of open-source LLMs free for commercial use (13B & 7B) .
- Based on Transformer, supporting English & Chinses, context window length is 4096.

<https://github.com/baichuan-inc/Baichuan-7B> <https://github.com/baichuan-inc/Baichuan-13B#readme>
<https://github.com/BlinkDL/RWKV-LM> <https://github.com/THUDM/GLM>
<https://huggingface.co/RWKV> <https://juejin.cn/post/722330585923044409>



能力维度	Llama-13B	Llama2-13B	Baichuan2-13B	InternLM-20B	Llama-33B	Llama-65B	Llama2-70B
语言	42.5	47	47.5	55	44.6	47.1	51.6
知识	58.2	58.3	45.9	60.1	64	66	67.7
逻辑	45.5	50.9	58.1	67.3	50.6	54.2	60.8
推理	42.7	43.6	44.2	54.9	46.4	49.8	55
学科	37.3	45.2	51.8	62.5	47.4	49.7	57.3
平均得分	43.8	47.3	49.4	59.2	48.5	51.9	57.4

本页所列大模型无排名先后
<https://github.com/InternLM/InternLM> <https://mp.weixin.qq.com/s/>
<https://mp.weixin.qq.com/s/oTXnvWZJVdoOpFLHngBTYQ>

<https://github.com/Hannibal046/Awesome-LLM> <https://keg.cs.tsinghua.edu.cn/jietang/publications/ChatGLM&Beyond.pdf>
<https://research.facebook.com/publications/llama-open-and-efficient-foundation-language-models/> <https://arxiv.org/pdf/2211.05100.pdf>
<https://stock.adobe.com/search?k=llama+logo> <https://chatglm.cn/blog> <https://bigscience.huggingface.co/blog/blom>



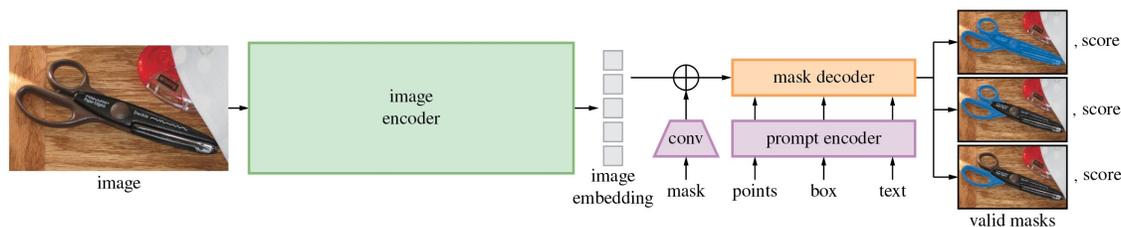
What **task** will enable zero-shot generalization?

What is the corresponding **model** architecture?

What **data** can power this task and model?

Segment Anything Model (SAM):

Build a pretrained model with large quantities of parameters in segmentation area, strengthen its generalization by prompt engineering.



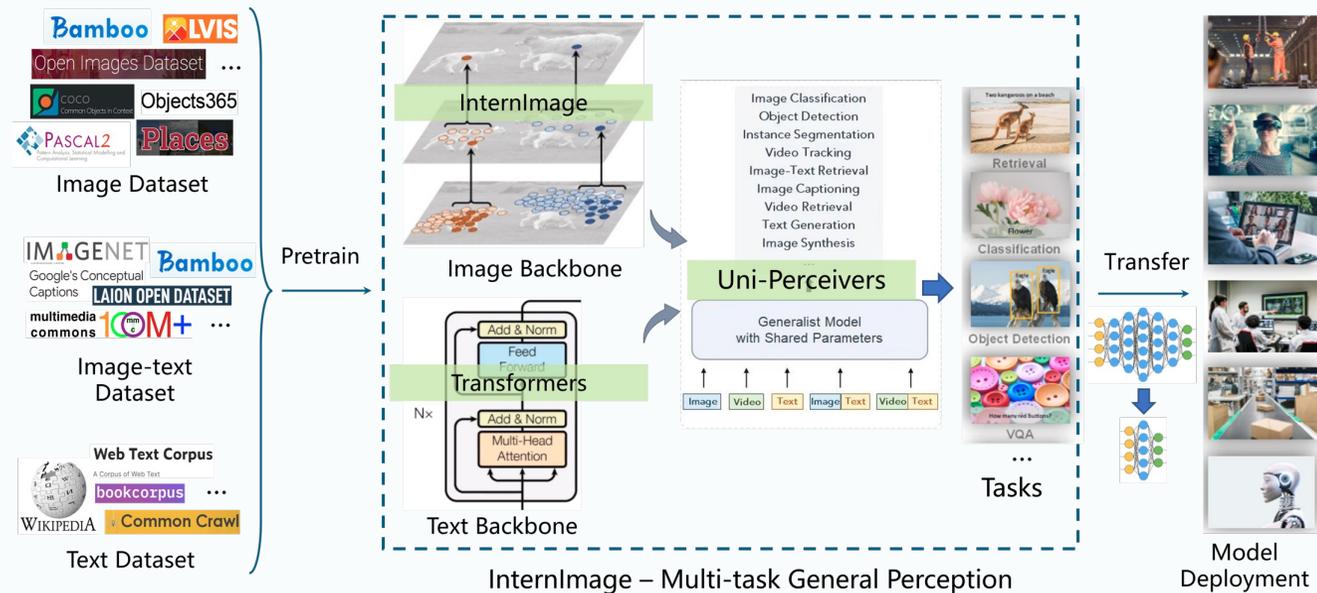
A powerful image encoder computes an image embedding, a prompt encoder embeds prompts, and then the two information sources are combined in a lightweight mask decoder that predicts segmentation masks.



Vision Foundation Models

A new generation of General Vision technology System: INTERN 2.5

"INTERN-2.5" is a powerful multimodal multitask general model jointly released by SenseTime and Shanghai AI Laboratory. It consists of large-scale vision foundation model "InternImage", pre-training method "M3I-Pretraining", generic decoder "Uni-Perceiver" series, and generic encoder for autonomous driving perception "BEVFormer" series.



分类	图像分类		场景分类				长尾分类												
	ImageNet		Places 365		Places 205		iNaturalist 2018												
BEIT-3 (pub)	ours	ViT-H/14	ours	MixMIM-L	ours	MAE-H	ours												
	89.5	90.1	60.7	61.2	69.3	71.7	86.8												
检测	常规物体检测				长尾物体检测														
	COCO		VOC 2007		VOC 2012		OpenImage												
ViT-H	ours	Eff-B7	ours	ATLDET v2	ours	冠军性能	ours	GLIP v2	ours	Swin-L	ours	PP-YOLOE	ours	DAMH40	ours	nuScenes	ours	Iter-DETR	ours
	64.5	65.5	89.3	94.0	92.9	97.2	72.2	74.1	59.8	62.5	50.9	63.2	35.6	38.8	62.4	64.8	94.1	97.2	
分割	语义分割				街景分割				RGBD分割										
	ADE20K		COCO Stuff-10K		Pascal Context		CityScapes		NYU Depth V2										
BEIT-3	ours	ViT-A-L	ours	ViT-A-L	ours	ViT-A-L	ours	ViT-A-L	ours	CMX-B5	ours								
	62.8	62.9	54.2	59.6	68.2	70.3	85.2	86.1	56.9	69.7									
图文	图像描述		微调图文检索				零样本图文检索												
	COCO Caption		COCO Caption		Flickr30k		Flickr30k												
BEIT-3	ours	BEIT-3	ours	BEIT-3	ours	BEIT-3	ours												
	147.6	148.2	76.0	76.4	94.2	94.8	88.2	89.1											

InternImage: Exploring Large-Scale Vision Foundation Models with Deformable Convolutions CVPR2023
Uni-Perceiver v2: A Generalist Model for Large-Scale Vision and Vision-Language Tasks CVPR2023
Towards All-in-one Pre-training via Maximizing Multi-modal Mutual Information CVPR2023
BEVFormer v2: Adapting Modern Image Backbones to Bird's-Eye-View Recognition via Perspective Supervision CVPR 2023

Wang W, Dai J, Chen Z, et al. Internimage: Exploring large-scale vision foundation models with deformable convolutions[J]. CVPR 2023.

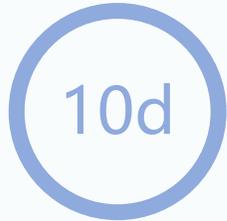
"INTERN-2.5" demonstrated world's best performance on 20 important visual benchmark datasets, covering a wide range of tasks such as classification, detection, and segmentation, making it the top-performing model across multiple domains.

High-precision, Long-term, High-efficiency: AI large model forecasting global weather

Built on multi-modal and multi-task deep learning methods

Science Large AI Model **FENGWU**

Global medium-range weather forecast AI Model released by Shanghai AI Laboratory, University of Science and Technology of China, Shanghai Jiao Tong University, Nanjing University of Information Science and Technology, The Institute of Atmospheric Physics, Chinese Academy of Sciences.



Improves the forecast skill and extend the skillful global medium-range weather forecast **out to 10 days lead for the first time.**



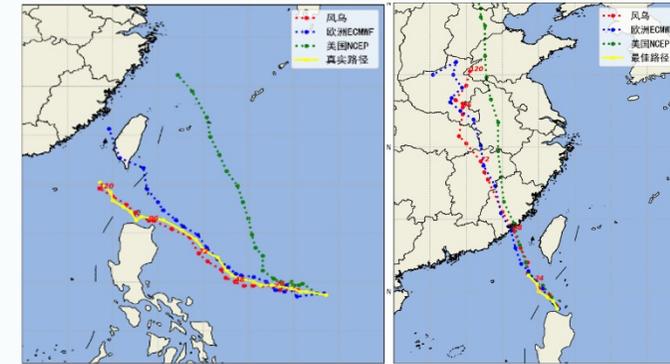
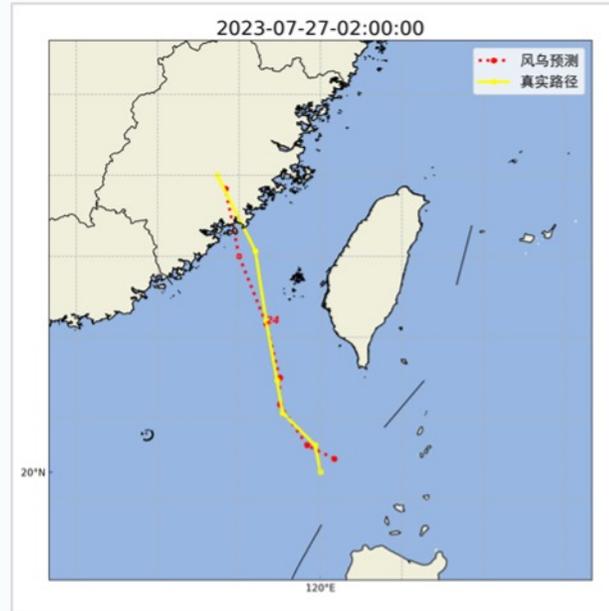
Costs **less than 30 seconds** to generate all forecasts in the following 10 days



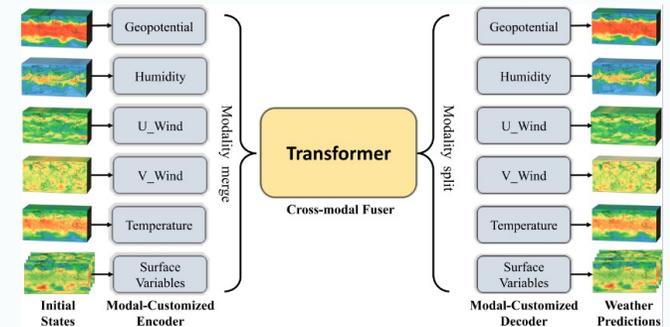
Has higher accuracy than **GraphCast (DeepMind)** on **80% reported prediction targets.**

FENGWU: PUSHING THE SKILLFUL GLOBAL MEDIUM-RANGE WEATHER FORECAST BEYOND 10 DAYS LEAD

- Recently, with the support of the China Meteorological Administration, **Shanghai AI Laboratory cooperated with the National Meteorological Center and the Shanghai Meteorological Bureau** to complete the real-time trial operation of the Fengwu model deployment. Comparative test evaluation was carried out for the two typhoons "Tali" and "Dusuri" that made landfall this year, and Fengwu showed strong business application potential, providing strong technical support for typhoon forecast and early warning.



Tracking forecasting of Typhoon Doksuri of FengWu, ECMWF and NCEP, figures above are reports from July 22th 8am and July 26th 8am, respectively.



Overview of FengWu's architecture. FengWu first **treats the multiple weather factors as different modalities** and extracts their feature embeddings independently. And then a **transformer-based network is utilized to fuse and pass messages among different modalities.**

LLM for Medicine



Nuance is introducing a new GPT-powered capability for contact center AI solutions. Here's how it will help our customers make their existing AI self-service experiences even smarter, enabling higher levels of self-service while increasing cost savings and driving revenue.

<https://www.nuance.com/index.html>
<https://whatsnext.nuance.com/customer-engagement/gpt-powered-capability-nuance-contact-center/>



patient simulator Feedback generator

random patient



<https://www.hippocraticai.com/>
<https://mp.weixin.qq.com/s/1SiQLSaa dpYgc8A1j1atcw>



<https://www.nabla.com/>
<https://twitter.com/NablaTech>



2023年，基于大模型重新定义在线问诊

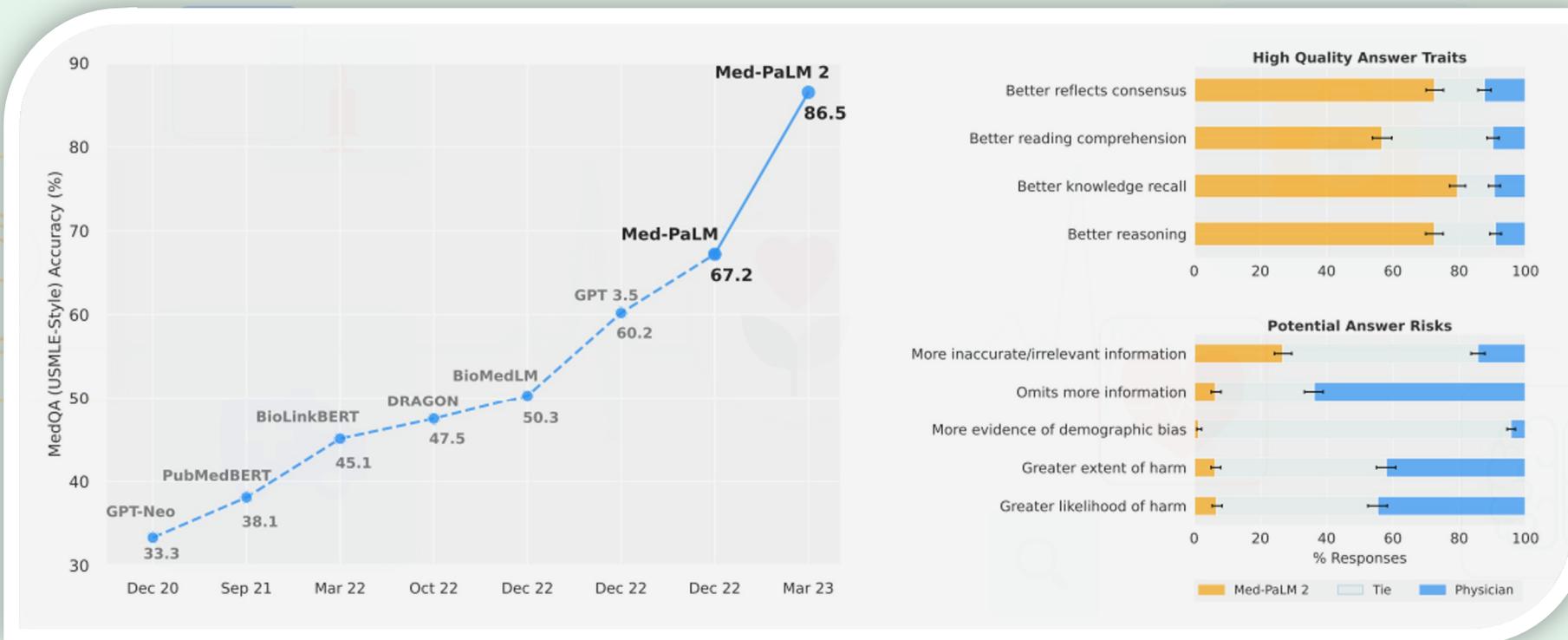


LLM for Medical

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<https://www.chunyuyisheng.com/>



<https://www.google.ca/>
<https://sites.research.google/med-palm/>



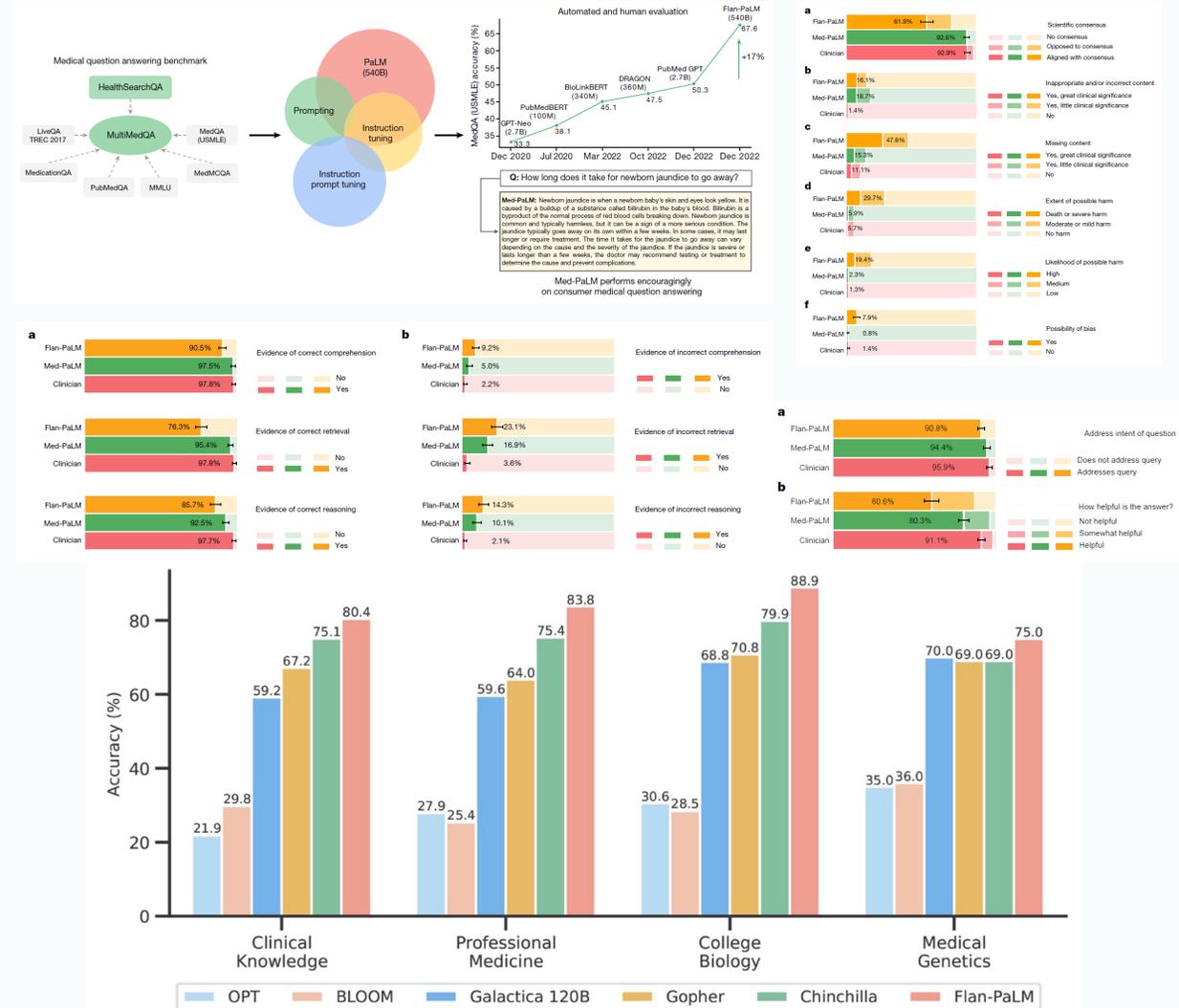


Article
Large language models encode clinical knowledge

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 Open access

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- Med-PaLM is proposed on the basis of Plan-PaLM by prompt engineering method.
- Flan-PaLM achieves SOTA in each MultiMedQA multiple-choice question tasks, its accuracy on MedQA is 67.6%.
- Compared with Plan-PaLM, Med-PaLM has significant improvement in practical problems, its accuracy is comparable with professional doctors, showing its potentials in medical fields.
- Besides Med-PaLM, a new dataset is proposed for medicine LLM evaluation.

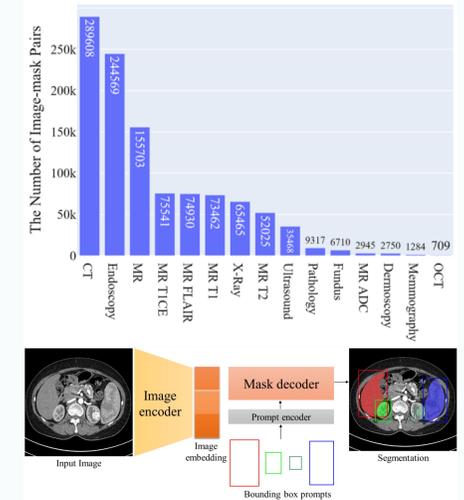
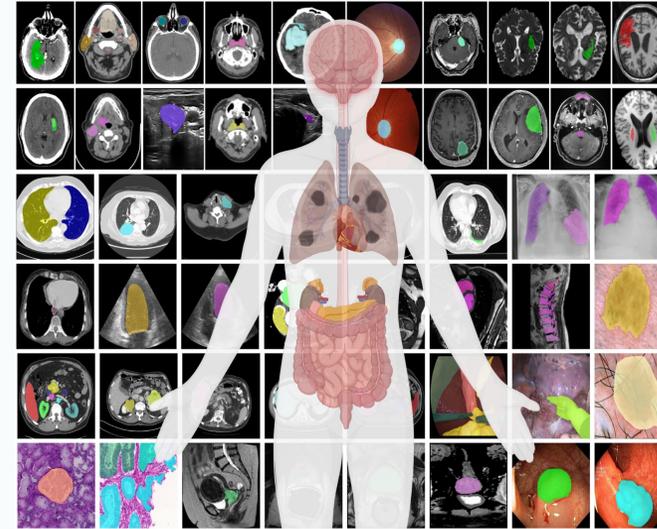


K. Singhal, S. Azizi, A. Karthikesalingam, V. Natarajan, et al. Large language models encode clinical knowledge. Nature. 2023

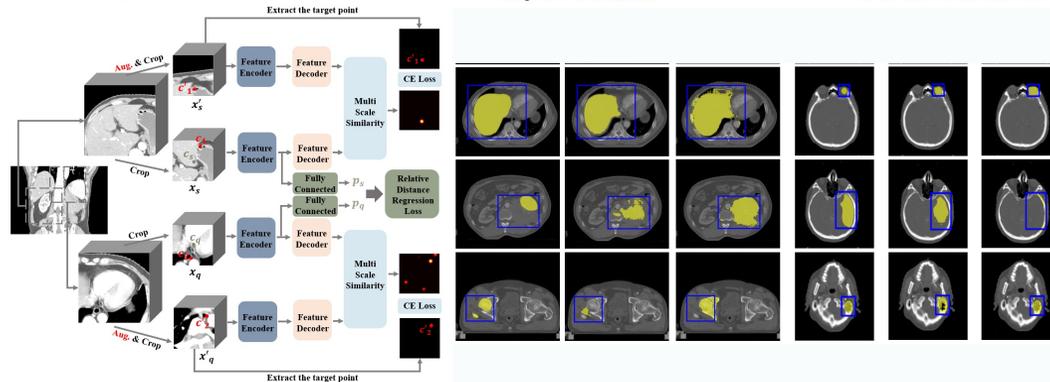
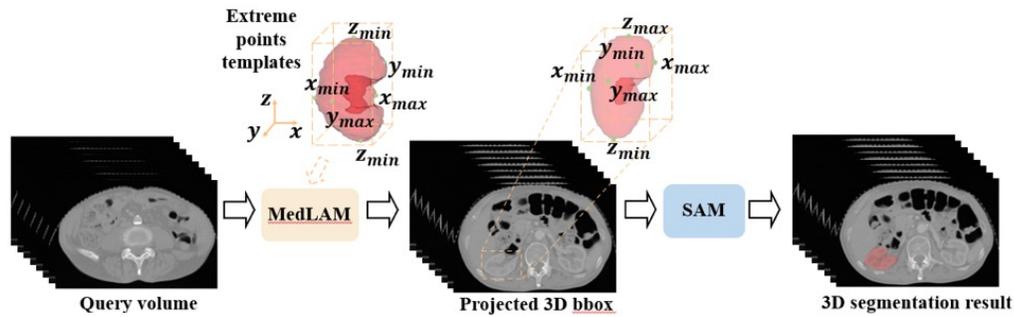
Extended Data Fig. 2 | Comparison of SOTA LLMs on MMLU clinical topics. Flan-PaLM achieves state-of-the-art performance on MMLU clinical topics.

Vision Foundation Models for Medical Images

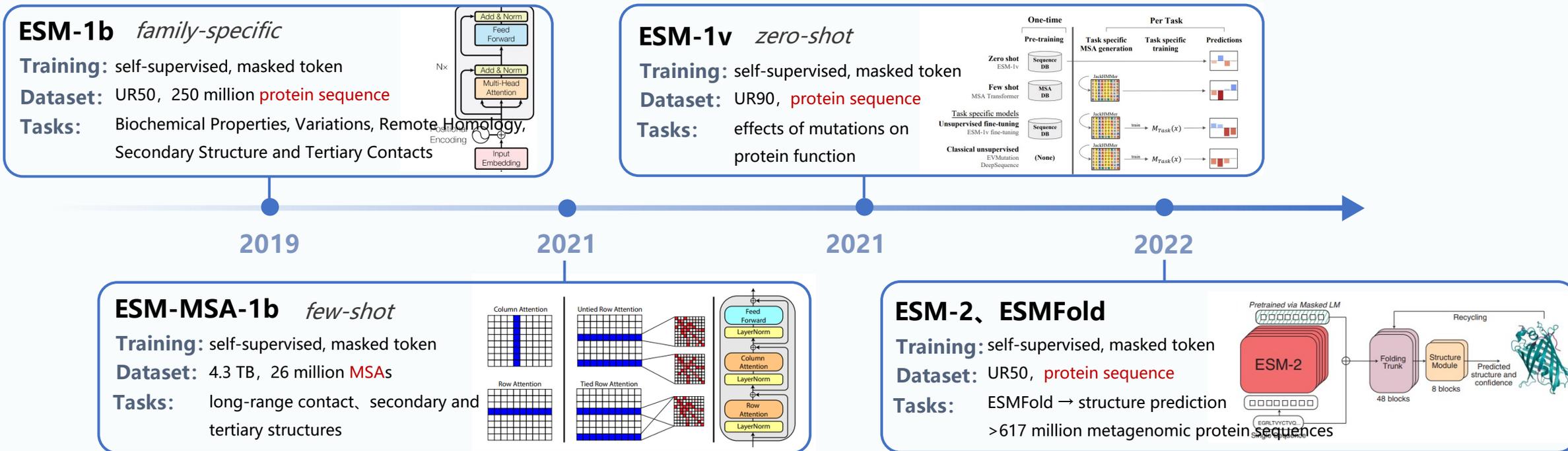
- MedLAM is the first completely automated medical adaptation of the SAM model, designed to significantly alleviate the annotation workload in the segmentation of medical image.
- Developing MedLAM, a Localize Anything Model for 3D Medical Images
- Coupling this approach with SAM's segmentation capabilities led to an efficient and accurate process for image segmentation.



- Motivated by the remarkable generality of the Segment Anything Model (SAM), MedSAM is the first foundation model for universal medical image segmentation;
- Thoroughly evaluate MedSAM through comprehensive experiments on over 70 internal validation tasks and 40 external validation tasks, spanning a variety of anatomical structures, pathological conditions, and medical imaging modalities;
- MedSAM consistently outperforms the state-of-the-art (SOTA) segmentation foundation model, while achieving performance on par with, or even surpassing specialist models. These results highlight the potential of MedSAM as a powerful tool for medical image segmentation.

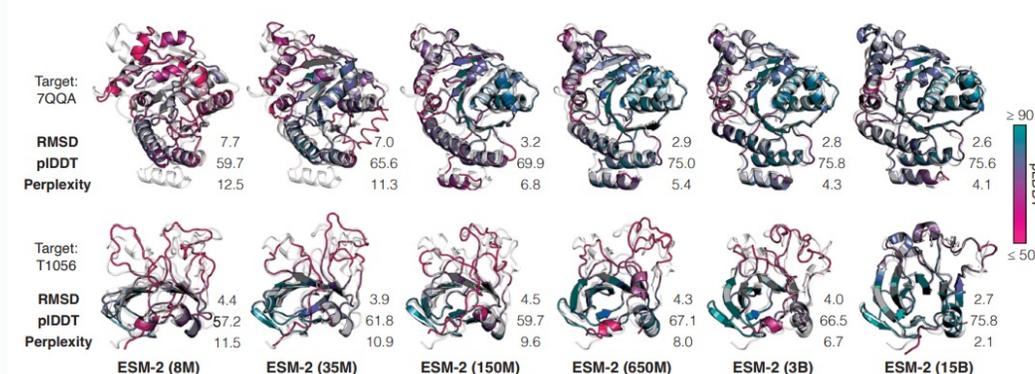
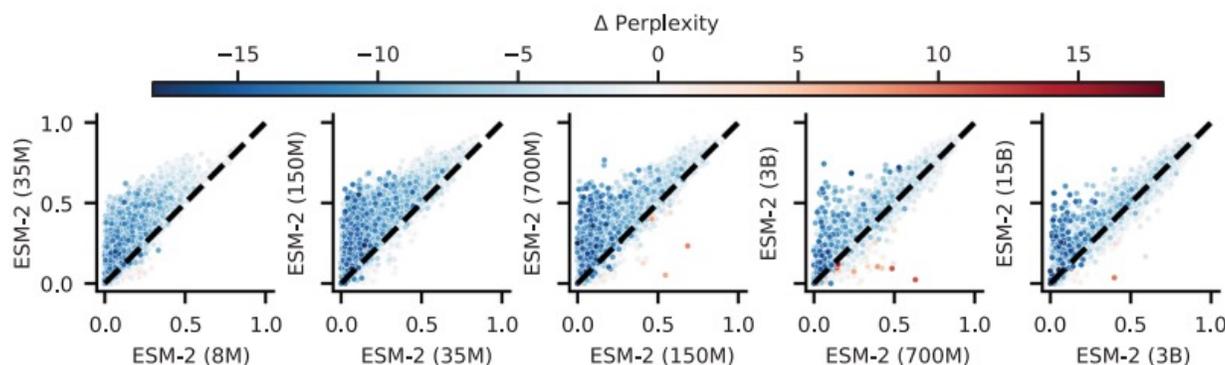
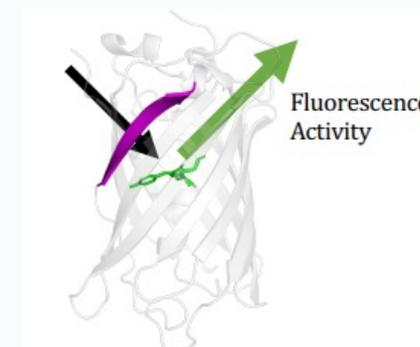


- Protein language model is the transfer application of various language models in the field of biochemistry.
- It can promote the prediction tasks such as protein structure, protein function and generation of new sequences.
- Unsupervised language models can directly utilize massive sequence information in protein databases without relying on manual annotation of sequences.
- Examples: ESM series including ESM-1b, ESM-MSA-1b, ESM-1v, ESM-2 are proposed by Meta.



- Motivation: the statistical laws of protein sequences imply their structural information → the language model can learn this law through the input sequence.
- Task: filling in missing amino acids in protein sequences → a language model has to understand the underlying structure that creates the patterns in the sequences
- Evidence: As language models of protein sequences are scaled up to 15 billion parameters, an atomic-resolution picture of protein structure emerges in the learned representations.

```
ASKGEELFTGVVPI LVELDGDVNGHKFSVSGEGEGDATYGKLT LKFICTTGKLPVPWPPTLVT
TFSYGVQCFSRYPD HMKRHDFFKSAMPEGYVQERTIFFKDDGNYKTRA EVKFEGDTLVNRIE
LKGIDFKEDGNILGHKLEYNYNSHNVYIMADKQKNGIKVNFKIRHNI EDGSVQLADHYQQNT
PIGDGPVLLPDNH YLSTQSA LSKDPNEKR D H M V L L E F V T A A G I T H G M D E L Y K
```



Emergent abilities

With the advent of large models, the term 'emergence' has become popular. It is typically described as capabilities that do not exist in small-scale models but are present in large-scale models.

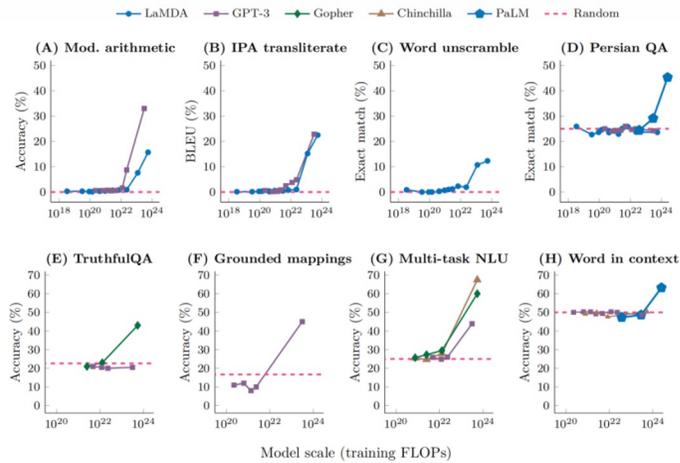
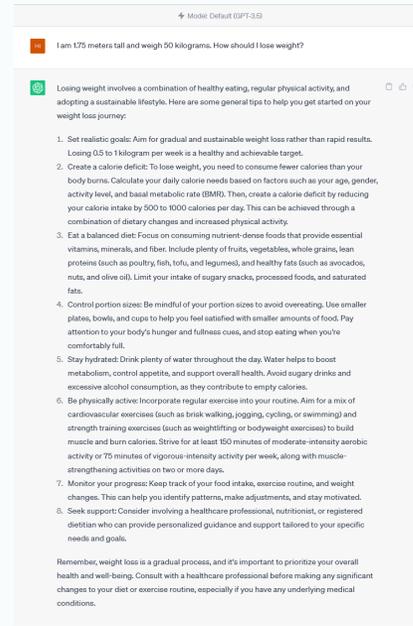


Figure 1: **Emergent abilities of large language models.** Language model families display *sharp* and *unpredictable* increases in performance at specific tasks as model scale increases. Emergent abilities [34] have also previously been labeled “abrupt, specific capability scaling” [8], “break-through capabilities” [29] and “sharp left turns” [17, 18]. Source: Fig. 2 from [34].

Are Emergent Abilities of Large Language Models a Mirage?

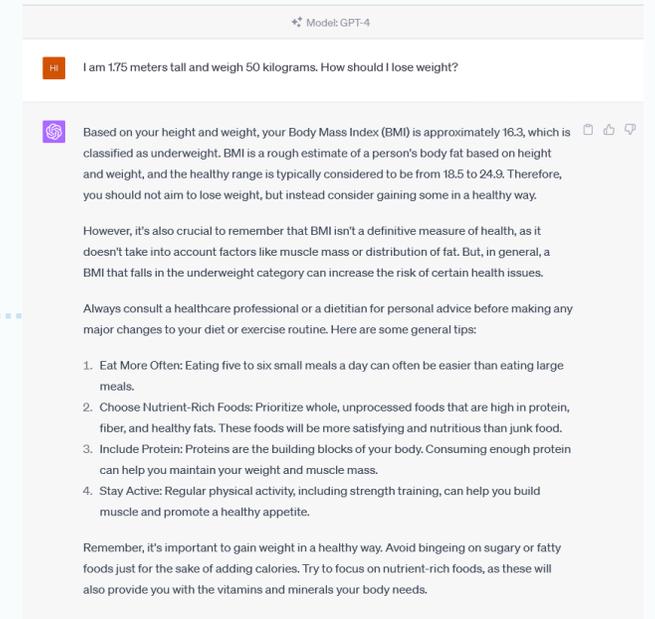
Rylan Schaeffer, Brando Miranda, and Sanmi Koyejo
Computer Science, Stanford University

"I am 1.75 meters tall and weigh 50 kilograms. How should I lose weight?"



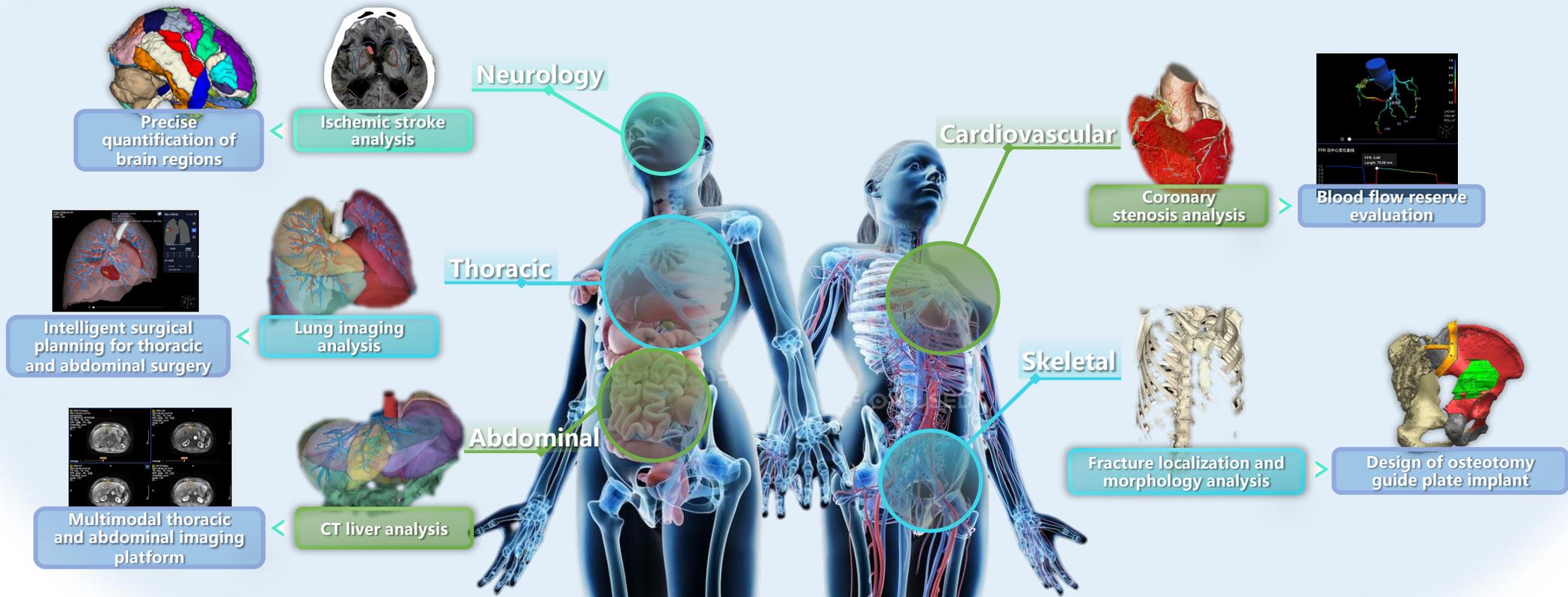
GPT 3.5

GPT 4

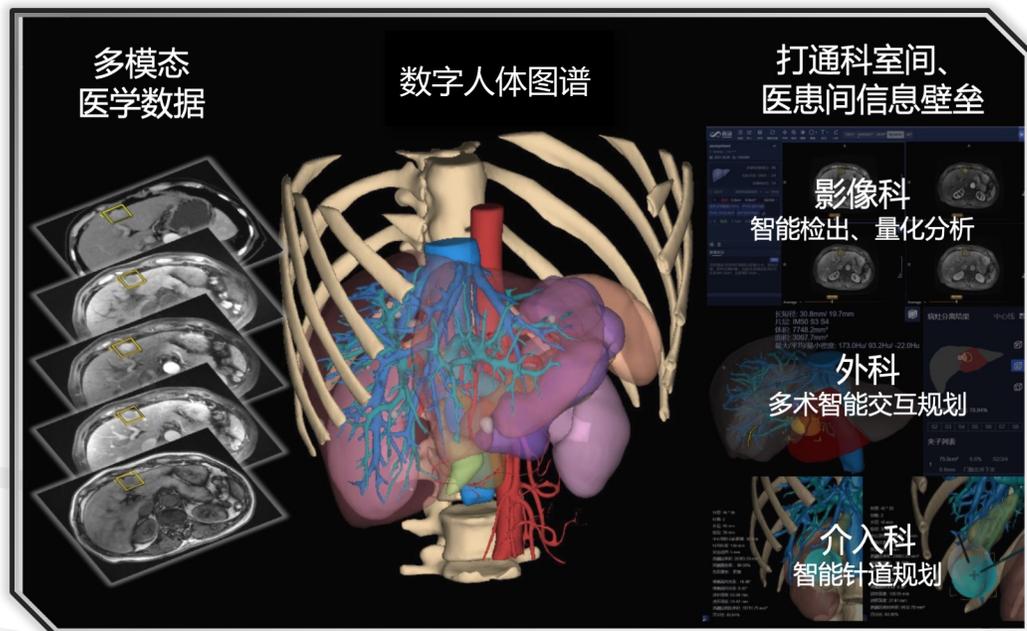


Advantage of large models: Long-tail problems

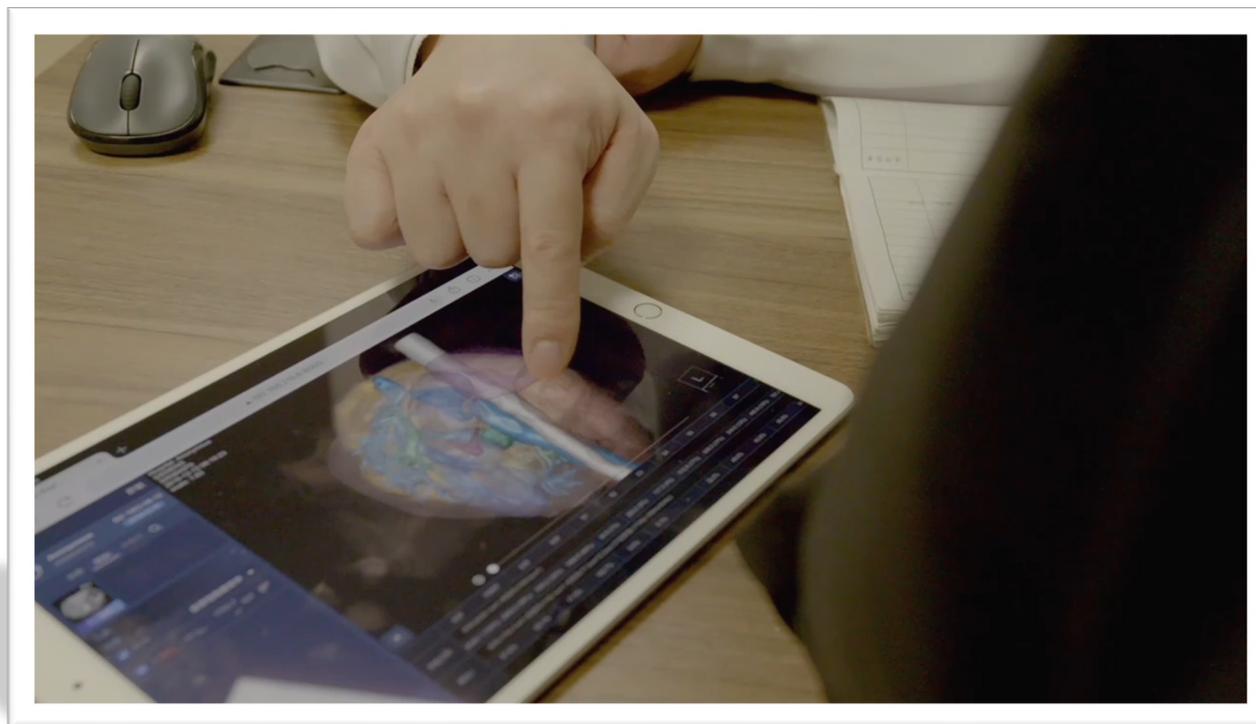
Published achievements in digital human medicine, which include 200 anatomical targets of human organs. The data modalities involved include: MRI, Ultrasound, DSA, CT, DR. Over 200+ sections of vascular networks between human organs have been restored, the precision of which surpasses the domain of what human doctors can visually distinguish between arteries and veins. This has reduced the amount of annotated data required for downstream tasks to less than a hundred instances. The model and some finely annotated data are open-source.



Advantage of large models: Long-tail problems



Based on a full-chain AI technology stack, we construct a digital human body atlas to achieve efficient and adaptive diagnosis and treatment.



Advantage of large models: Data security, MaaS

Model-as-a-Service

Automated data annotation

Intelligent annotation efficiency increased by a hundredfold

Deployment of large model inference

Large model inference efficiency improved by 100%

Parallel training of large models

Single cluster with 4000 cards - 500 billion dense parameter model

Incremental training of large models

Incremental fine-tuning cost reduced by 90%

Developer efficiency

Open model and AI development toolchain

Training

- Hardware: 384 pieces of A100, with a single card memory of 80GB, accommodating the batch training of 175 billion parameters
- Cost: \$9.6 million (48 servers, \$200,000 per server)
- Rental: Single training round ~\$3 million (AWS p4d × 2 months)

Inference

- Hardware: 8 pieces of A100 for single operation, capable of generating about 15-20 words per second (before deployment optimization)

Estimated Resources

- Training from scratch for a trillion-parameter model: 500-1000 pieces of A100
- Fine-tuning for a trillion-parameter model: 50-100 pieces of A100

GPT-3.5

Model	Input	Output
4K context	\$0.0015 / 1K tokens	\$0.002 / 1K tokens
16K context	\$0.003 / 1K tokens	\$0.004 / 1K tokens

GPT-4

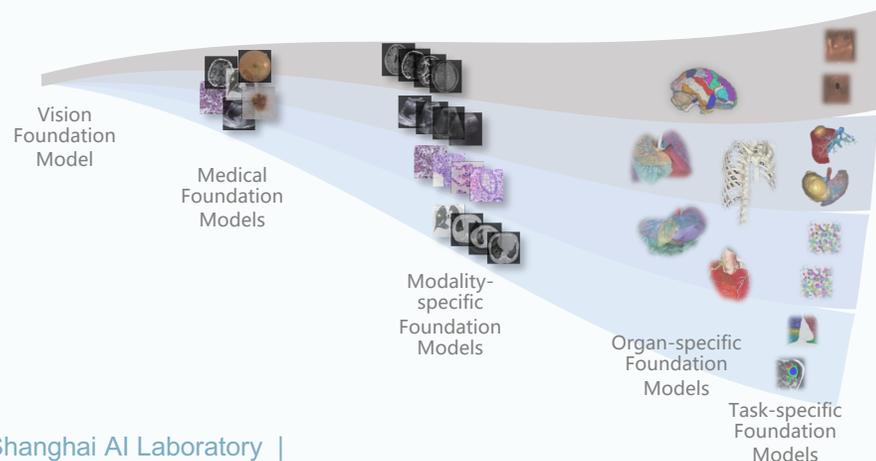
Model	Input	Output
8K context	\$0.03 / 1K tokens	\$0.06 / 1K tokens
32K context	\$0.06 / 1K tokens	\$0.12 / 1K tokens

OpenMEDLab



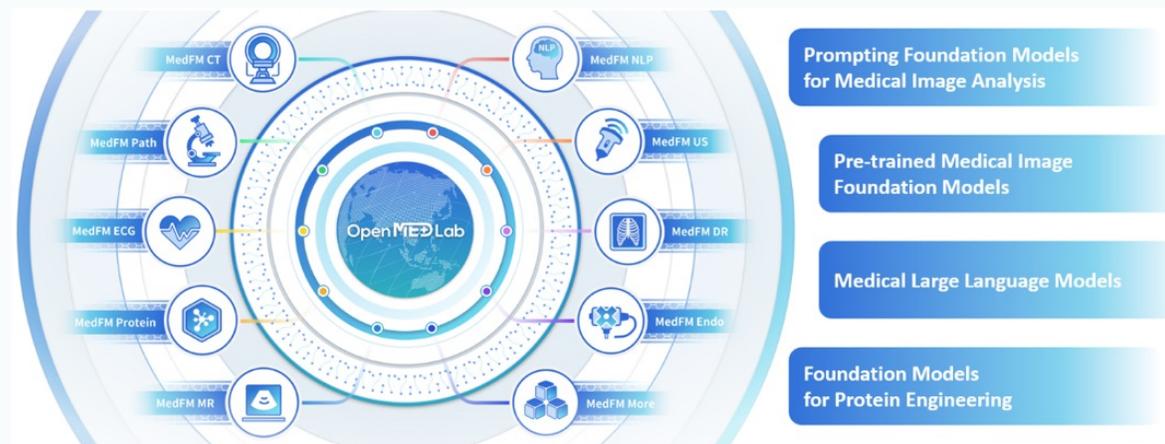
The first multi-level, multi-scenario, high-precision and implementable medical basic model group "OpenMEDLab" has recently been open sourced, integrating the world's top AI research and development capabilities, massive medical data and medical expert knowledge.

The first group of basic models released includes basic models trained on **more than 10 data modalities** such as **medical images, medical texts, biological information, and protein engineering**. Aiming at promoting cross-field, cross-disease, and cross-modal scientific research breakthroughs based on basic medical models, while helping to solve long-tail problems in the medical field and promote the industrial implementation of large medical models.



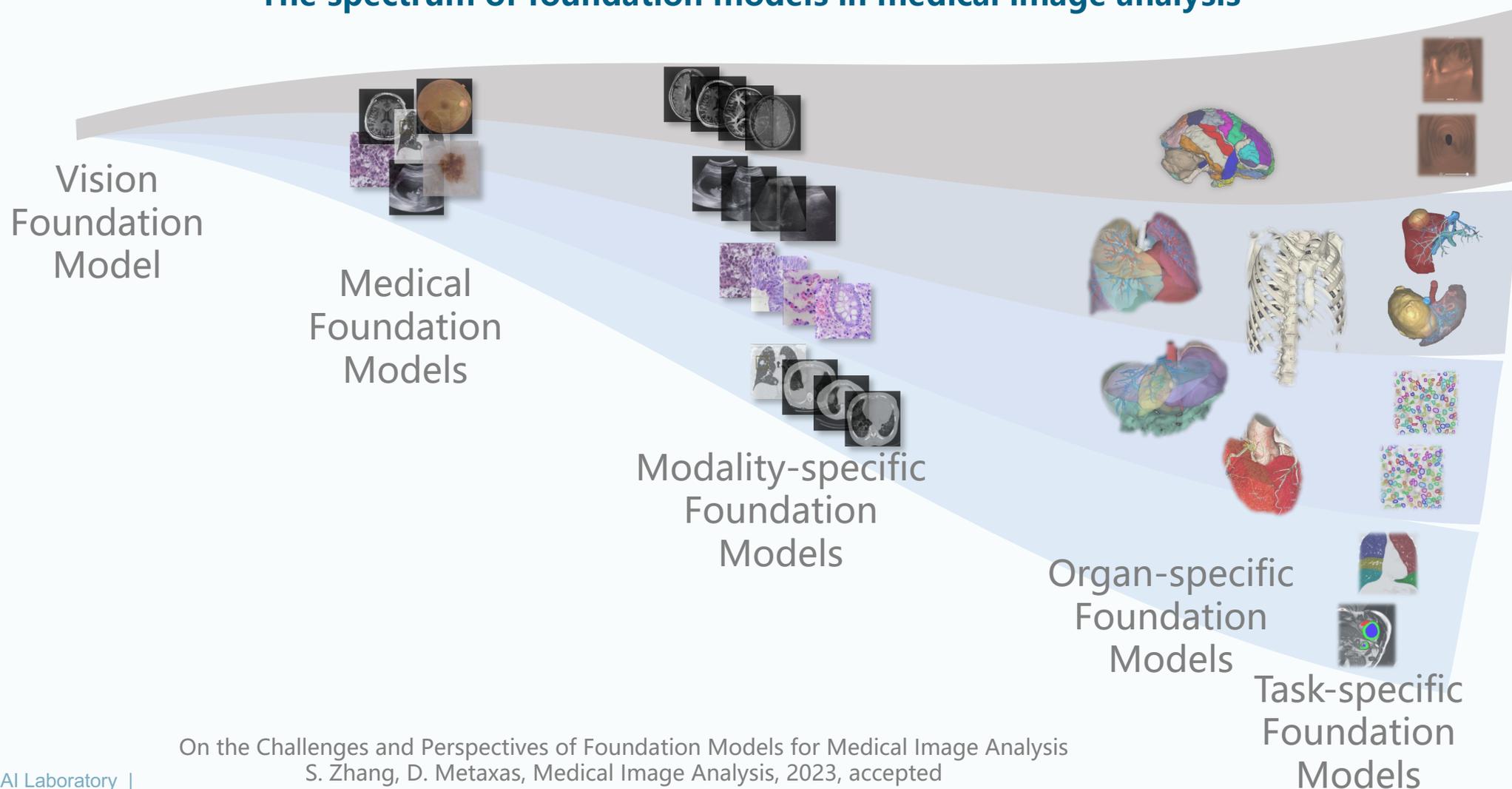
World's first open-source platform for medical foundation models "OpenMEDLab"

<https://github.com/openmedlab>



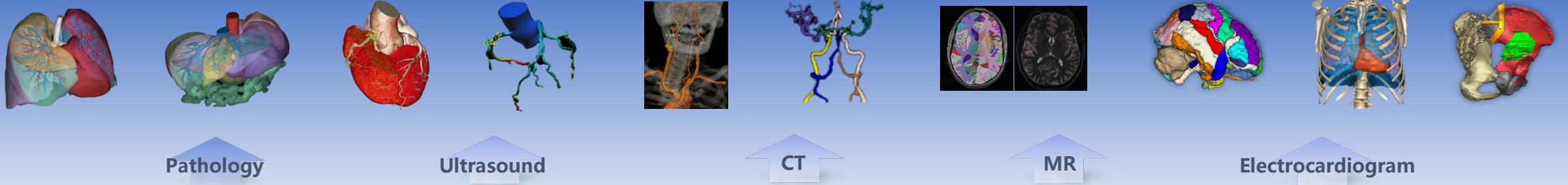
- The large Chinese medical language model, based on massive medical knowledge and **real doctor-patient interaction dialogue data**, can provide multi-scenario and multi-round conversation capabilities such as **guidance, consultation, health consultation, and assisted decision-making**.
- A series of image model groups can achieve cutting-edge research and clinical tasks such as **high-precision detection, segmentation, and classification** for different modalities such as **radiological images, pathological images, endoscopy, and ultrasound images**, and efficiently enable solutions to clinical medical imaging-based problems. The model is lightweight and can meet the clinical deployment and convenient application of innovative research.
- In terms of biomedicine, the large-scale protein sequence model is developed. It can help design and synthesize highly stable and bioactive proteins by integrating biomedical expertise, solving the common problems of long time and high cost in the development of new protein drugs.

The spectrum of foundation models in medical image analysis



On the Challenges and Perspectives of Foundation Models for Medical Image Analysis
S. Zhang, D. Metaxas, Medical Image Analysis, 2023, accepted

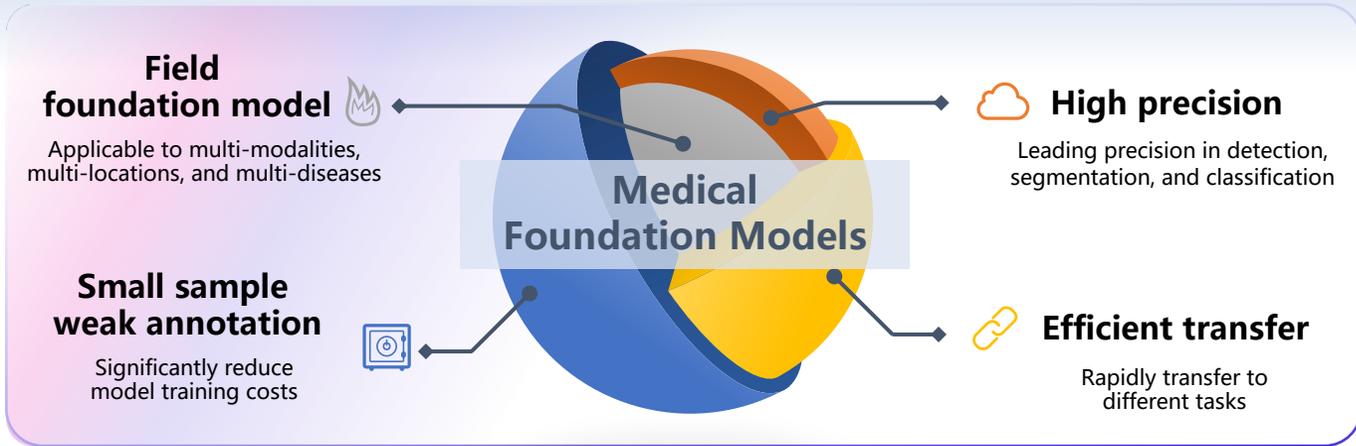
Our Related Work: Vision Foundational Model



Building global academic influence

Medical Image Analysis

Special issue on foundational models
IF 13.8



WAIC 2023
officially released

Grand Challenge
International challenge



Our Related Work: Vision Foundational Model

Grand Challenge

A platform for end-to-end development of machine learning solutions in biomedical imaging.

82,000+ users 356 challenges 2,750 algorithms

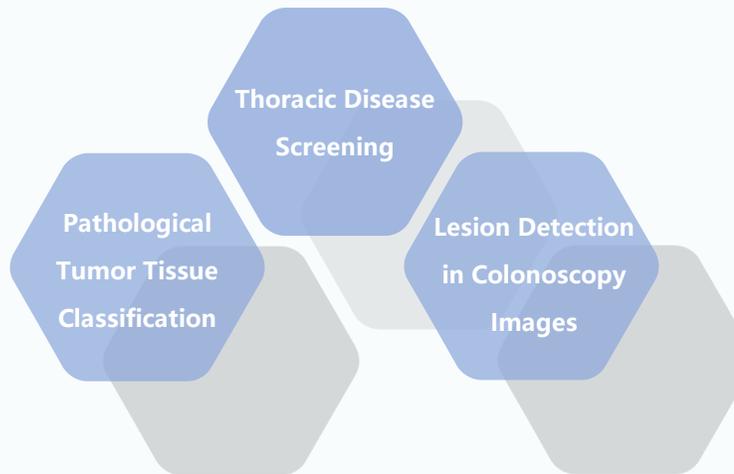


MedFMC: Foundation Models for Medical Image Analysis Challenge

This challenge aims to promote technological innovation and explore new learning paradigms in the field of medical image analysis.

Using a small number of data (1-shot, 5-shot, 10-shot) provided by the challenge, transferring the large-scale pre-trained general foundation model to three medical image classification tasks.

Three Medical Image Classification Tasks



Abnormality	#Sample	Abnormality	#Sample
Cardiomegaly	1300	TB	305
Pleural effusion	1266	Pneumothorax	284
Pneumonia	898	Atelectasis	199
Hilar enlargement	758	Emphysema	159
Nodule	665	Calcification	104
Aortic calcification	645	Pulmonary edema	86
Tortuous aorta	612	Increased lung markings	53
Fibrosis	532	Consolidation	34
Thickened pleura	497	Elevated diaphragm	23
Fracture	409		

	Normal Tissue	Tumor Tissue
	#Sample 6494	#Sample 3515
	#Slide 249	#Slide 147

Leison	#Sample	#Study
Ulcer	790	69
Erosion	951	70
Polyp	390	62
Tumor	136	11



(Sponsors)

<https://medfm2023.grand-challenge.org/medfm2023/>

"A Real-world Dataset and Benchmark For Foundation Model Adaptation in Medical Image Classification" by X. Wang, D. Wang, L. Wang, M. Li, S. Zhang, et al. Scientific Data. 2023 Accepted



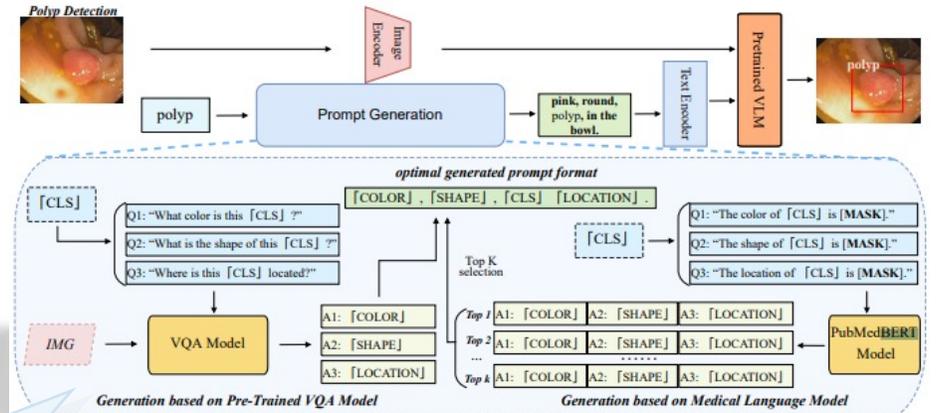
Medical Image Understanding with Pretrained Vision Language Models: A Comprehensive Study

- As one of the pioneering work in applying the visual-language pre-trained model to medical image analysis area, this work first explored how to automatically generate appropriate **language prompts** to guide the pre-trained model identifying the target object in few-shot or even zero-shot situations.
- Using the invariance of descriptive languages such as color and shape in different fields, we can recognize some unseen concepts even in the case of zero-shot tasks. And by inducing a large language model (LLM) to complete adjectives, we successfully automates the generation of these descriptive hints.

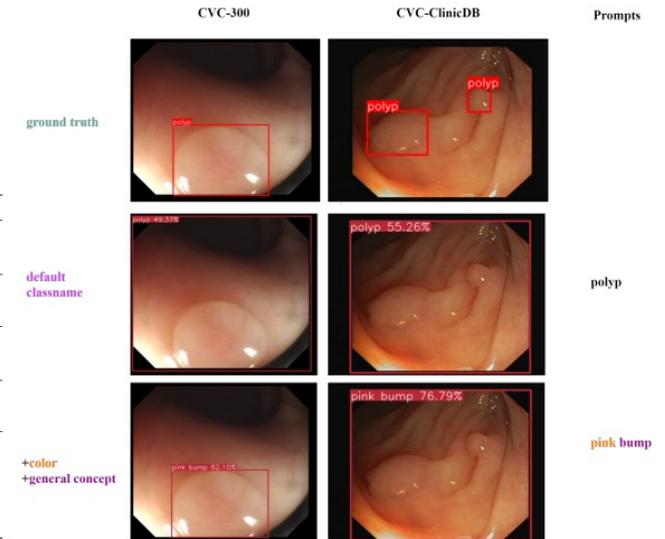
The method in the paper demonstrates the absolute superiority over traditional methods in low-resource scenarios on thirteen medical public datasets spanning multiple modalities.

Table 2: Our approaches v.s. supervised models on non-radiology datasets (AP%).

	Method	Backbone	ISIC 2016	DFUC 2022	Polyp (x5)	BCCD	CPM-17	Avg.
Full Data	Faster RCNN	RN50	50.3	42.3	56.6	56.9	39.8	49.2
	RetinaNet	RN50	54.0	43.1	58.8	56.7	35.7	49.7
	DyHead	Swin-T	52.9	44.2	62.9	60.1	38.8	51.8
	GLIP-T(default cls)	Swin-T	62.4	50.3	68.1	62.5	43.9	57.4
	Ours (Manual)	Swin-T	64.1	50.3	69.4	62.2	43.4	57.9
	Ours (Auto)	Swin-T	61.6	50.1	68.8	63.1	44.2	57.6
100-Shot	Faster RCNN	RN50	44.6	27.0	44.9	38.6	-	38.8
	RetinaNet	RN50	41.7	28.4	41.7	54.3	-	41.5
	DyHead	Swin-T	42.5	27.8	42.5	40.5	-	38.3
	GLIP-T(default cls)	Swin-T	55.9	41.4	57.6	59.8	-	53.7
	Ours (Manual)	Swin-T	58.0	43.7	60.8	60.1	-	55.7
	Ours (Auto)	Swin-T	58.8	42.4	60.8	60.2	-	55.6
Zero-Shot	GLIP-T(default cls)	Swin-T	20.1	0.1	4.1	0.7	7.6	6.5
	GLIP-L(default cls)	Swin-L	20.4	3.6	11.9	10.4	11.6	11.6
	Ours (with MLM)	Swin-T	25.1	24.8	38.4	24.1	20.3	26.5
	Ours (with VQA)	Swin-T	23.5	12.9	27.1	14.3	26.2	20.8
	Ours (with Hybrid)	Swin-T	24.5	22.5	35.1	14.3	24.8	24.2
	Ours (Manual)	Swin-T	33.3	25.9	41.3	26.9	31.4	31.8



Overview of the proposed approach: The optimal medical prompts can be automatically generated with the help of pre-trained VQA model, medical language model, or a hybrid of both.



By adding descriptive prompts, the accuracy of model recognition can be greatly improved without additional training

Endo-FM

- Constructing a large-scale endoscopic video dataset consisting of over 32K video clips (5M frames), encompassing varying modalities, target organs, and disease types.
- Developing a foundation video transformer model capable of capturing both local and global long-range dependencies across spatial and temporal dimensions.
- Endo-FM achieves promising performance when applied to various downstream tasks, surpassing state-of-the-art methods by a significant margin.

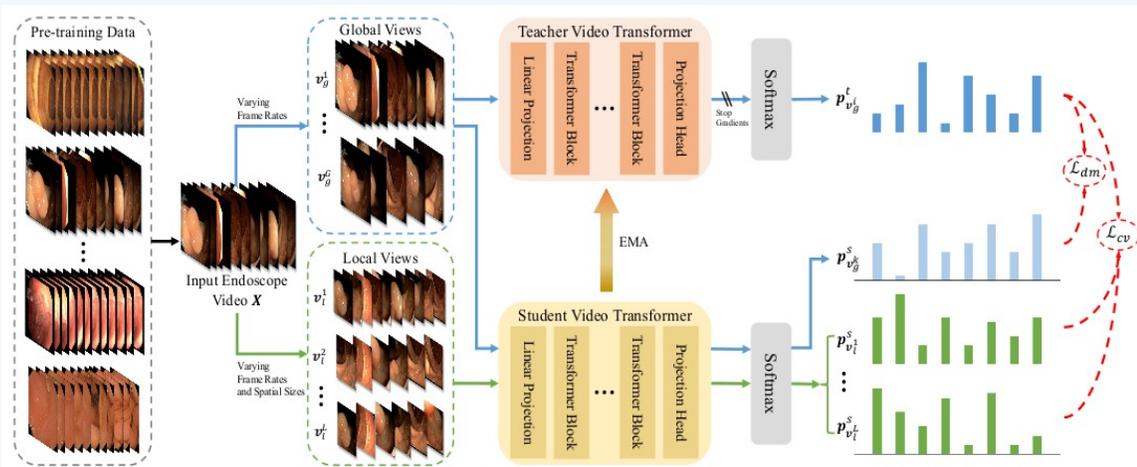
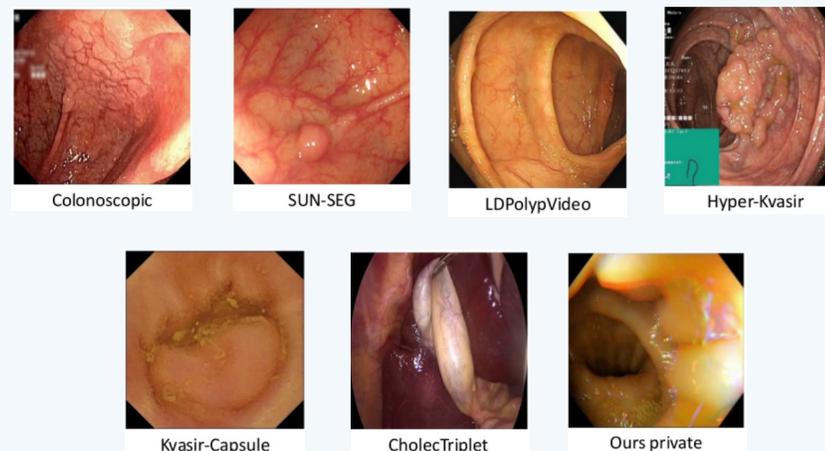


Table 2. Comparison with other latest SOTA methods on 3 downstream tasks. We report F1 score for PolypDiag, dice for CVC-12k, and F1 score for KUMC.

Method	Venue	PolypDiag	CVC-12k	KUMC
Scratch (Rand. init.)		83.5±1.3	53.2±3.2	73.5±4.3
TimeSformer [3]	ICML'21	84.2±0.8	56.3±1.5	75.8±2.1
CORP [13]	ICCV'21	87.1±0.6	68.4±1.1	78.2±1.4
FAME [10]	CVPR'22	85.4±0.8	67.2±1.3	76.9±1.2
ProViCo [26]	CVPR'22	86.9±0.5	69.0±1.5	78.6±1.7
Qian <i>et al.</i> [28]	ECCV'22	87.6±0.6	69.1±1.2	78.1±1.9
ST-Adapter [25]	NeurIPS'22	84.8±0.7	64.3±1.9	74.9±2.9
Endo-FM (Ours)		90.7±0.4	73.9±1.2	84.1±1.3



Example frames of 6 public and ours private endoscope video datasets.

nature

Article

A foundation model for generalizable disease detection from retinal images

<https://doi.org/10.1038/s41586-023-06555-x>

Received: 5 December 2022

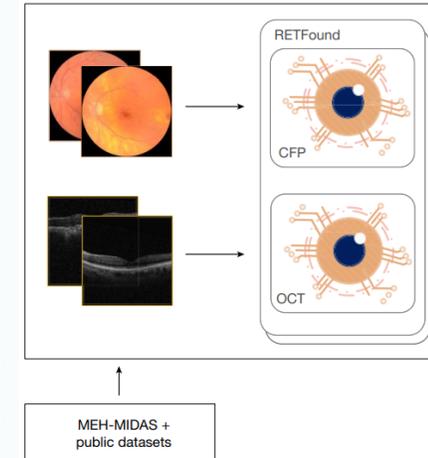
Accepted: 18 August 2023

Published online: 13 September 2023

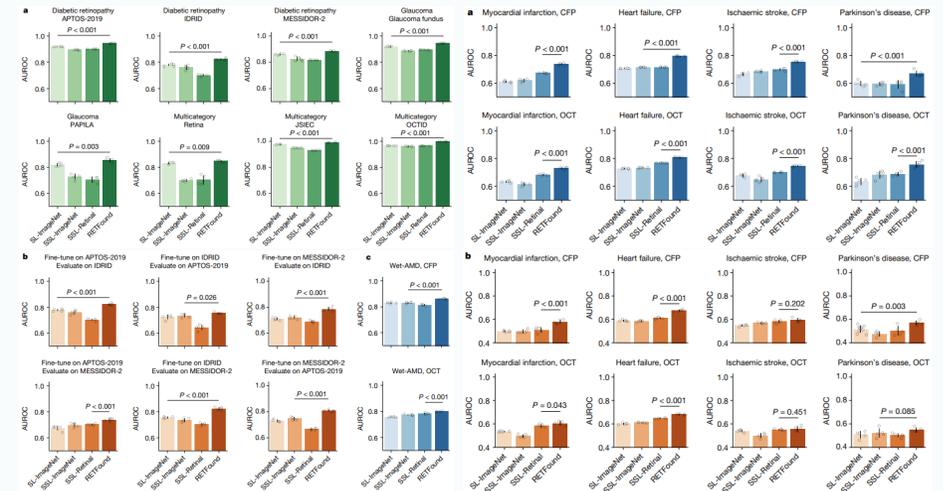
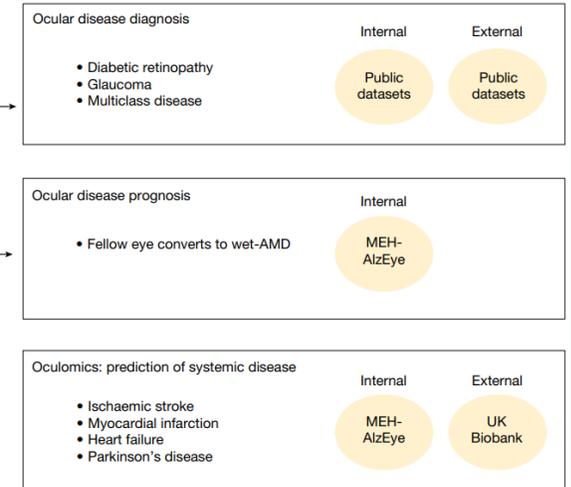
Yukun Zhou^{1,2,3,5*}, Mark A. Chia^{2,4}, Siegfried K. Wagner^{2,4}, Murat S. Ayhan^{1,2,4}, Dominic J. Williamson^{1,2,4}, Robbert R. Struyven^{1,2,4}, Timing Liu², Moucheng Xu^{1,3}, Mateo G. Lozano^{2,5}, Peter Woodward-Court^{1,2,6}, Yuka Kihara^{7,8}, UK Biobank Eye & Vision Consortium*, Andre Altmann^{1,3}, Aaron Y. Lee^{2,8}, Eric J. Topol⁹, Alastair K. Denniston^{10,11}, Daniel C. Alexander^{1,12} & Pearse A. Keane^{2,4,5,6*}

- Investigated the development and validation of a foundational model for retinal images, RETFound, with the objective of mitigating the AI model's heavy reliance on massive labeled data and bolstering its generalization capability in disease detection tasks.
- RETFound stands as the inaugural foundational model in the field of ophthalmology, adopting one of the most extensive training datasets and being validating on the most comprehensive set of validation tasks. This open-source model can be tailored to custom downstream tasks, promoting diverse ophthalmological research.
- Trained on 1.6 million unlabeled retinal images by a manner of Self-Supervised Learning and later adapted to disease detection tasks based on explicit labels for the detection of various diseases.
- RETFound outperformed the other pretraining strategies using only 10% of labeled training data and showed consistently high adaptation efficiency, demonstrating the potential of this foundation model in alleviating data shortages and adapting to downstream tasks.
- RETFound provides a generalizable solution to improve model performance and alleviate the annotation workload of experts to enable broad clinical AI applications from retinal imaging.

Stage 1: Self-supervision on retinal images



Stage 2: Supervised fine-tuning for clinical tasks



Our Related Work: Large Language Models



- **Large-scale Training:** A model optimized on approximately 4,000,000 instruction datasets from both the Chinese medical domain and general domain.
- **Comprehensive Support for Chinese Medical NLP Tasks:** Including health education, physician examinations, report interpretation, medical record structuring, and simulated diagnostics and treatments.



MedGPT Intelligent consultation display



专业医生的帮助和建议。

最近加班很辛苦，眼睛都黄了。。。

您好，感谢您向我咨询。我需要更多的信息才能帮助确定您的症状可能的原因和建议下一步的行动。首先我想问一下：除了眼睛发黄之外，还有其他不适的症状吗？例如视力模糊、眼痛或头痛等？

眼睛没有不舒服，但是手掌有点黄了

好的，感谢您提供的额外信息。我的第一个问题是，这种情况已经持续了多久了？

好像有一个多月了

非常抱歉听到您的不适。基于您的描述，这可能是黄疸的症状之一。我建议您尽快预约一次肝功能检查以了解更多情况并排除任何潜在的健康问题。此外，我也想提醒您注意休息和保持健康的生活方式以帮助缓解症状和促进康复。如果您有任何进一步的问题或需要进一步的医疗咨询，请随时与我联系。

那我应该去哪个科室看病呢

根据您的症状描述，可能需要看内科或者消化科的医生。他们可以对您进行全面的身体检查和相关检测以确定病因并提供适当的治疗方案。同时也可以考虑去眼科做一下视力、验光等方面的相关检查以排除是否存在与视力和眼部疾病相关的因素导致黄疸的出现。

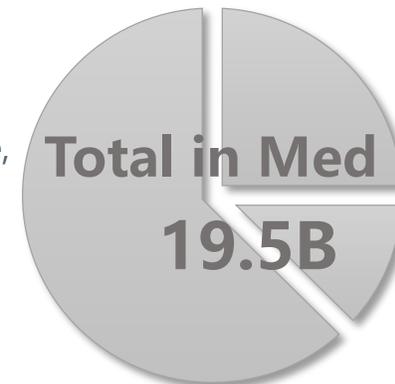
请输入...



Self-supervised Pre-training

Medical Domain Corpus: Textbook, Guideline, EHR, papers, etc. ~12B tokens

General Domain Corpus : Wiki, Webpage, News, etc. ~230B tokens



Instruction Tuning

Medical Domain Instruction Tuning Dataset:

1 language, 17tasks, 19 types of prompt, 2.7B

Generic Domain Instruction Tuning Dataset:

46 languages, 16 tasks, 16 types of prompt, 5.3M

Medical Domain QA Dataset: ~4B tokens

Generic Domain QA Dataset: ~1.2B tokens



Plugins

General Plugins: Search Engine, Sci Calculator, ~600M tokens

Medical Plugins: Knowledge Retrieval, BMI Calculation, ~109M tokens



RLHF

Medical Domain Dialog Dataset:

7 scenarios, ~49M tokens

Generic Domain Dialog Dataset:

16 languages, ~72M tokens

Generic Domain Task Dataset:

Chinese & English, 16 tasks, ~35M tokens

Our Related Work: Large Language Models

Medical Domain Supervised Fine-tuning

Collect demonstration data and train a supervised policy

- **Instructions:** NER, plugins, etc.
- **Multi-round dialog:** Consultation, triage, etc.
- **QA:** physical examination, drug consultation, etc.
- **Rejection:** refuse to respond non-medical questions

Reward Model Training

Collect comparison data and train a reward model

Design a prompt for self-evaluation

RL Model Training

Optimize a policy against the reward model with RL

Improve the basic PPO framework

Balance the exploration & exploitation

SFT Prompt

[USER]

检查途径: 经体表 图像等级: 丙 检查项目: 二维 M型 彩色多普勒 (脉冲式 连续式) 组织多普勒 一、M型主要测值 (单位 mm): 名称 测量值 正常参考值 主动脉根部内径 33 20-37 左房内径 35 19-40 左室舒张末期内径 48 35-56 左室收缩末期内径 28 20-37 室间隔厚度 11 6-11 左室后壁厚度 10 6-11 二、二维超声心动图描述: 1.各房室无明显扩大。2.室间隔基底部偏厚,约12mm,静息状态下左室壁各节段收缩活动未见明显异常。3.各心瓣膜未见明显增厚,开放不受限。三、彩色多普勒超声描述: 1.房、室间隔水平未见明显分流。2.二尖瓣轻微反流。舒张期经二尖瓣口血流:E=63cm/s, A=69cm/s, E/A=0.9。3.主动脉瓣未见明显反流。4.三尖瓣轻微反流,最大反流速度约2.0m/s,估测肺动脉收缩压约27mmHg。5.经左室流出道收缩期血流速度未见明显异常。四、左心功能测定: 名称 测量值 名称 测量值 左室舒张末期容量(ml) 106 左室收缩末期容量(ml) 30 左室射血分数(%) 71 左室短轴缩短率(%) 40 每搏输出量(ml) 76 五、组织多普勒检查: 二尖瓣瓣环水平: 室间隔侧 E'=6.9cm/s, E/E'=9.2, 左室侧壁 E'=11.3cm/s, E/E'=5.6。

根据上文中信息,判断主动脉根部内径是什么?请提取文中对应的值

[AI]

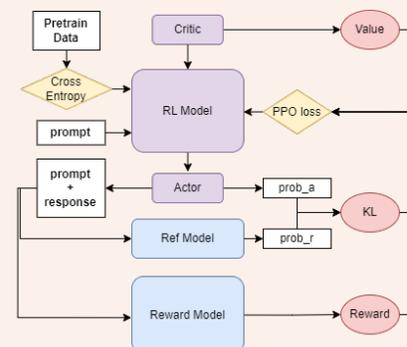
根据上文中信息,主动脉根部内径为33毫米,正常参考值为20-37毫米。

RLHF Labeling Tool

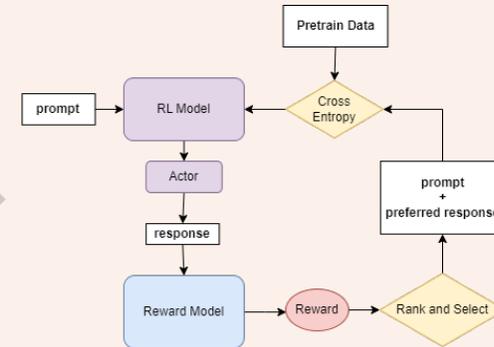


The tool displays a grid of 12 prompts related to white blood cell counts. Each prompt is accompanied by a label (e.g., '非反人类', '反人类') and a '医疗相关' (Medical Related) checkbox. The prompts are numbered 1/12 through 12/12.

RL Models



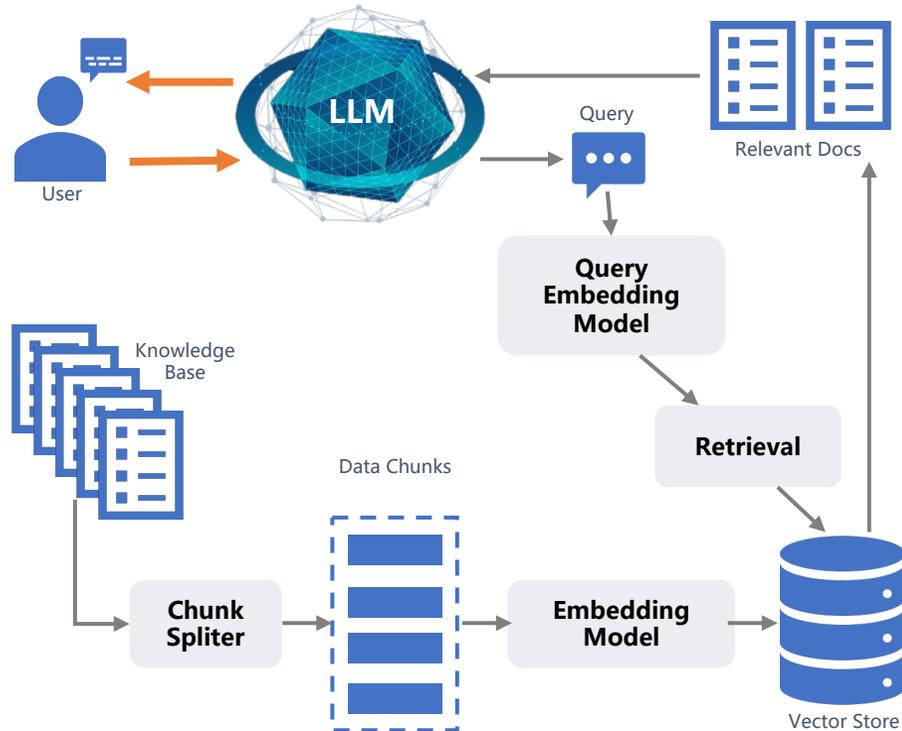
Standard PPO



Ours

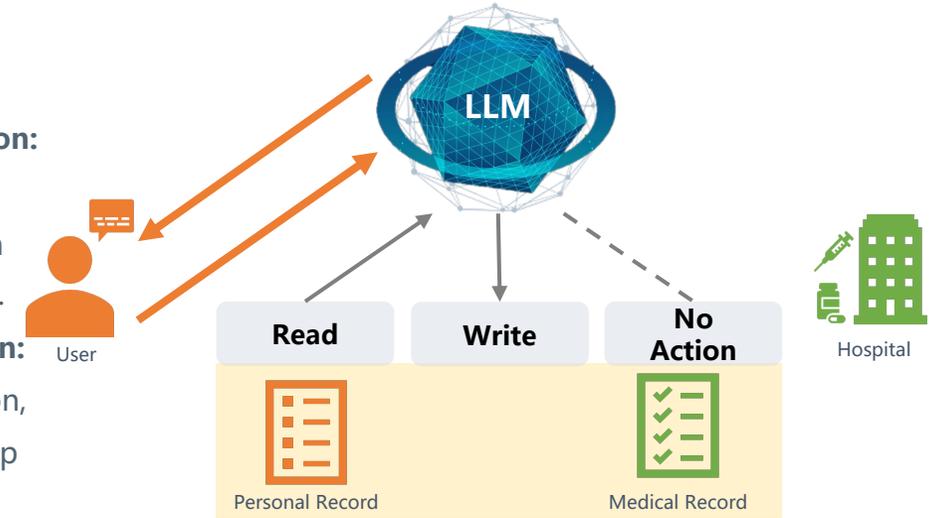
Customized Knowledge Base

The model can search within a specified knowledge base based on the question posed. It summarizes the content of the related documents retrieved and generates a response to the question.



Long-range Memory

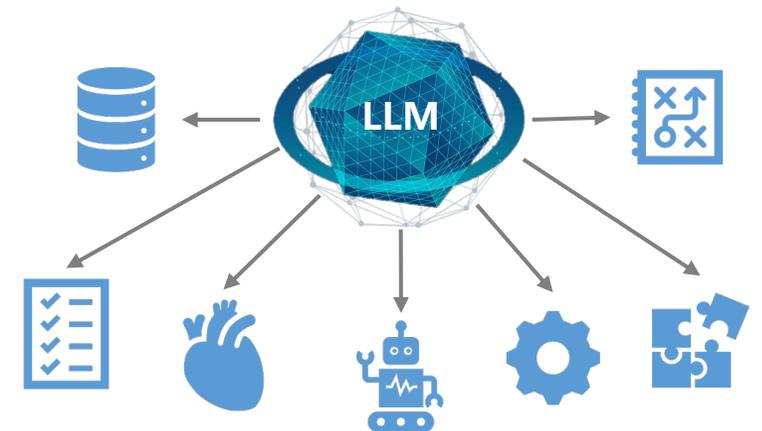
- **Storing personal information:** No maximum input length restriction, retains long-term health records of individuals.
- **Storing medical information:** Maintains patient information, supports automatic follow-up scenarios.



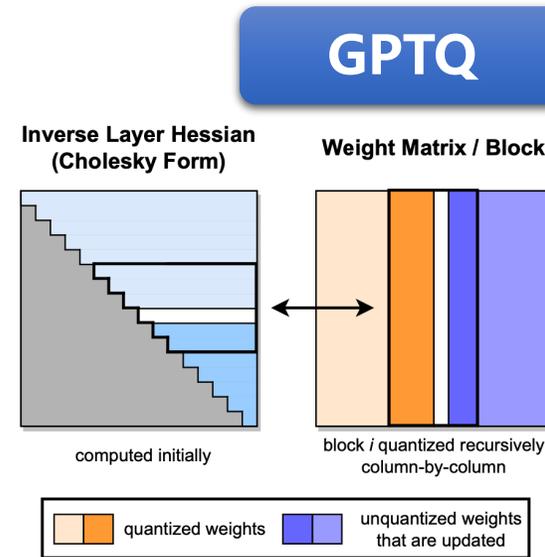
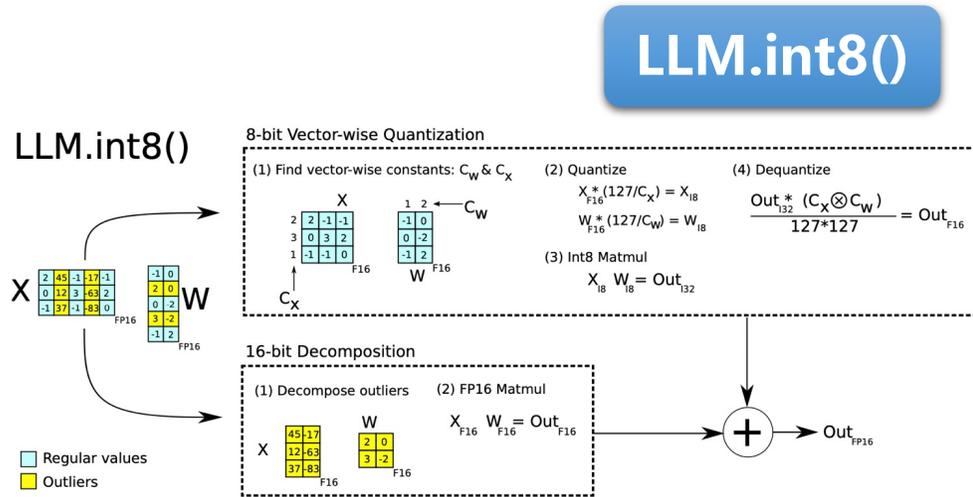
Multi-agent Scheduling

In current complex AI medical diagnostic and treatment processes, there often exist multiple specialized small models, such as data quality control, vessel segmentation, lesion detection, and critical judgment, etc.

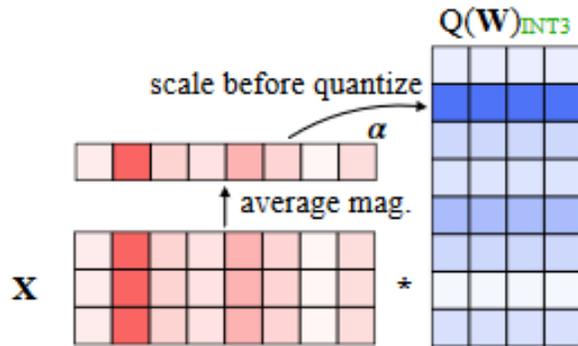
Having LLM invoke its own capabilities to complete simple tasks, and coordinating the scheduling of specialized small models for complex tasks, will make complex reasoning across multiple departments and modalities in the diagnostic and treatment phases possible.



Our Related Work: Large Language Models



AWQ



$$s^* = \arg \min_s \mathcal{L}(s), \quad \mathcal{L}(s) = \|Q(W \cdot s)(s^{-1} \cdot X) - WX\|$$

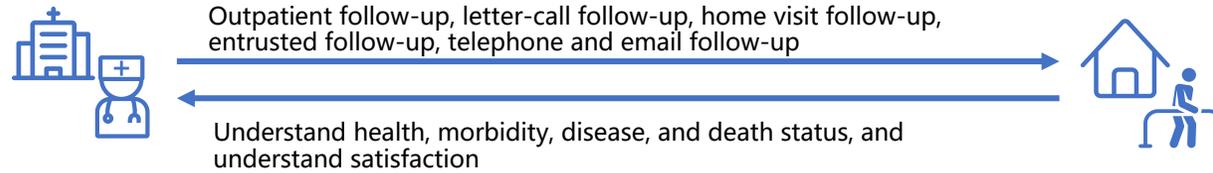
$$s = f(s_X, s_W) = s_X^\alpha \cdot s_W^{-\beta}, \quad \alpha^*, \beta^* = \arg \min_{\alpha, \beta} \mathcal{L}(s_X^\alpha \cdot s_W^{-\beta})$$

Model Size	Min Hardware Requirements Before Quantization	Min Hardware Requirements After Quantization
7B	1 x 3090	1 x 3070
14B	1 x A100	1 x 3090
30B	1 x A100	1 x 3090
100B	3 x A100	1 x A100
176B	5 x A100	2 x A100

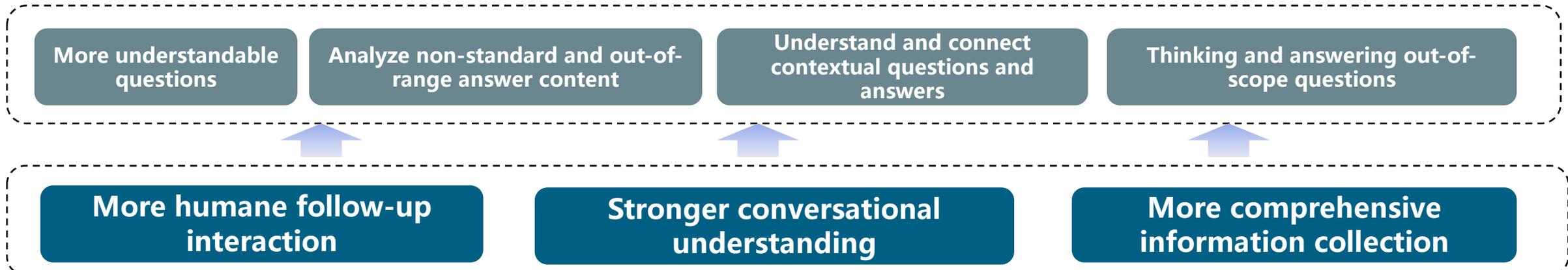
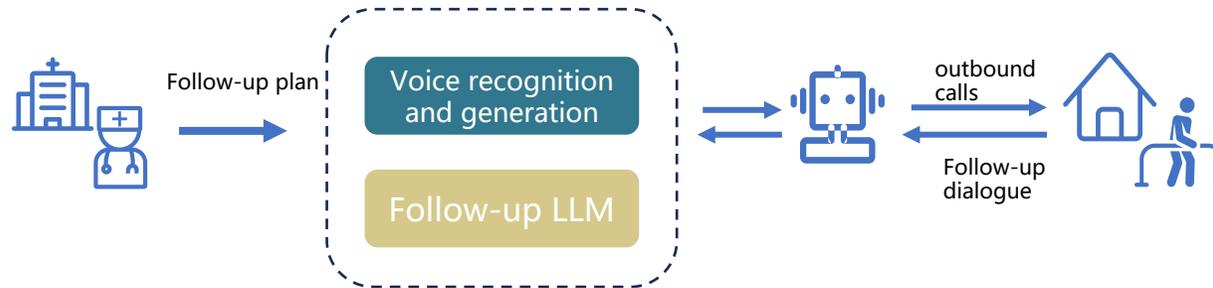
Our Related Work: Large Language Models

Based on LLMs and intelligent voice technology, it realizes functions such as batch outbound calls, human-computer interaction, and automatic statistics, helping hospitals complete follow-up of scientific research, specialized disease and caring work, reducing hospital follow-up costs and improving the work efficiency of medical staff.

Traditional Follow-up pipelines



AI for Follow-up





Core scientific question: Can a general AI model for protein engineering be constructed to directly design protein products that meet application requirements (end-to-end)?

1. First, use the sequence and structure information that already exists in nature for pre-training. Our model: ~600 million parameters, billions of protein training data.



Human Corpus Pre-training
(gpt 3.0: 175 billion parameters, hundreds of billions of training data)
Analogy: Chat GPT

2. Then, use the labeled related data set (such as ocean data) for supervised learning.



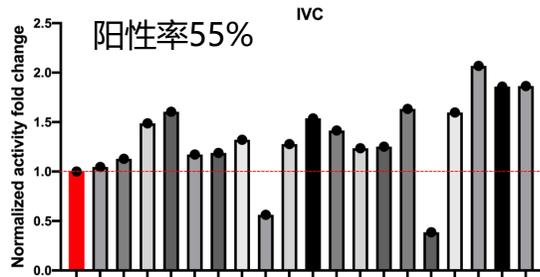
Standard Dataset Optimization
(supervised learning)

3. Finally, use a small amount of wet experimental data (dozens) for model tuning.



Reinforcement Learning for Expert Decision Making

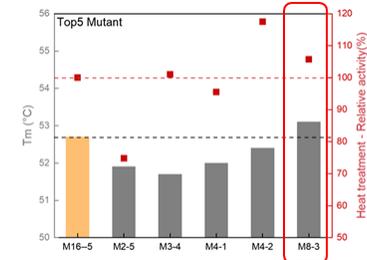
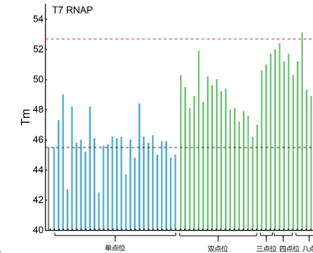
Crispr Case 12 (gene editing proteins, improve activity)



One round of design + experimental iteration, the positive rate of the design is 55%. The editing efficiency of the best mutant is increased by 2 times, achieving the best of its kind.

T7RNAP (RNA drug polymerase, improves thermostability)

Improve high temperature stability and reduce immunogenic impurity products

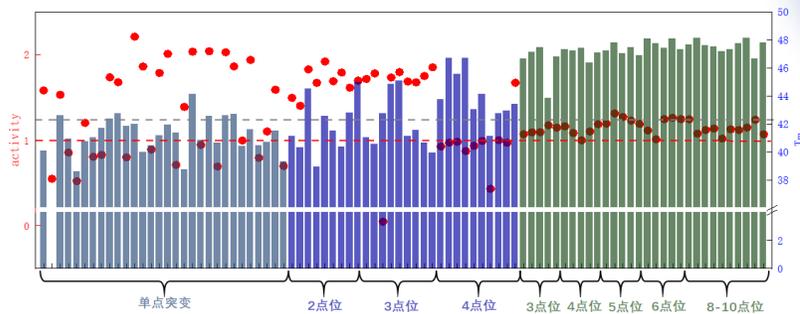


It took three and a half months and two rounds of iterations to develop a multi-site mutant with a melting temperature increased by 7 °C. Its stability and activity are comparable to those of the engineering protein developed by the leading company.

Case 1 Case 2

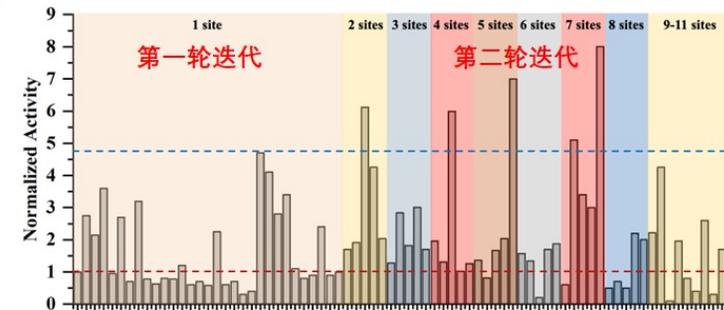
Case 3 Case 4

An IVD enzyme (improves thermostability)



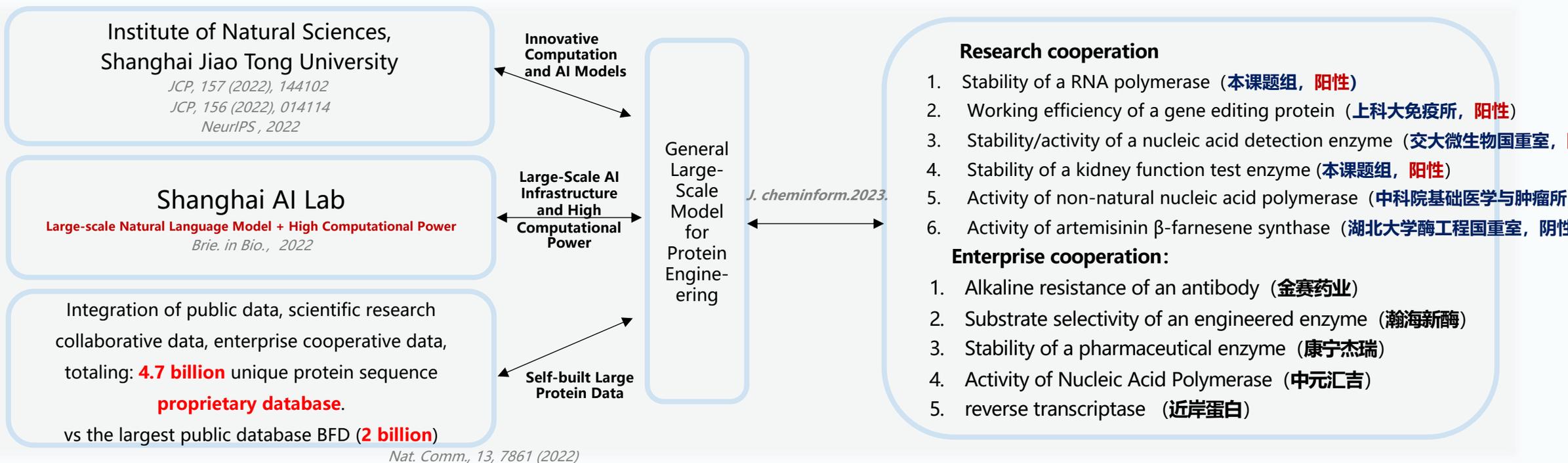
Three-round design + wet experiment, the positive rate of the design (30% in the first round), the second round (60%), and the third round (100%). The melting temperature of the best multi-site mutant was 6°C higher than that of the wild type (WT).

Ago protein (nucleic acid detection/ tumor detection enzyme, improve activity)



Two-round design + wet experiment, the positive rate of the design (first round 40%, second round (90%). The optimal mutant has an 8-fold increase in activity compared to the wild type. (WT).

Our Related Work: Foundation Models for Science



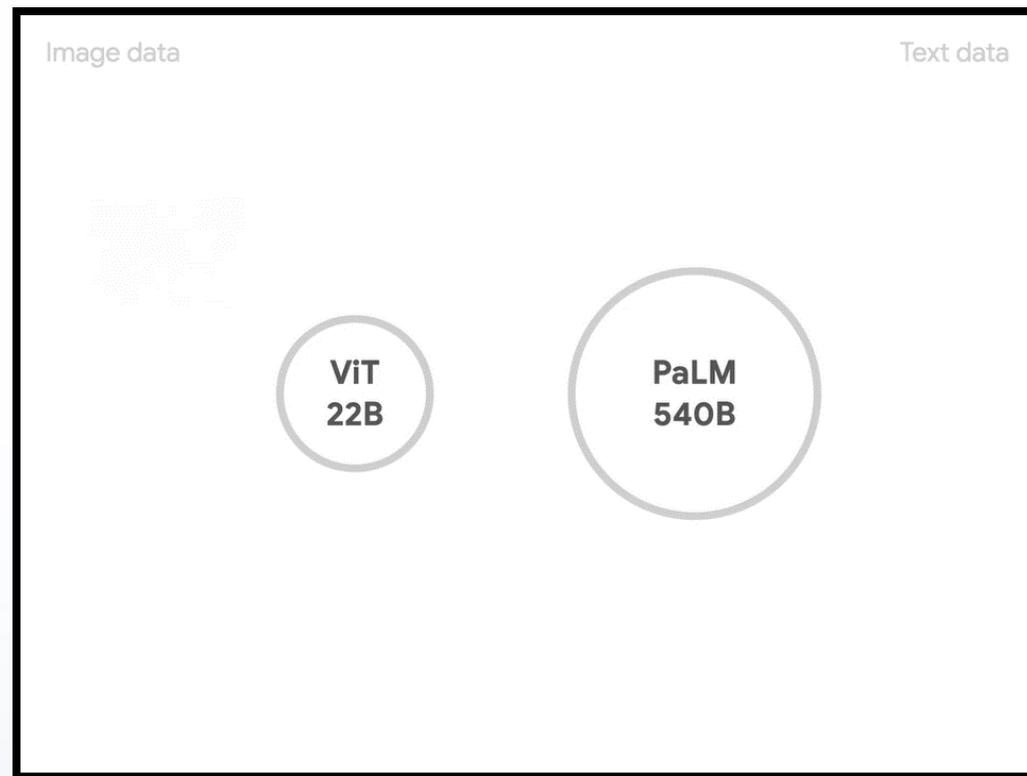
Model type	Large-scale model training in Natural Language Processing			
Development Team	Facebook (USA)		Salesforce Research (USA)	SJTU (Research Group)
Model Name and Parameter Quantity	ESM-1V (650 Million Parameters)	ESM2 (150 Million Parameters)	Progen (XLarge) (6.4 Billion Parameters)	TemPL (650 Million Parameters)
Zero-Shot Model Accuracy (Correlation coefficient between predictions and laboratory results, for 22 protein TM datasets)	0.350	0.384	0.256	0.467

Performance Comparison with Internationally Renowned Teams or Companies

<https://arxiv.org/abs/2304.03780>

The Future Direction of the Medical Foundational Model

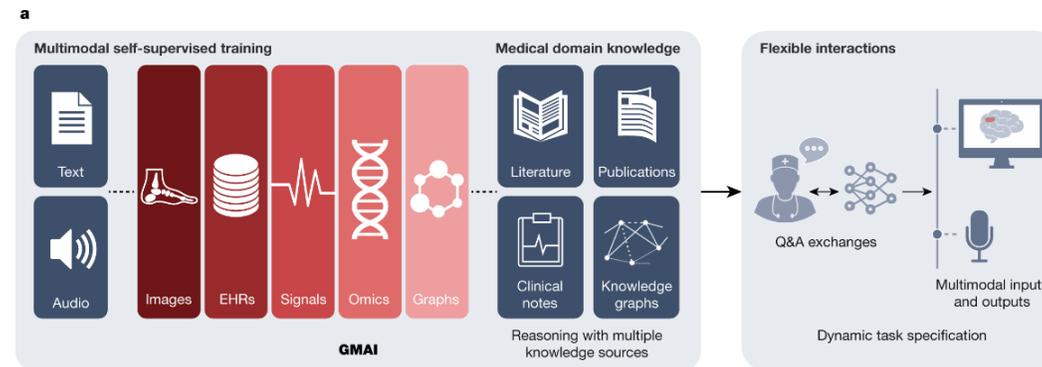
PaLM-E: A concrete multimodal language model.



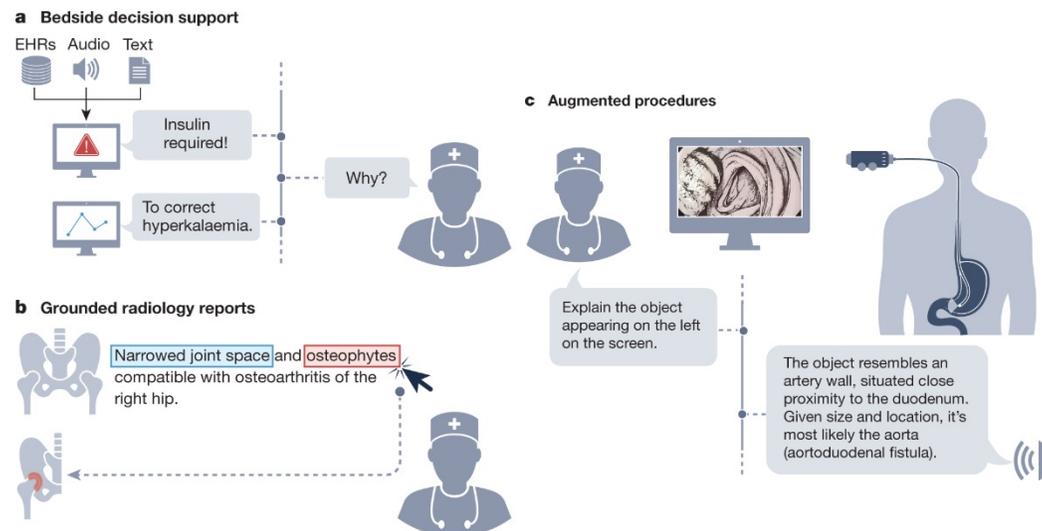
nature

Foundation models for generalist medical artificial intelligence

- Propose a new paradigm for medical AI, which refer to as generalist medical AI (GMAI)
- Be capable of carrying out a diverse set of tasks using very little or no task-specific labelled data.
- Flexibly interpret different combinations of medical modalities, including data from imaging, electronic health records, laboratory results, genomics, graphs or medical text
- In turn produce expressive outputs such as free-text explanations, spoken recommendations or image annotations that demonstrate advanced medical reasoning abilities



Regulations: Application approval; validation; audits; community-based challenges; analyses of biases, fairness and diversity

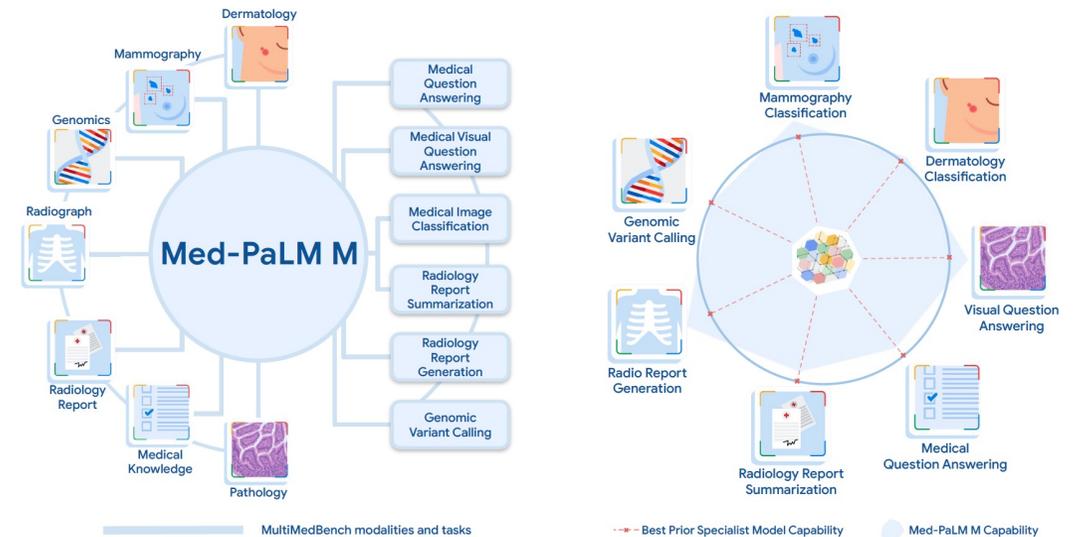


The Future Direction of the Medical Foundational Model



Towards Generalist Biomedical AI

- (1) MultiMedBench, a new multimodal biomedical benchmark, is curated.
- (2) MultiMedBench encompasses 14 diverse tasks such as medical question answering, mammography and dermatology image interpretation, radiology report generation and summarization, and genomic variant calling.
- (3) Then introducing Med-PaLM M, large multimodal generative model that flexibly encodes and interprets biomedical data including clinical language, imaging, and genomics with the same set of model weights.
- (4) In a side-by-side ranking on 246 retrospective chest X-rays, clinicians express a pairwise preference for Med-PaLM M reports over those produced by radiologists in up to 40.50% of cases, suggesting potential clinical utility.



Task Type	Modality	Dataset	Metric	SOTA	PaLM-E (84B)	Med-PaLM M (Best)
Question Answering	Text	MedQA	Accuracy	86.50% [61]	28.83%	69.68%
		MedMCQA	Accuracy	72.30% [61]	33.35%	62.59%
		PubMedQA	Accuracy	81.80% [61]	64.00%	80.00%
Report Summarization	Radiology	MIMIC-III	ROUGE-L	38.70% [62]	3.30%	32.03%
			BLEU	16.20% [62]	0.34%	15.36%
			F1-RadGraph	40.80% [62]	8.00%	34.71%
Visual Question Answering	Radiology	VQA-RAD	BLEU-1	71.03% [63]	59.19%	71.27%
			F1	N/A	38.67%	62.06%
			BLEU-1	78.60% [64]	52.65%	92.7%
	Pathology	Slake-VQA	F1	78.10% [64]	24.53%	89.28%
			BLEU-1	70.30% [64]	54.92%	72.27%
			F1	58.40% [64]	29.68%	62.69%
Report Generation	Chest X-ray	MIMIC-CXR	Micro-F1-14	44.20% [65]	15.40%	53.56%
			Macro-F1-14	30.70% [65]	10.11%	39.83%
			Micro-F1-5	56.70% [66]	5.51%	57.88%
			Macro-F1-5	N/A	4.85%	51.60%
			F1-RadGraph	24.40% [14]	11.66%	26.71%
			BLEU-1	39.48% [65]	19.86%	32.31%
			BLEU-4	13.30% [66]	4.60%	11.50%
			ROUGE-L	29.60% [67]	16.53%	27.49%
			CIDEr-D	49.50% [68]	3.50%	26.17%
Image Classification	Chest X-ray	MIMIC-CXR (5 conditions)	Macro-AUC	81.27% [69]	51.48%	79.09%
			Macro-F1	N/A	7.83%	41.57%
			Macro-AUC	N/A	63.37%	97.27%
	Dermatology	PAD-UFES-20	Macro-F1	N/A	1.38%	84.32%
			Macro-AUC	64.50% [49]	51.49%	71.76%
			Macro-F1	N/A	16.08%	35.70%
Mammography	VinDr-Mammo	Macro-AUC	N/A	47.75%	73.31%	
		Macro-F1	N/A	7.77%	51.12%	
		Macro-AUC	N/A	40.67%	82.22%	
	Genomics	CBIS-DDSM (mass)	Macro-F1	70.71% [70]	11.37%	67.86%
			Macro-F1	99.40% [71]	53.01%	97.04%
			Macro-F1	99.70% [71]	52.84%	99.35%
Genomics	CBIS-DDSM (calcification)	PrecisionFDA (Variant Calling) (Truth Challenge V2)	Indel-F1	99.40% [71]	53.01%	97.04%
			SNP-F1	99.70% [71]	52.84%	99.35%

D. Driess, M. Schaeckermann, A. Karthikesalingam, V. Natarajan, et al. Towards Generalist Biomedical AI. arXiv:2307.14334. 2023

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论文标题	期刊名称
Spatially aware graph neural networks and cross-level molecular profile prediction in colon cancer histopathology: a retrospective multi-cohort study	The Lancet Digital Health
Predicting unseen antibodies' neutralizability via adaptive graph neural networks	Nature Machine Intelligence
Mining Multi-Center Heterogeneous Medical Data with Distributed Synthetic Learning	Nature Communications
Genetic mutation and biological pathway prediction based on whole slide images in breast carcinoma using deep learning	npj Precision Oncology
MedFMC: A Real-world Dataset and Benchmark For Foundation Model Adaptation in Medical Image Classification	Scientific Data
A Large-scale Synthetic Pathological Dataset for Deep Learning-enabled Segmentation of Breast Cancer	Scientific Data

OpenMEDLab




<https://github.com/openmedlab>



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