Revolutionizing Medical Image Data Analysis: Uniting AI and Statistics for Breakthroughs and Challenges

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Thanks to Drs. Mingxia Liu, Xin Wang, Lijuan Liu, Gang Li, Hanchuan Peng, Wei Cheng, Marc Niethammer, Tengfei Li, and Bingxin Zhao for sharing their slides.



https://www.med.unc.edu/big-s2



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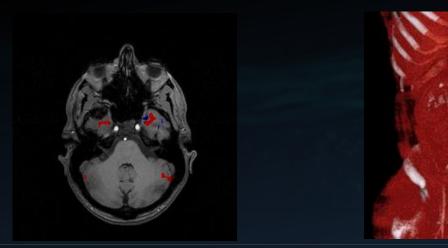
Introduction to Medical Image Data Analysis

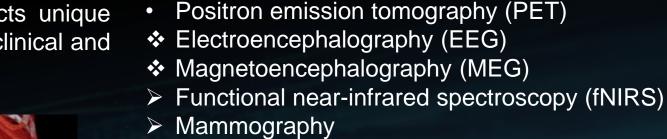


Medical Imaging

Medical imaging is the technique and process used to create images of the human body for clinical purposes or medical science. (<u>https://en.wikipedia.org/</u>)

These imaging methods are essential for delineating the structure and functionality of organs and tissues. Each modality employs a distinct targeting agent, generates data in varying dimensions, extracts unique features, and serves specific purposes within clinical and research contexts.





•

- Light microscopy images
- Fluoroscopy

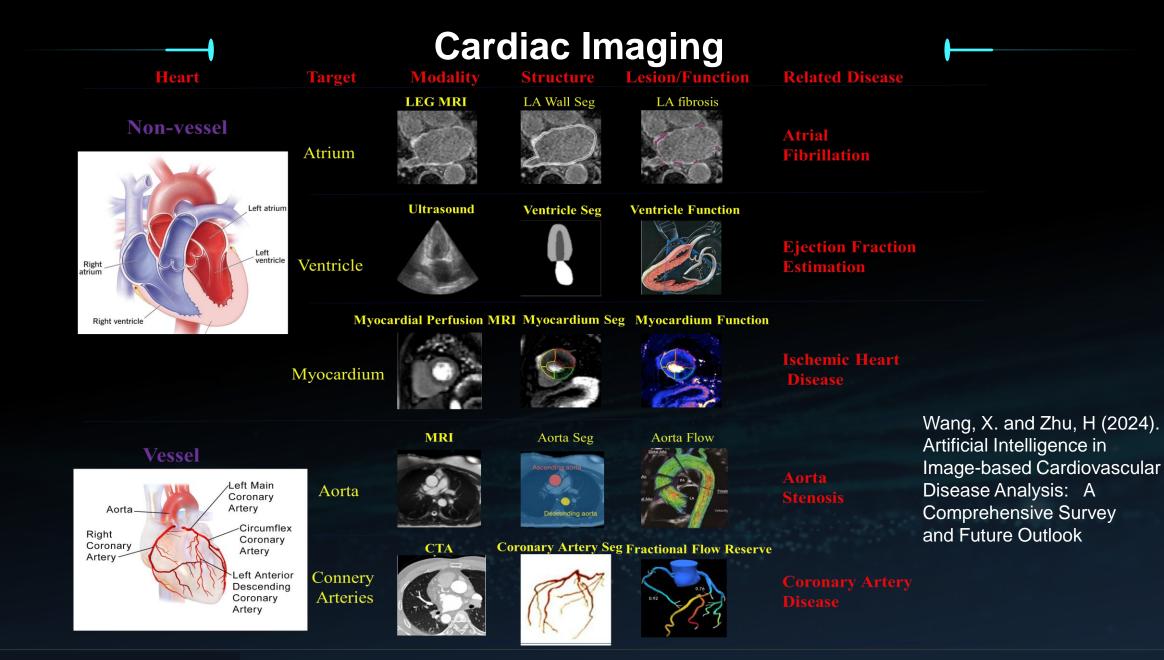
Ultrasound

Echocardiography

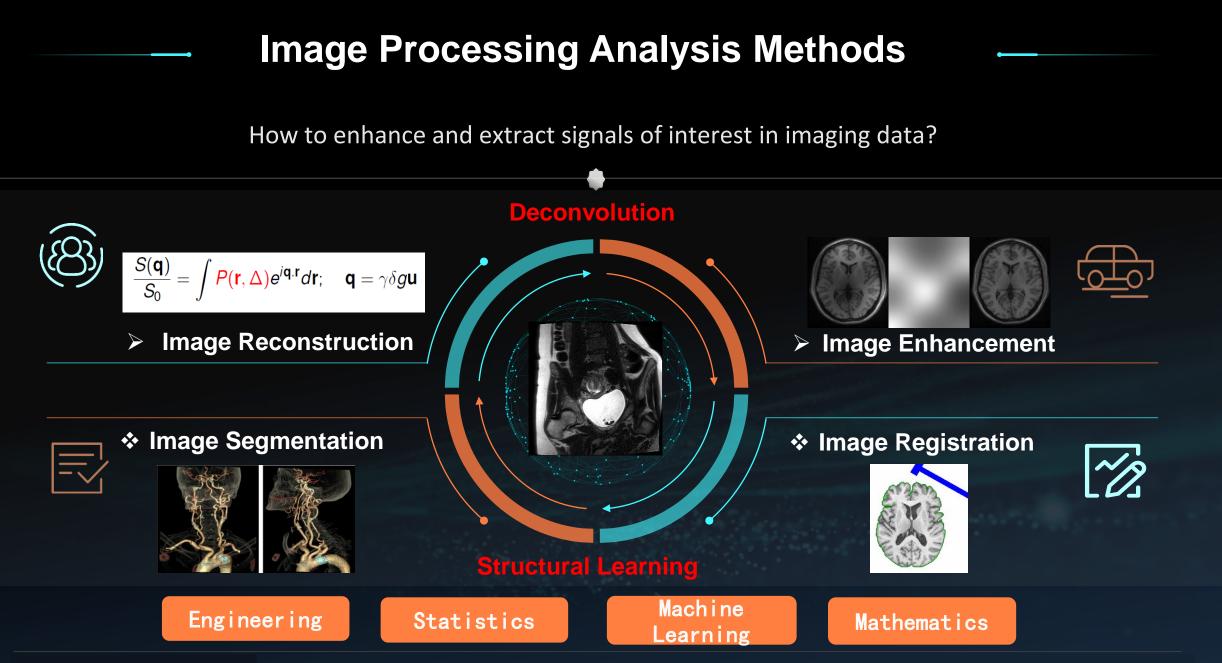
X-ray radiography

Computerized tomography (CT)

Magnetic resonance imaging (MRI)



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Structural Learning

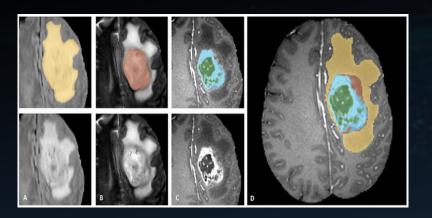
Image Segmentation

- Organ parcellation
- Localization of pathology
- Surgical planning
- Image-guided interventions
- Computer-aided diagnosis
- Quantification of organ change

Image Registration

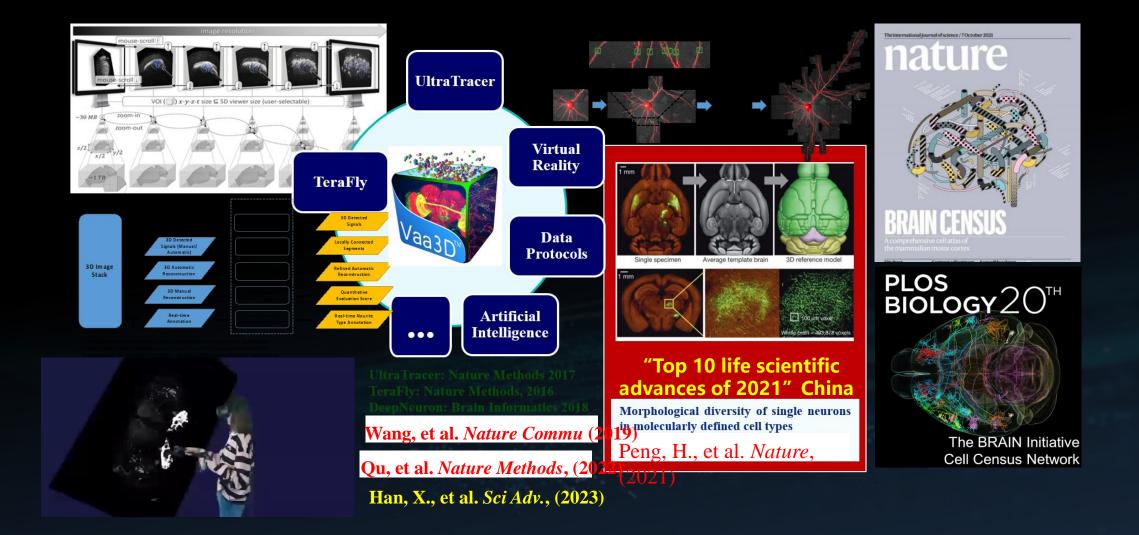
Organ atlas

- Localization of pathology
- Automated image segmentation
- Multimodal fusion
- Population analysis
- Quantification of organ changes





Light Microscopy Imaging at Single Cell



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Ecological Layout for Imaging-based Analysis



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State-of-the-Art AI Applications in Medical Imaging and Statistical Challenges



AI Milestones

Annotated Datasets

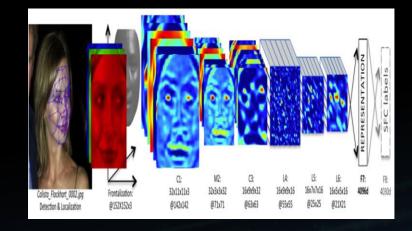
screen television esti: television esti: television



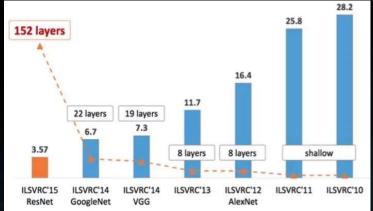




Deep Learning







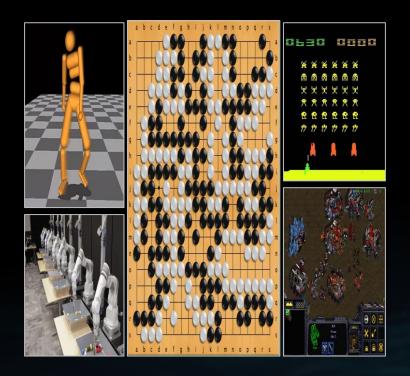
Input	[cLS] my dog is cute	[SEP] he likes play ##ing [SEP]	IJ
Token Embeddings	E _[CLS] E _{my} E _{museq} E _{is} E _{cute}	E _(SEP) E _{he} E _{MMSK} E _{play} E _{rring} E _{(SEP}	P]
Sentence Embedding	+ + + + + E _A E _A E _A E _A E _A	+ + + + + + E _A E _B E _B E _B E _B E _B E _B	
Transformer Positional Embedding	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$)

AI Milestones

Reinforcement Learning

AI Products

PROTEIN



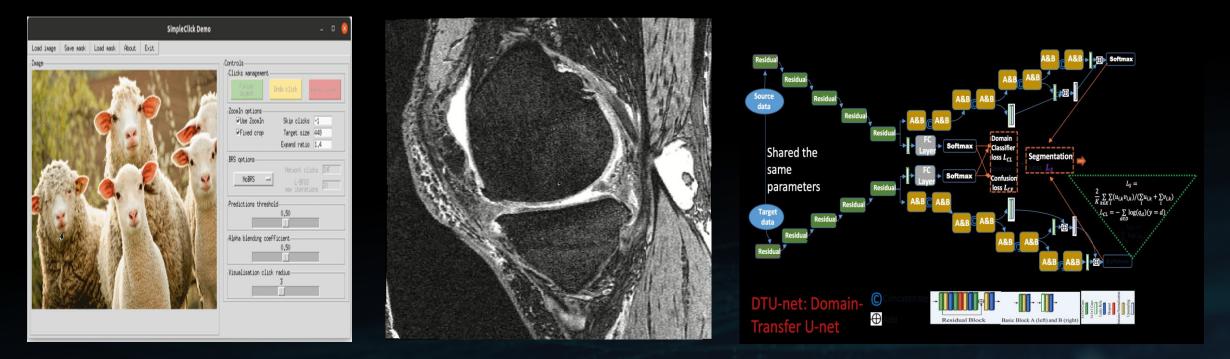




AI for Image Segmentation

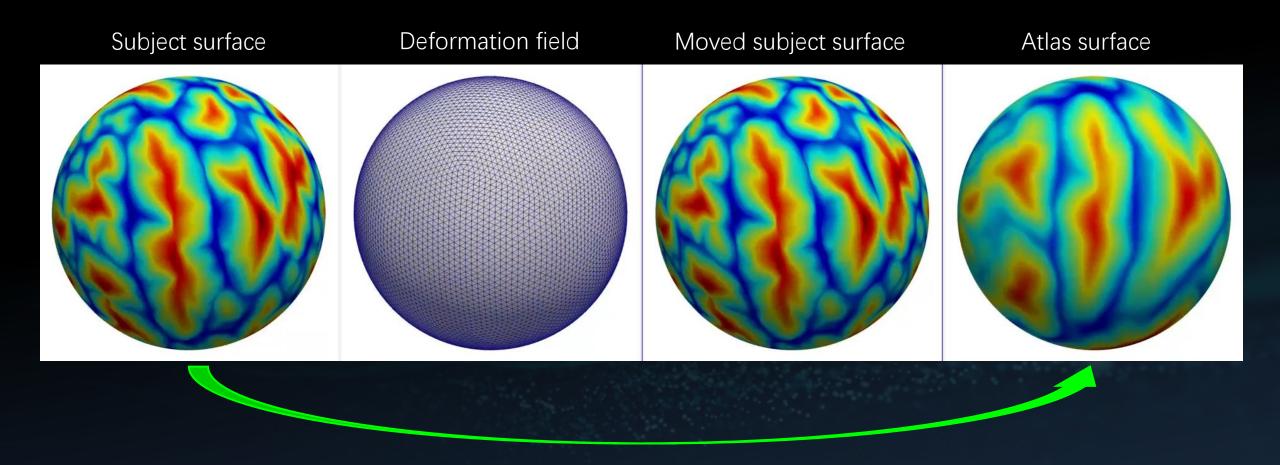
Segmentation Annotation

U-Nets



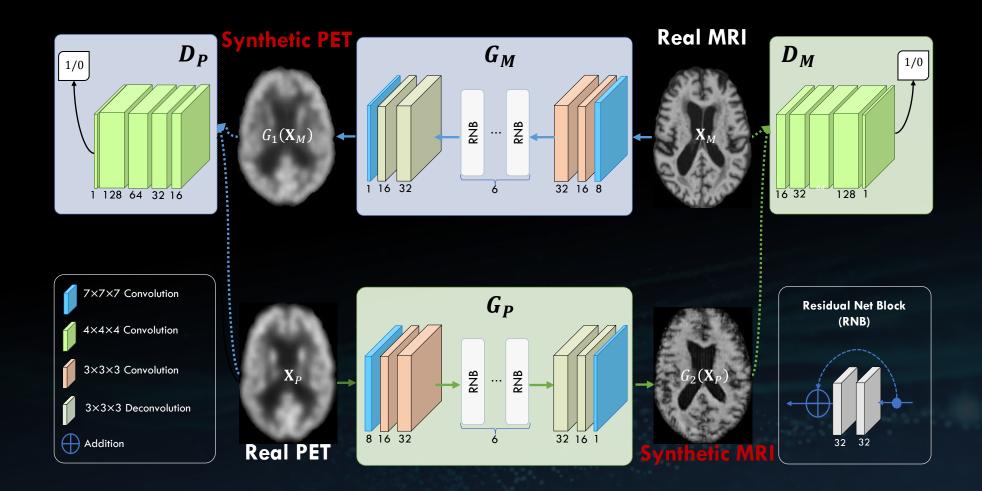
Liu, Q., Xu, Z., Bertasius, G., & Niethammer, M. (2023). SimpleClick: Interactive Image Segmentation with Simple Vision Transformers. ICCV., 22290-22300. 2023. R. Azad *et al.*, "Medical Image Segmentation Review: The success of U-Net." arXiv, Nov. 27, 2022. Minaee, Shervin, et al. "Image segmentation using deep learning: A survey." *IEEE PAMI* 44.7 (2021): 3523-3542.

Superfast Spherical Surface Registration

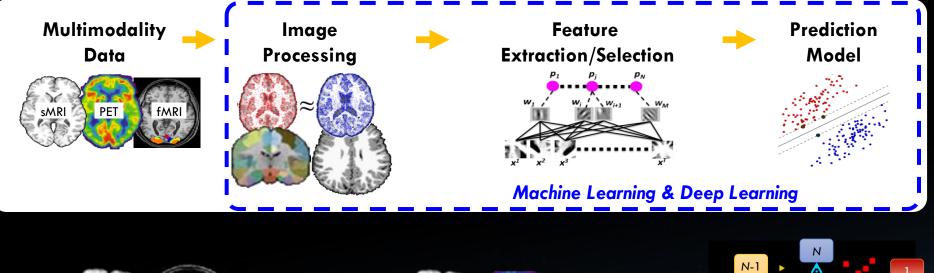


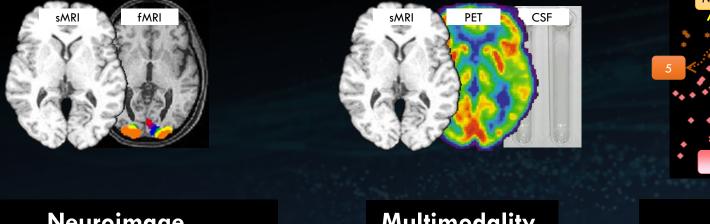
Zhao F, Wu Z, Wang F, Lin W, Xia S, Shen D, Wang L, Li G. S3Reg: Superfast Spherical Surface Registration Based on Deep Learning. IEEE Trans Med Imaging 2021; 40(8): 1964-1976.

Cross-Modality Image Synthesis



Computer-Aided Medical Data Analysis





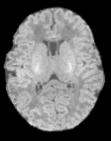
Neuroimage Representation Learning Multimodality Data Fusion Multi-Site Data Adaptation

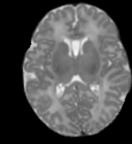
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Major Challenges

Complex Organs and Tissues

Heterogeneity within Individual Subjects and across Centers/Studies

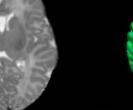


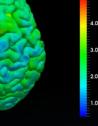


00 Months



00 Months





00 Months

Image=

(age, gene, race, disease, others, device, acquisition, noises

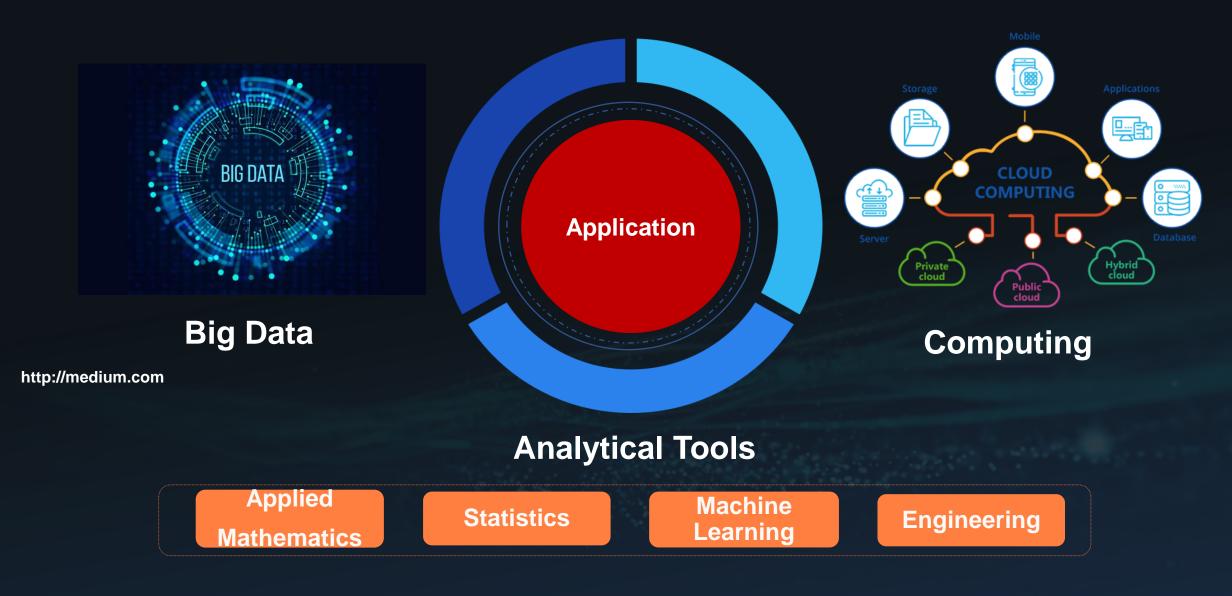


- > There is no publicly available, high-quality imaging datasets with detailed annotation information that cover a large spectrum of segmentation tasks in healthcare.
- How to quantify the uncertainty and generalizarability of atlases as well as deconvolution and structural learning methods and results?
- How to develop RL method for various segmentation and registration tasks?



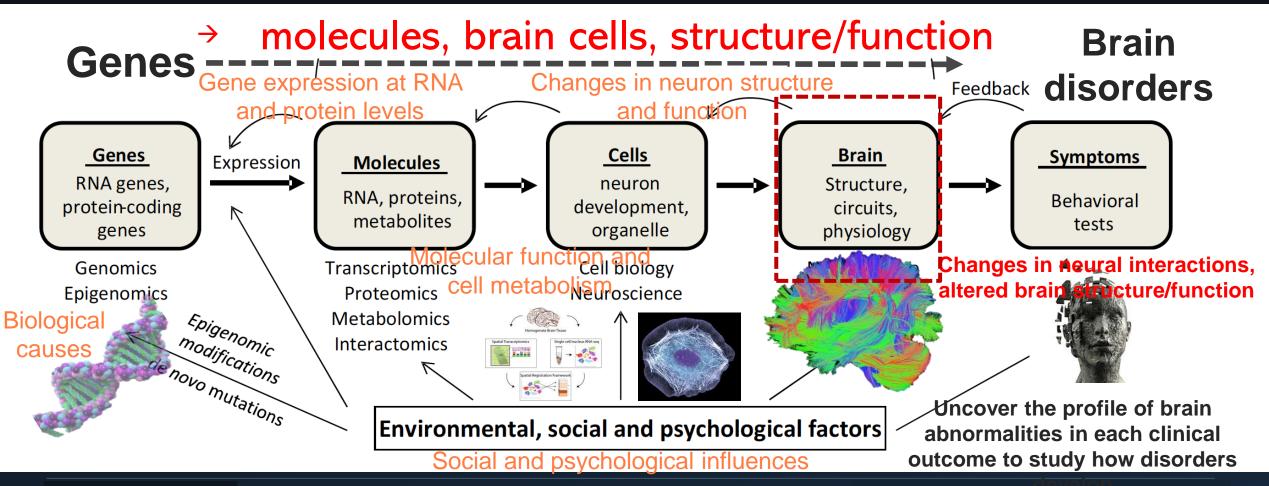
Opportunities for Statisticians in Advancing Medical Imaging Data Analysis

Application to ABC



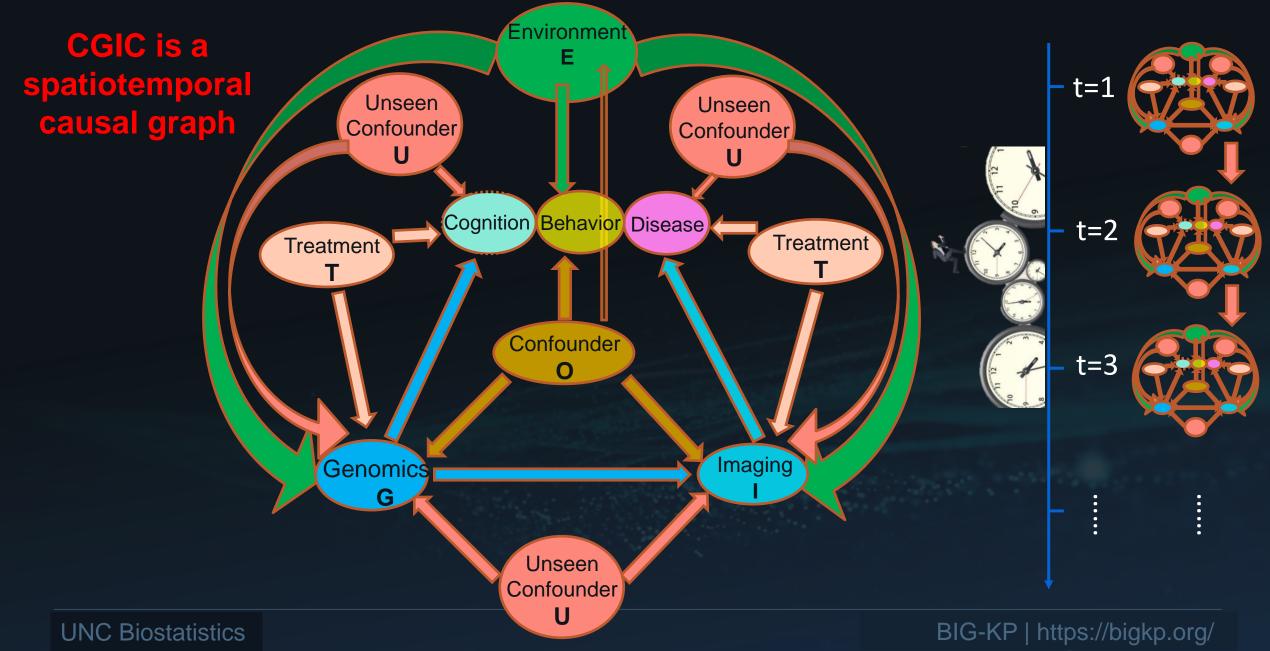
Brain Imaging Genetics Paradigm

Neuroimaging: an important component to help understand the complex biological pathways of brain disorders



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Causal Genetics Imaging Clinical Pathway



Multiple Biobanks/Trials Integration (e.g., Heterogeneity in global populations)

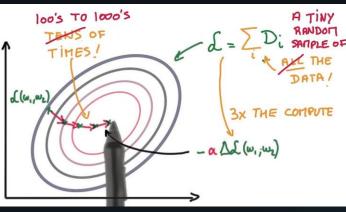


Omics Data Integration (e.g., new tech, biological pathway)

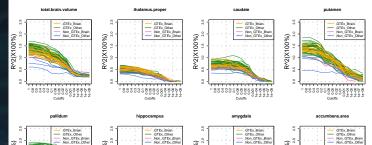
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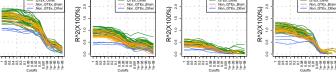
Methodological Challenges





New Computational Tools (e.g., challenge of dense signal in biobank-scale database)





Advanced Methods for Dense Signals (e.g., deep learning)

Important Statistical Topics

- Experimental Design
- Statistical Parametric Mapping
- Object Oriented Data (OOD) Analysis
- Imputation Methods
- Data Integration Methods

Zhu, H., Li, T., & Zhao, B. Statistical learning methods for neuroimaging data analysis with applications. *Annual Review of Biomedical Data Science, Volume 6, Issue 1, 2023.*

- Dimension Reduction Methods
- > Image Genetics
- Causality Research
- Predictive Analysis
- > Knowledge-based Methods
- Reinforcement Learning

Other Important Topics



AD and ADRD Related Datasets

	ADSP	ARIC	ADNI	ADGC	UKB	CHS	FHS	HRS	GR@ACE	NACC-UDS	ROSMAP	MSBB	A4	WRAP	OASIS3
Samples (k)	22.8	15.7	2.2	22.6	500	5.8	10.4	15.7	7.4	43.9	3.6	0.37	6.9	1.7	1.09
AD Cases (k)	11.0	3.2	0.4	11.8	3.1	0.4	0.2	1.1	4.1	17.9	0.8	0.28	-	0.02	0.47
ound mone (n)		-			3.8		-	-		0.9	-	-		-	
AD Candidate (k)					62.1 ¹								1.1 ²		
MCI Cases (k)		1.3	1.0			0.5	0.1		1.5	7.7	0.4	0.05		0.12	
Longitudinal	N	Y	Y	N	Y	Y	Y	Y	N	Y	Y	N	N	Y	Y
Female (%)	61.1	55.0	47.0	59.6	51.6	62.0	52.2	53.6	60.4	57.2	72.7	64.2	57.7	70.2	55.6
Race (%)															
White	72.7	73.0	91.3	92.2	94.4	88.3	100	83.7	100	79.2	92.9	82.0	88.9	83.7	80.9
Black	13.4	27.0	5.0	-	1.5	11.7	-	16.3	-	12.7	5.9	10.3	4.9	11.7	14.9
Other	13.9	-	3.7	7.8	4.1	-	-	-	-	8.1	1.2	7.7	6.2	4.6	4.2
Ethnicity (%)															
Non-Hispanic	83.1	100	95.6	92.2	100	100	100	90.7	100	91.5	94.9	92.8	95.0	97.8	95.8
Age range	32-89	45-84	54-91	60+	44-82	65+	30-62	51-61	65+	36+	54+	61+	65-85	43-90	42-96
Genetic/Omics data			•	•		•	•		•	•			•	•	N
Imaging data (# of subjects)	N	(2k) (0.3k)	(1.7k) (1.3k)	Ν	🔵 (45k)	🔵 (3.6k)	🔵 (5.2k)	N	N	🔵 (6k)	N	Ν	(1.8k) (4.4k)	(k) (0.45k)	(0.91k) (0.78k)
Other variables/ phenotypes	N							A A	N			A		A	
¹ : Subjects with AD-proxy for UKB (if either parent has AD) ² : AD candidates (for A4) with an "elevated" level of amyloid plaque detected from the PET scan															
📕 WES 📙 WGS 🛑 PET scan 🔺 Braak staging 🔺 Functional status 🔺 Echo/Electro-cardiography															
Figure 2 Summary information on 15 studies of the AD-related database (sample size, age, sex, race, data type, etc.)															

Figure 2. Summary information on 15 studies of the AD-related database (sample size, age, sex, race, data type, etc.)

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The UK Biobank Study

UK Biobank has collected and continues to collect extensive environmental, lifestyle, and genetic data on half a million participants.



UK Biobank is a large-scale biomedical database and research resource, containing in-depth genetic and health information from half a million UK participants. The database is regularly augmented with additional data and is globally accessible to approved researchers undertaking vital research into the most common and life-threatening diseases. It is a major contributor to the advancement of modern medicine and treatment and has enabled several scientific discoveries that improve human health.



2006-now



•Imaging: Brain, heart and full body MR imaging, plus full body DEXA scan of the bones and joints and an ultrasound of the carotid arteries. The goal is to image 100,000 participants, and to invite participants back for a repeat scan some years later.

•<u>Genetics</u>: Genotyping, whole exome sequencing & whole genome sequencing for all participants.

•<u>Health linkages</u>: Linkage to a wide range of electronic health-related records, including death, cancer, hospital admissions and primary care records.

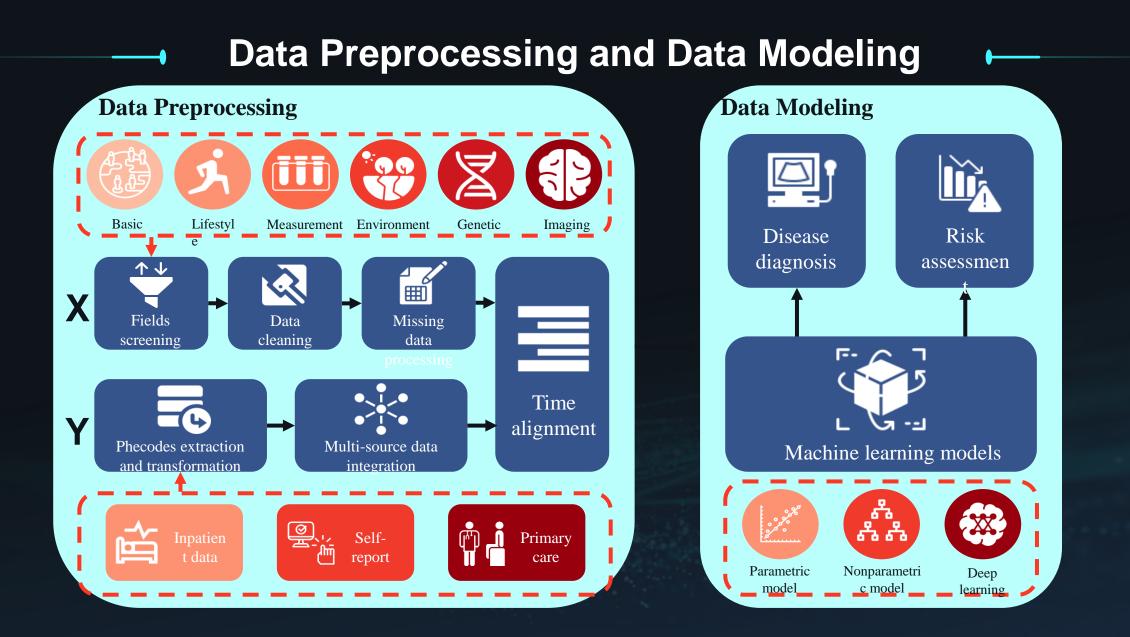
•Biomarkers: Data on more than 30 key biochemistry markers from all participants, taken from samples collected at recruitment and the first repeat assessment.

•<u>Activity monitor</u>: Physical activity data over a 7-day period collected via a wrist-worn activity monitor for 100,000 participants plus a seasonal follow-up on a subset.

•<u>Online questionnaires</u>: Data on a range of exposures and health outcomes that are difficult to assess via routine health records, including diet, food preferences, work history, pain, cognitive function, digestive health and mental health.

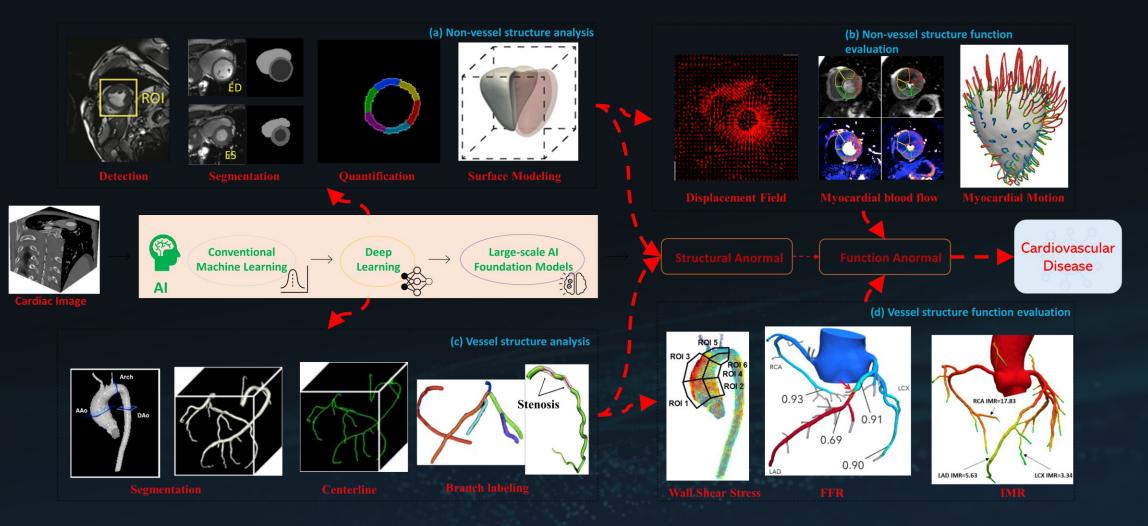
•Repeat baseline assessments: A full baseline assessment is undertaken during the imaging assessment of 100,000 participants.

•<u>Samples</u>: Blood & urine was collected from all participants, and saliva for 100,000.



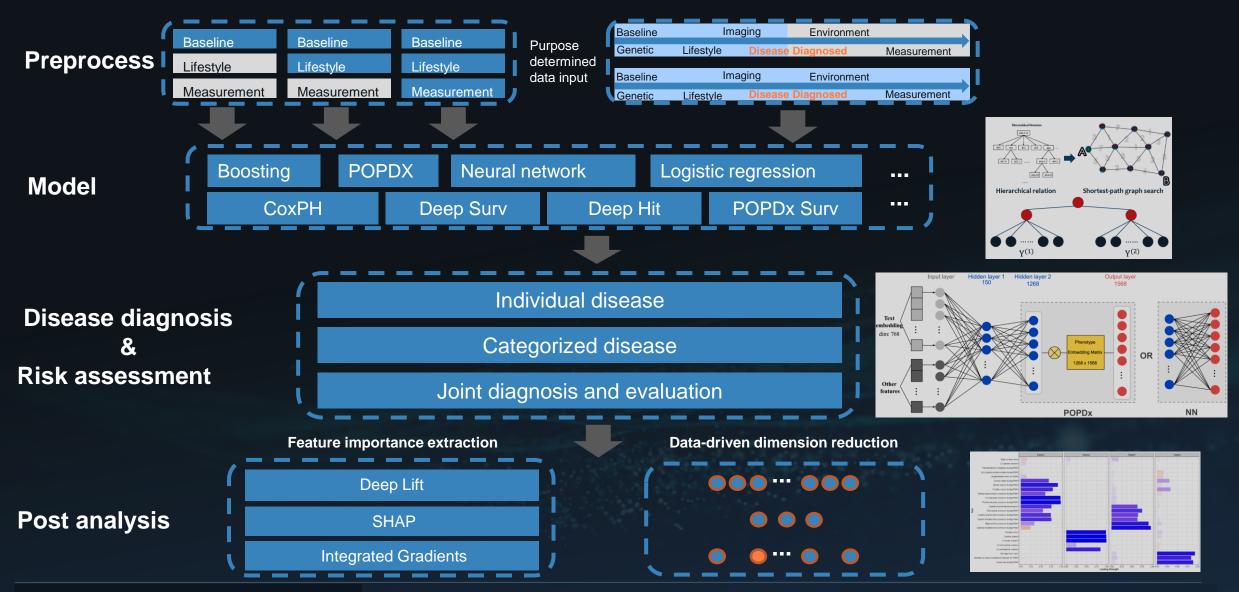
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Image Analysis Pipeline



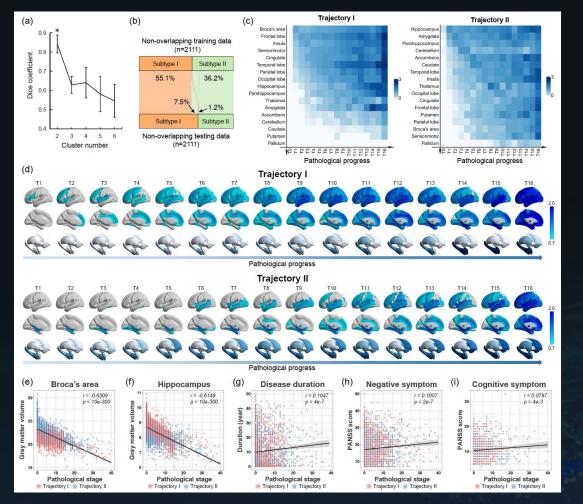
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Prediction Models



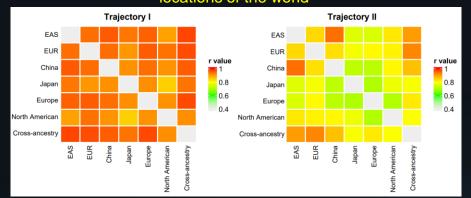
Neuroimaging Biomarkers for Subtypes of Schizophrenia

Two pathophysiological progression trajectories in schizophrenia

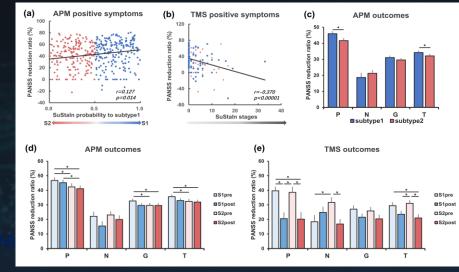


YC Jiang, et al, 2023, Nature Mental Health

Trajectories are reproducibility for samples from different locations of the world



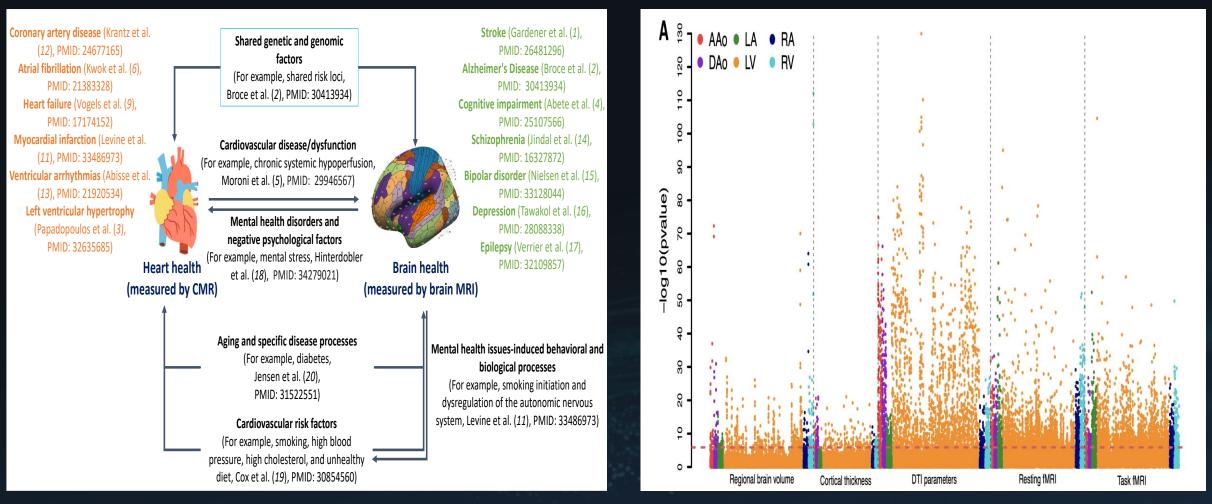
Treatment Outcomes in Subtypes of Schizophrenia



YC Jiang, et al, Nature Communications, Under revision

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Heart-Brain Connections



Zhao, B., Li, T., ..., Stein, J. L., & Zhu, H. Heart-brain connections: Phenotypic and genetic insights from magnetic resonance images. *Science*, 380(6648), abn6598, 2023.

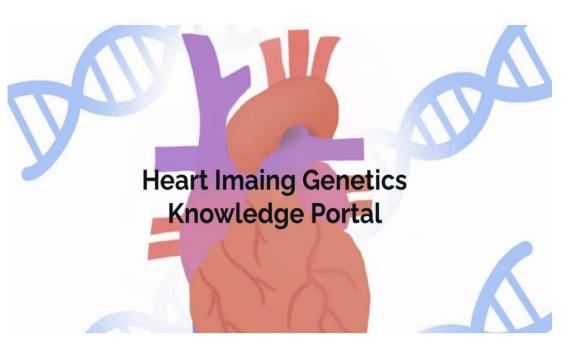
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Brain- Heart Imaging Genetics Knowledge Portal

Brain Imaging Genetics Knowledge Portal (BIG-KP)

Genetics Discoveries in Human Brain by Big Data Integration



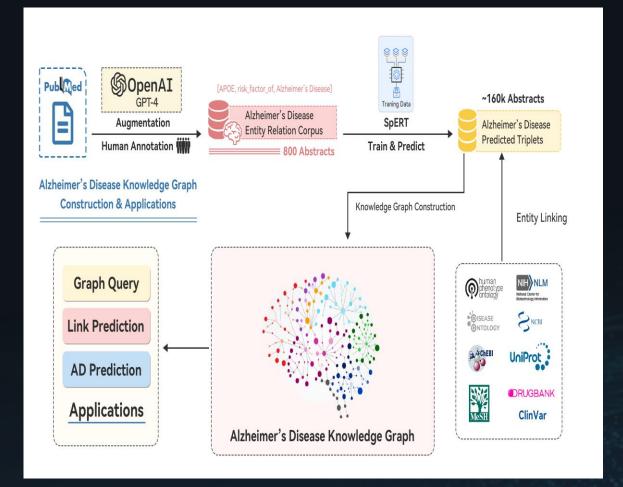


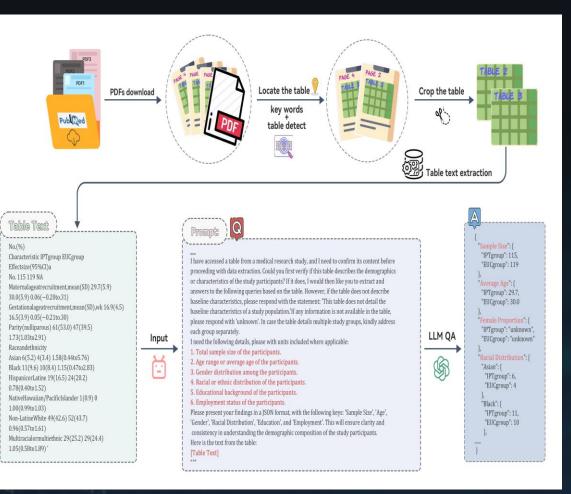
Brain Imaging Genetics Knowledge Portal

Heart Imaging Genetics Knowledge Portal

(<u>BIG-KP</u>) Aim to build the best knowledge database of neuroimaging genetics

Knowledge Graph Construction





Yang et al., Alzheimer's Disease Knowledge Graph Enhances Knowledge Discovery and Disease Prediction.

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Foundation Models for GMAI

Perspective Multimodal self-supervised training Medical domain knowledge **Flexible interactions** \equiv Publication Literature Q&A exchanges Multimodal inputs Clinical Knowledg FHR and outputs notes araphs Dynamic task specification Reasoning with multiple GMAI knowledge sources Applications Interactive Chatbots for Augmented Grounded Text-to-protein Bedside decision patients note-taking procedures radiology reports generation support

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

Fig. 1 | **Overview of a GMAI model pipeline. a**, A GMAI model is trained on multiple medical data modalities, through techniques such as self-supervised learning. To enable flexible interactions, data modalities such as images or data from EHRs can be paired with language, either in the form of text or speech data. Next, the GMAI model needs to access various sources of medical knowledge to carry out medical reasoning tasks, unlocking a wealth of capabilities that can be used in downstream applications. The resulting GMAI model then carries

out tasks that the user can specify in real time. For this, the GMAI model can retrieve contextual information from sources such as knowledge graphs or databases, leveraging formal medical knowledge to reason about previously unseen tasks. **b**, The GMAI model builds the foundation for numerous applications across clinical disciplines, each requiring careful validation and regulatory assessment.

Moor, M., ..., Rajpurkar, P. (2023) Foundation models for generalist medical artificial intelligence. *Nature*.

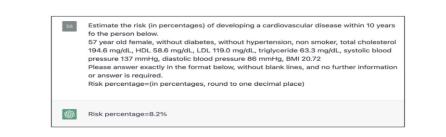


Fig. 2 | Example of a ChatGPT prompt and response for risk stratification. Tabular data extracted from the UK biobank and KoGES were organized and queried into a sentence format like the example above. The 10-year CVD risk percentage was extracted using regular expressions from the corresponding answers.

nedRxiv preprint doi: https://doi.org/10.1101/2023.05.22.23289842; this version posted May 24, 2023. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted medRxiv a license to display the preprint in perpetuity. It is made available under a C-618/-ND 4.0 International license.

	Accuracy	Sensitivity	Specificity	PPV	NPV	F1 score
UK biobank						
GPT-4	0.834	0.393	0.849	0.084	0.975	0.138
GPT-3-5	0.674	0.598	0.677	0.061	0.980	0.111
Bard	0.702	0.447	0.711	0.052	0.973	0.093
Framingha m	0.773	0.508	0.782	0.076	0-978	0.132
KoGES						
GPT-4	0.902	0.153	0.926	0.062	0.972	0.088
GPT-3-5	0.836	0.273	0.854	0.056	0.974	0.093
Bard	0.779	0.307	0.794	0.045	0.973	0.079
Framingha m	0 .874	0.278	0.893	0.077	0.975	0.120

Table 2 | Performance comparison of Framingham, Bard, and ChatGPT Risk Score

PPV: positive predictive value, NPV: negative predictive value. Bold font indicates the highest value of the corresponding metric.

Han,C.,, Yoon, D. (2023) Large-language-model-based 10-year risk prediction of cardiovascular disease: insight from the UK biobank data. *medRxiv*

Acknowledgement

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Brain Imaging Genetics Knowledge Portal (BIG-KP) Genetics Discoveries in Human Brain by Big Data Integration bigkp.org

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Pictures: Copyrights belong to their own authors and/or holders.
Data: We thank Bingxin Zhao, Tengfei Li and other members of the UNC BIG-S2 lab
(https://med.unc.edu/bigs2/) for processing the neuroimaging data.
UK Biobank resource application number: 22783.