University of Connecticut Economic Development Forum 2025



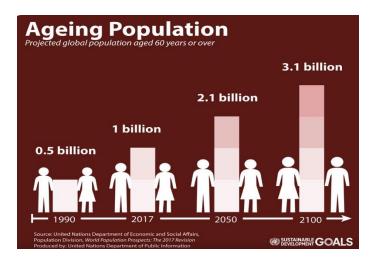
Agentic Generative AI for Precision Medicine

Martin Renqiang Min NEC Laboratories America Nov 13, 2025

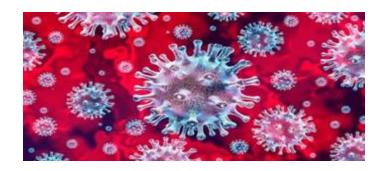


Modern Healthcare Needs Transformation

A growing elderly population, driven by increases in the expected human lifespan, will pressure the existing medical infrastructure.



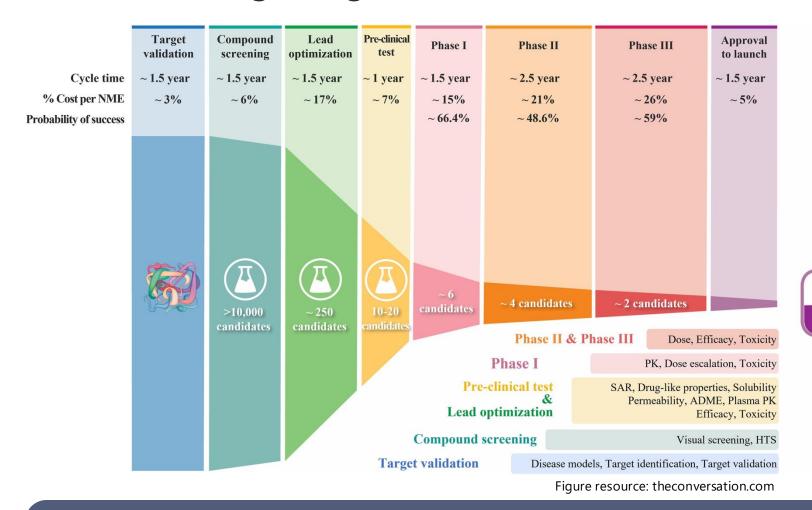
Widespread and low-cost world travel facilitates the quickly and easily spread of disease between countries, increasing the potential of frequent global pandemics.



Medical knowledge explosion will require an ever-increasing number of medical experts to manage and deliver life saving treatments.



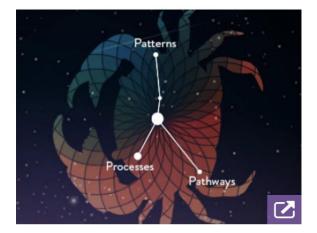
Traditional Drug Design Business is Not Sustainable



R&D cost, but still low success rate (<10%)

Average time/cost for designing one new drug = >10 years + \$~2B

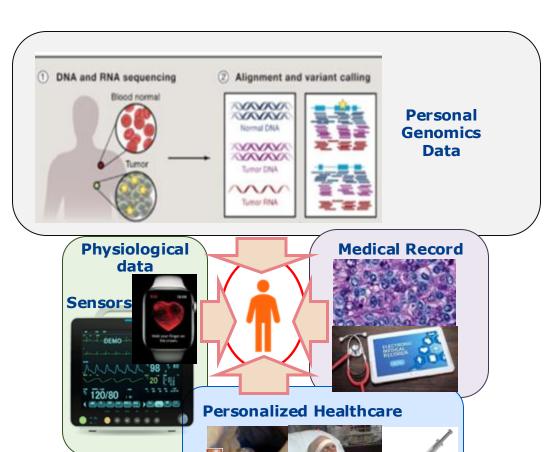
Large-Scale Biomedical Data Enables Precision Medicine and Healthcare

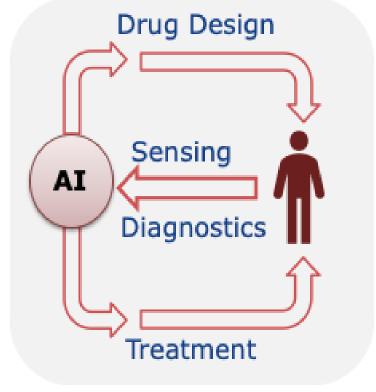


TCGA's Pan-Cancer Atlas



TCGA Outcomes & Impact





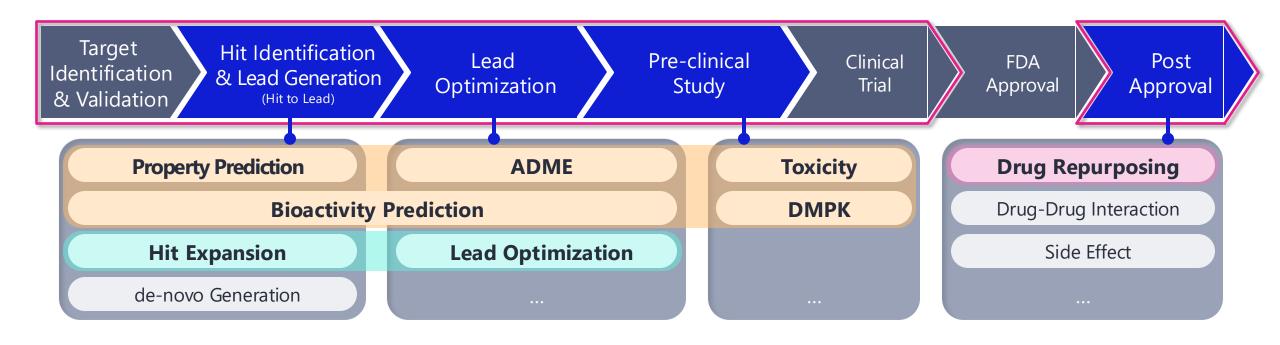
Al Accelerated Drug Development Process

Advances in **AI and high-throughput experiments** empower **accelerated drug development**

FDA NEWS RELEASE

FDA Announces Plan to Phase Out Animal Testing Requirement for Monoclonal Antibodies and Other Drugs

For Immediate Release: April 10, 2025

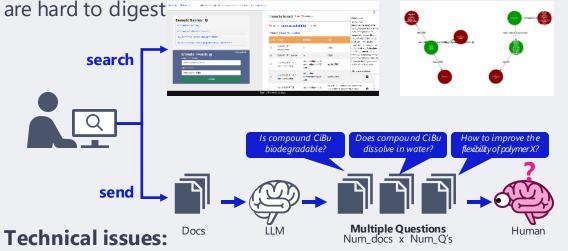




Human Collaborative Multi-Agents for Drug Screening

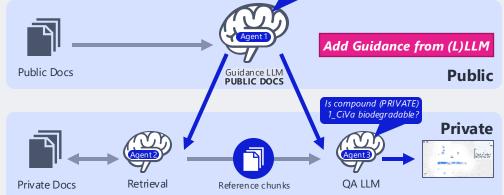
Enable interactive privacy-preserving insight discovery from multimodal data to significantly reduce drug investigation, screening, and design time

Traditional: Scientists manually search thousands of research papers to investigate/screen drug candidates or send a lot of private data to public LLMs to get lengthy textual answers that



Reading many literature papers and reports is time-consuming; Sending private company data to public LLMs poses data-leaking and privacy issues.

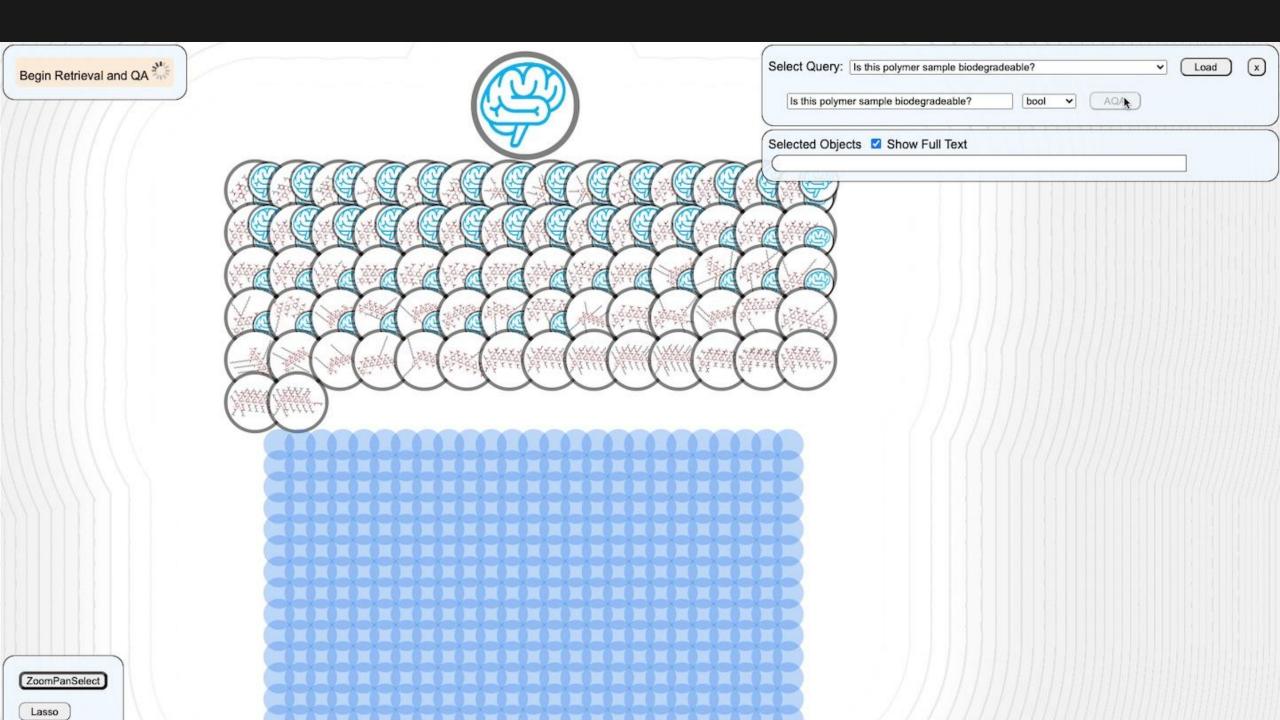
NEC: Human collaborative Al agents for interactively visualizing large-scale existing drugs, filtering candidates, answering questions, and generating manufacturing methods to increase efficiency in all phases of drug design.

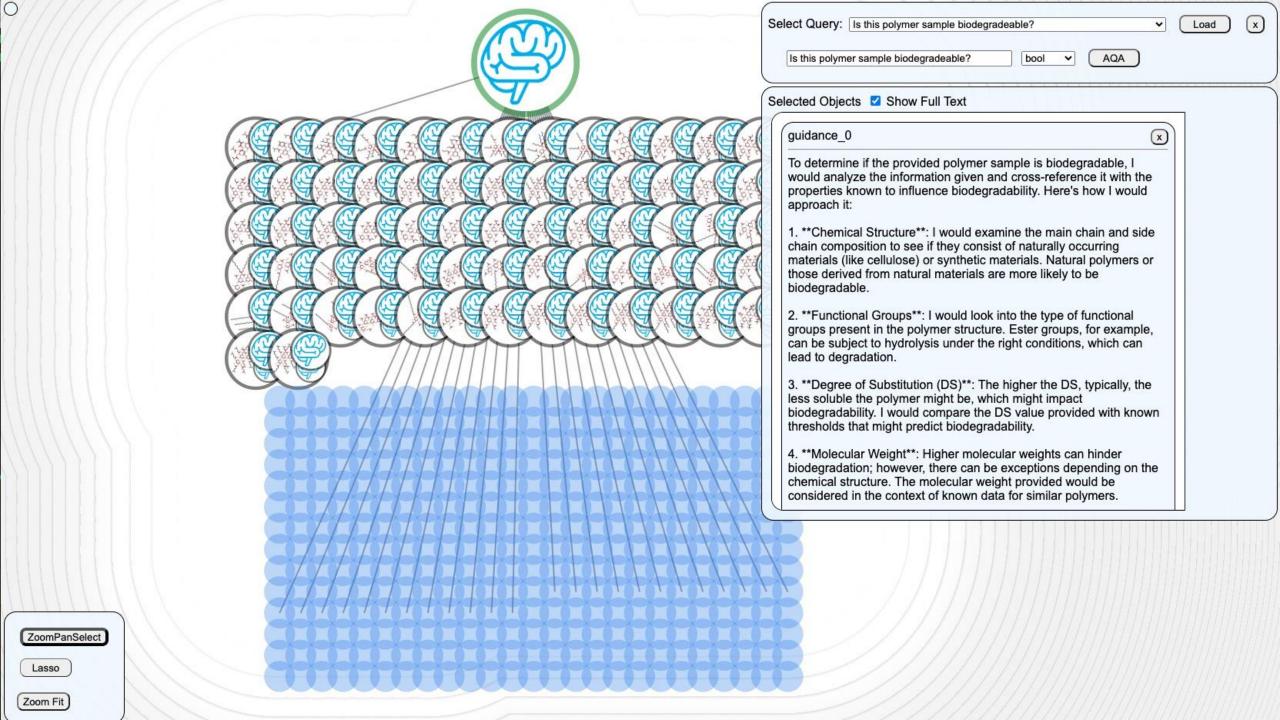


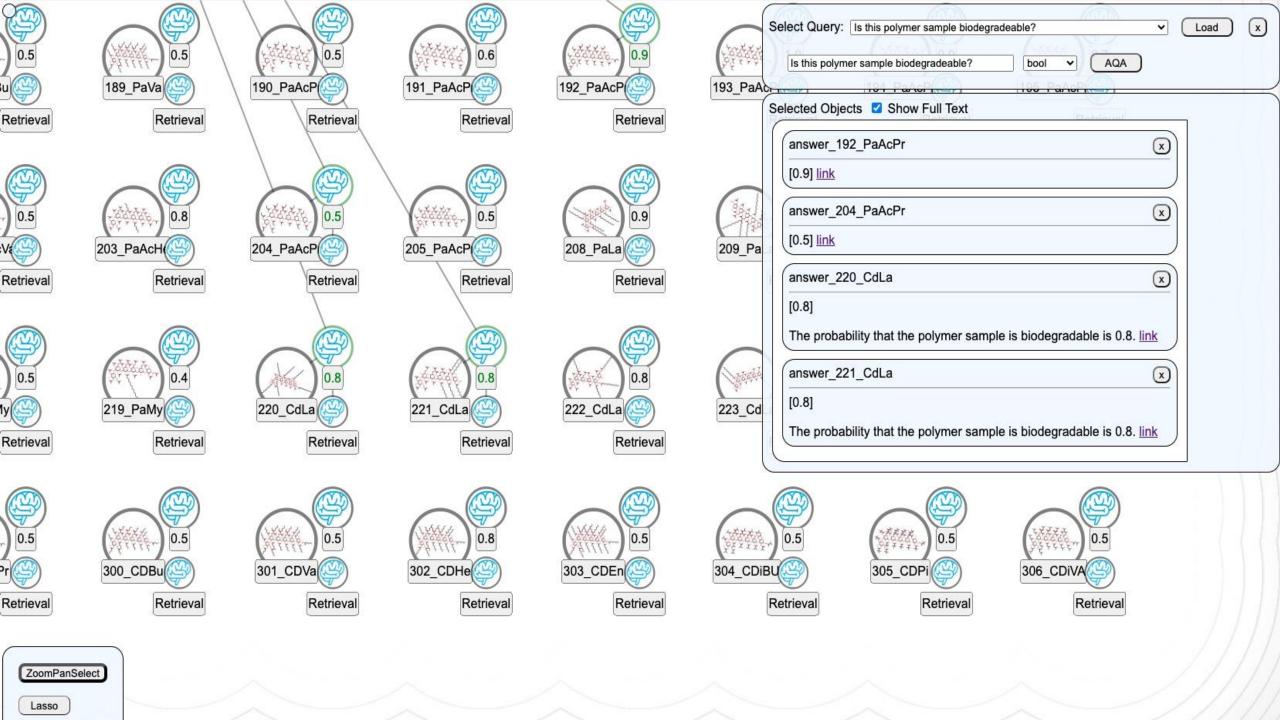
Efficient and safe drug insight discovery:

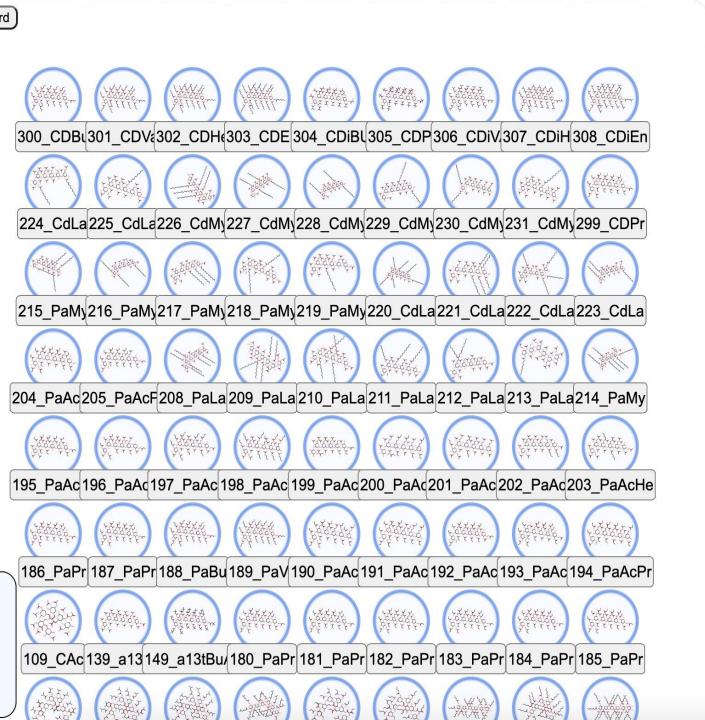
LLM-enhanced interactive data visualization and QA increases the efficiency of insight discovery for drug design; Local agents enable data safety.



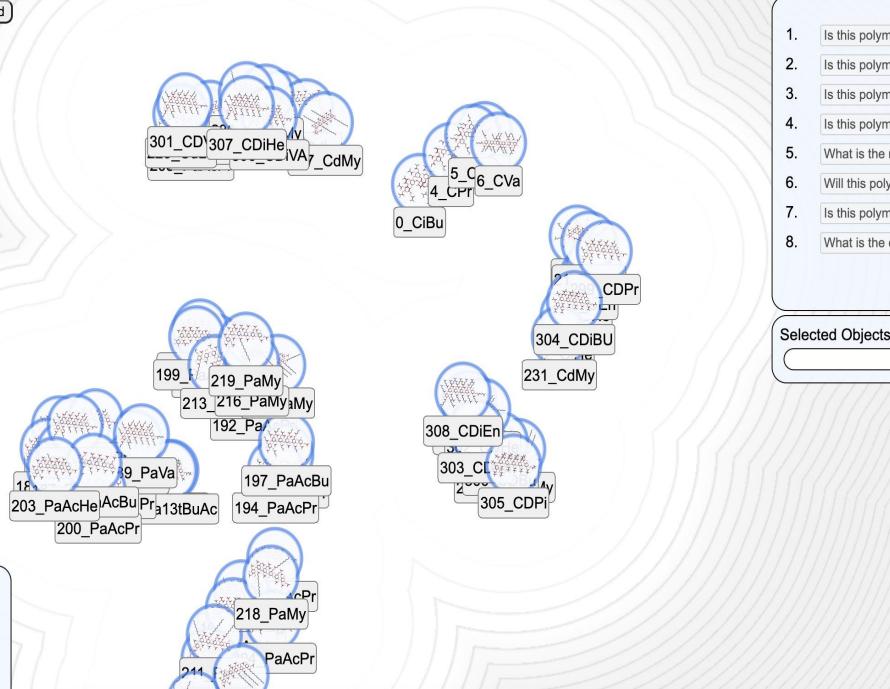




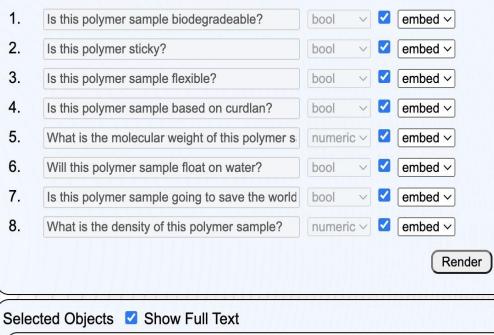


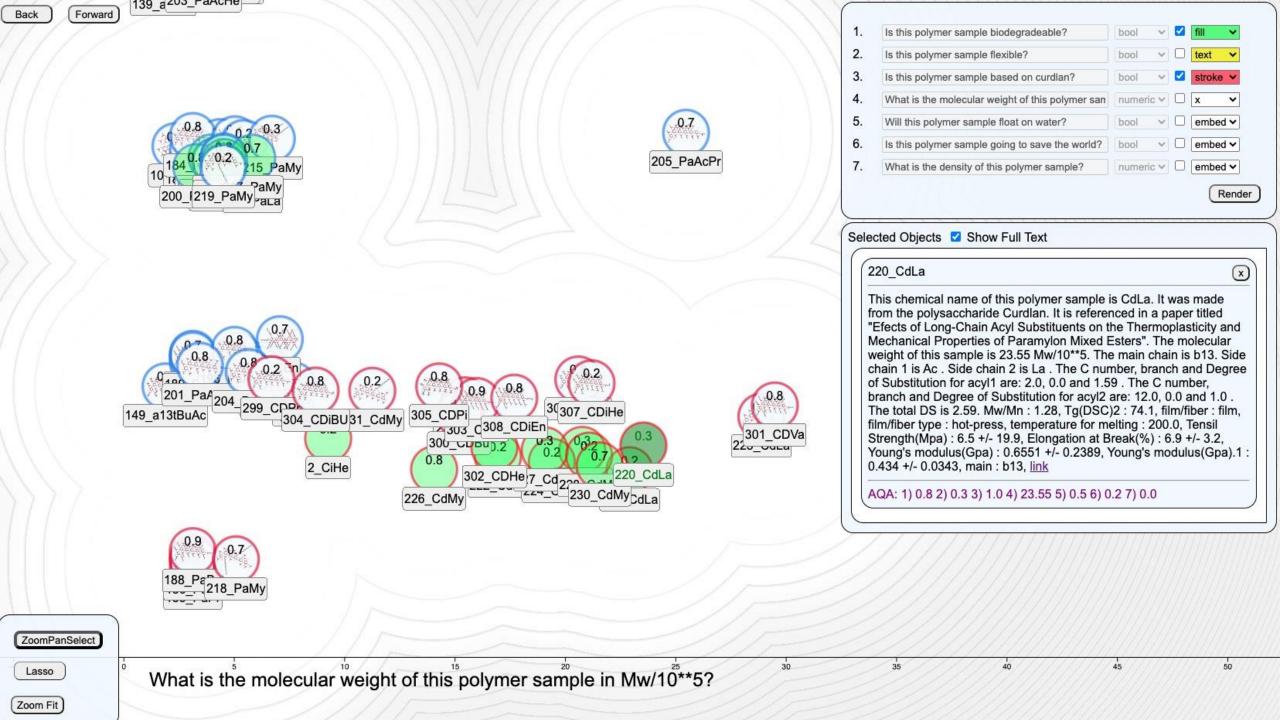


Is this polymer sample biodegradeable?	bool	~	\checkmark	embed ~
Is this polymer sticky?	bool	~	~	embed ~
Is this polymer sample flexible?	bool	~	~	embed >
Is this polymer sample based on curdlan?	bool	~	~	embed >
What is the molecular weight of this polymer s	numer	ic ×	~	embed ~
Will this polymer sample float on water?	bool	~	~	embed ~
Is this polymer sample going to save the world	bool	~	~	embed ~
What is the density of this polymer sample?	numer	ic v	V	embed ~



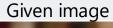
217 Dally





Compositional Generation with Diffusion Probabilistic Models

Compositionally manipulate multiple attributes in generation

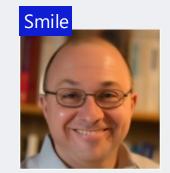








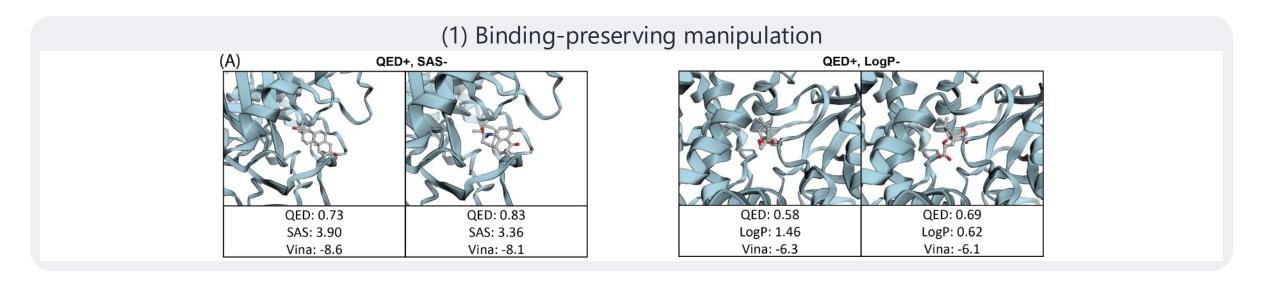


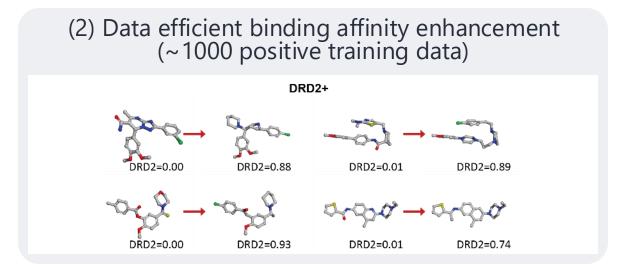


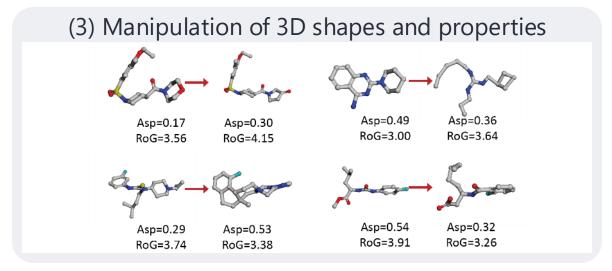


Shi et al., Exploring Compositional Visual Generation with Latent Classifier Guidance. CVPR Workshop 2023.

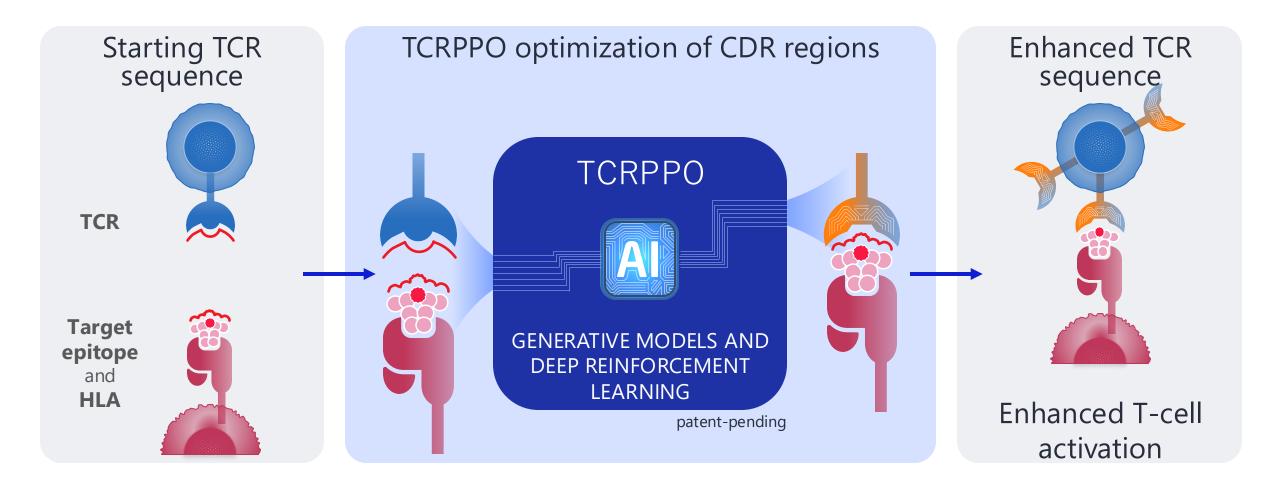
Efficient Multi-Objective Manipulation of 3D Molecules







Generative Models and RL for TCR Engineering



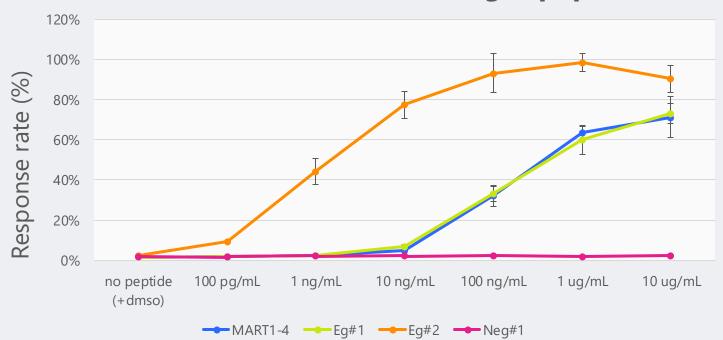
Chen et al., T-Cell Receptor Optimization with Reinforcement Learning and Mutation Polices for Precision Immunotherapy. RECOMB 2023.

Min et al., Design of enhanced TCR against cancer antigens using an Al system. SITC 2024.

In Vitro Validation Results

Mutate existing TCRs for better immunotherapy with reinforcement learning

Cell activation after stimulation with different concentration of target peptide



From over 6.4 billion mutated TCR candidates, we selected optimized TCRs targeting the melanoma antigen MART-1 for wet-lab validation, one engineered TCR is over 1000 times more sensitive than the given template.



PPDiff: Diffusing in Hybrid Sequence-Structure Space for **Protein-Protein Complex Design** Code Link

SSINC Network

 $p_{\theta}(\mathcal{B}_{t-1}|\mathcal{B}_t; \mathcal{T})$

 $q(\mathcal{B}_t | \mathcal{B}_{t-1}; \mathcal{T})$

Pretraining

target-protein mini-

binder complex design

Causal Attention Laver



Carnegie Mellon University School of Computer Science

Zhenqiao Song, Tianxiao Li, Lei Li, Martin Renqiang Min

kNN Equivariant Graph Layer Self-Attention Layer



Protein-Protein Complex Design



Given a target protein with its amino acid sequence and the alpha-carbon backbone structure, the goal is to designing proteins that bind to the target

Core idea

- Conditioning on target protein
- Co-designing binder protein sequence and backbone structure
- Hybrid (discrete-continuous) diffusion

General Protein-Protein Complex Design

PPDiff

Generative Process

antigen-antibody

complex design

Finetuning

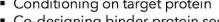
PPDiff

B denotes binder protein

- x: alpha-carbon based backbone structure
- s: amino acid sequence

 L_{attn} : 11 L_{I} : 3 L_{causal} : 1

- PPBench
- 734,032 complexes
- Training: 706149, Validation: 155, Test: 56
- Target Protein-Mini Binder Complex
- 10 target proteins
- 5 seen classes, 5 zero-shot
- Antigen-Antibody Complex
- 4261 complexes



PPDiff achieves the highest success rate and novelty General Protein-Protein Complex Design 0.2 ■ Structured Encoder+ProteinMPNN ■ InterleavingDiff ■ SSINC Network ■ PPDiff

Cases designed by PPDiff

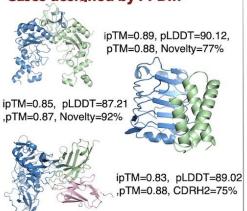
PDB

PPBench

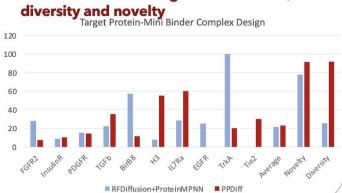
rotein-Protein

Complex

(a) PPDiff



Target Protein-Mini Binder Complex Design PPDiff achieves the higher success rate,





Orchestrating a brighter world

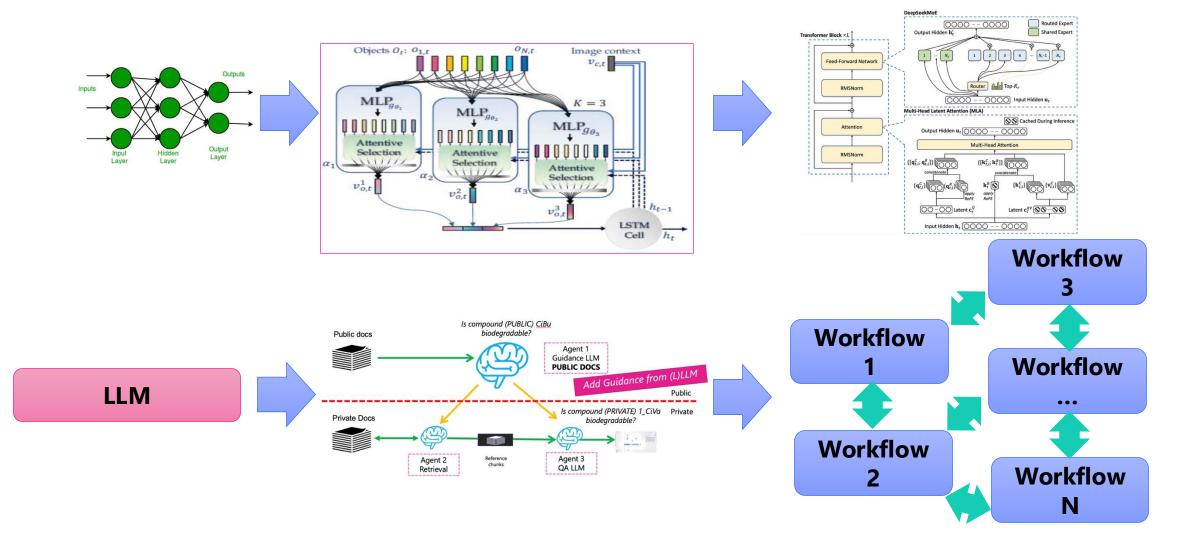
Future Directions

Al will make precision medicine scalable, accessible, and affordable worldwide.

What should we do?

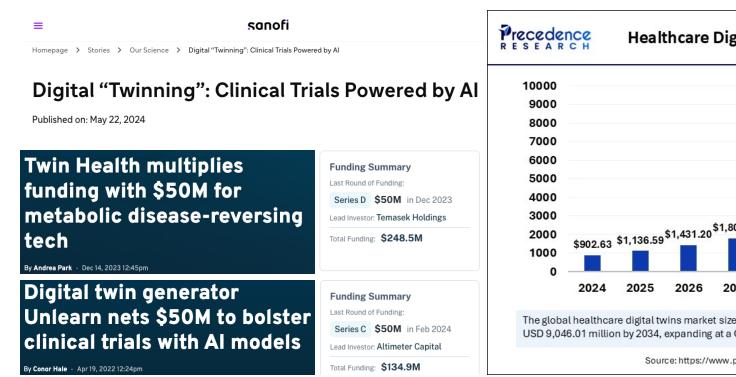


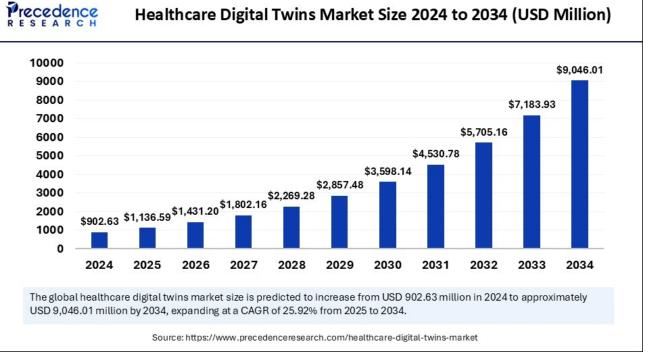
Capture Tacit Knowledge into Workflows: From LLM to Structured Workflow Al



Digital Cell and Patient

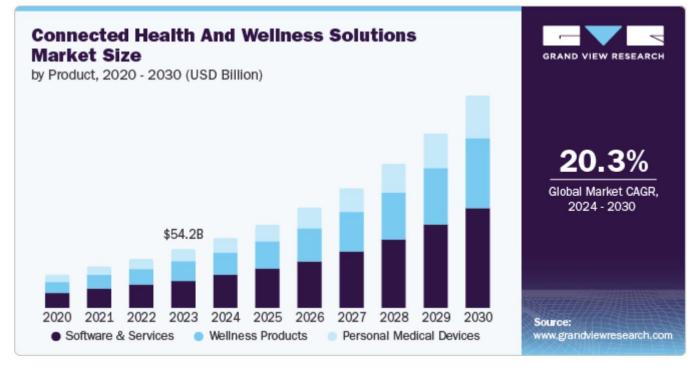
Increasing efforts from both **pharmaceutical and IT companies** in building and utilizing digital twins of cells and patient models to reduce the trial-and-error of clinical research





Connected Health

Primary care providers, virtual care providers, and specialist clinics will buy services to summarize incoming medical records, triage inquiries from patients and augment doctorpatient interaction

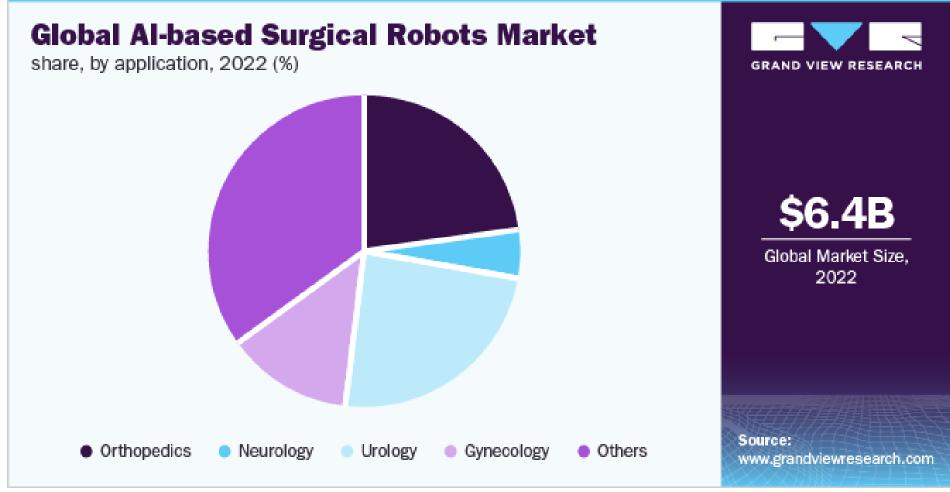


https://www.grandviewresearch.com/industry-analysis/connected-health-wellness-devices-market



Al-driven Robotic Surgery

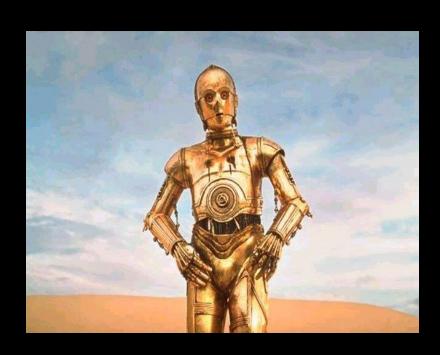
Al-based surgical robots' segment was valued at \$6.4 billion in 2022 and is expected to expand at a CAGR of **18.9% from 2023 to 2030**



The Ultimate Future of Machine Learning with "Simulated Consciousness"

From Generative Digital Cell and Digital Human to

Mike Peter Robot



Bill Gates Professor of Medicine, Molecular Biophysics & Biochemistry and Computer Science A.B. 2039, Harvard University Ph.D. 2043, Cambridge University Joined Yale Faculty 2047

Professor Robot does research in the new field of Rapid Human Disease Treatment and Prevention. He won the Nobel Prize in Physiology for discovering ABCBBCNBC007 for the no-pain one-smell treatment of cancer in 2067.

Most of this slide was shown during a talk at Yale in 2015

The Ultimate Future of Machine Learning with "Simulated Consciousness"

From Generative Digital Cell and Digital Human to

Mike Peter Robot

Of the Human, By the Human, For the Human



Bill Gates Professor of Medicine, Molecular Biophysics & Biochemistry and Computer Science

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Acknowledgements

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Prof. Xia Ning (OSU)

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Prof. Jing Zhang (UCI)

NEC

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