

The Smart Drug Discovery Alliance

Accelerating Decentralized AI-empowered Drug Discovery through Industry-Academia Coalitions





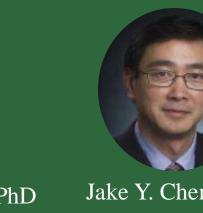


sets, drug

profile similarities

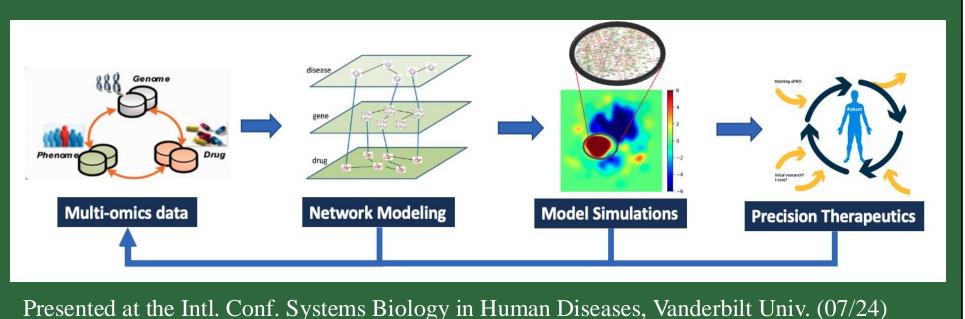
(drug-drug)





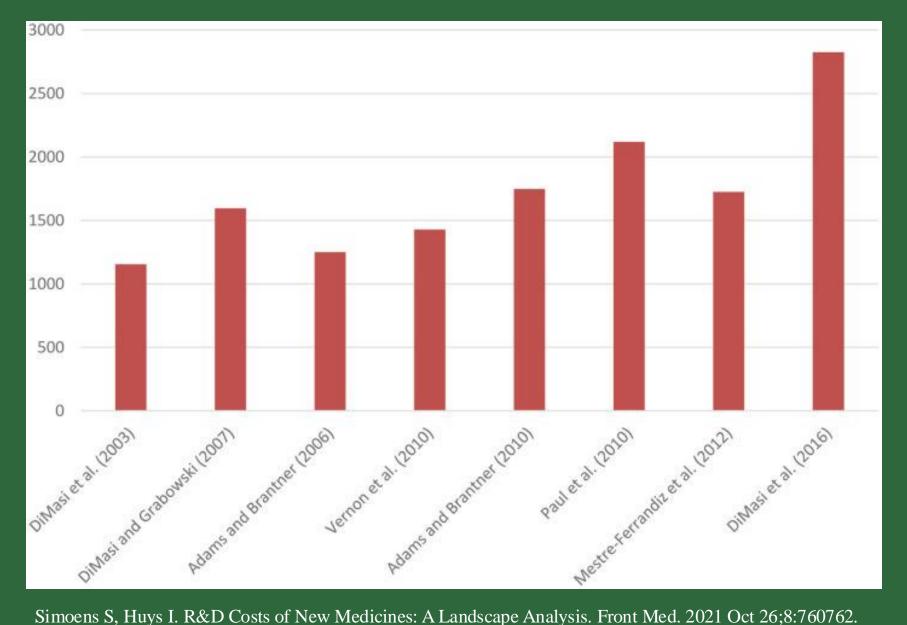
AI + Drug Discovery Manifesto

- 1. Curate AI-ready open data sets for drug discovery
- 2. Develop hierarchical accurate simulation models for efficacy/toxicity evaluations.
- 3. Nurture a decentralized AI drug discovery ecosystem
- 4. Train AI and drug discovery talents through expert team
- 5. Establish academia-industry collaborations
 - Platform Licensing
 - Innovation lab with data-driven insights
 - Sponsored research and development projects
 - Industry-academia program (Yale biomedical data science *fellowship example)*
 - Joint panel, workshop, symposium, report, white papers



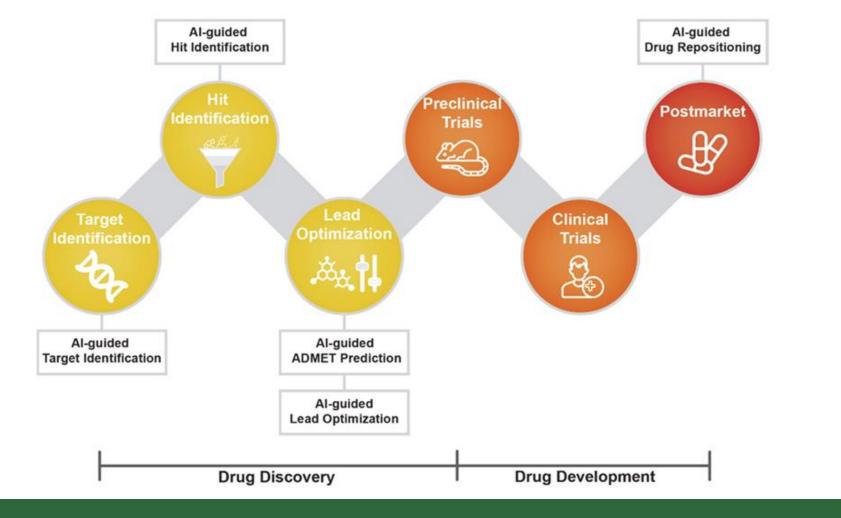
Can AI address drug discovery cost and speed issues?

~3x Drug Development Cost Increase over 20 years (2003-2023)



doi: 10.3389/fmed.2021.760762

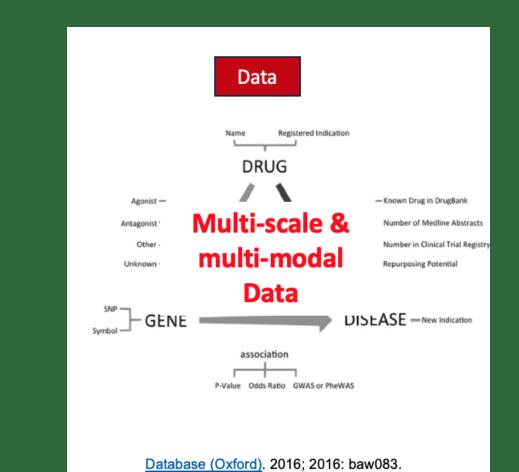
AI has been introduced in all stages of drug discovery today in the industry to optimize processes

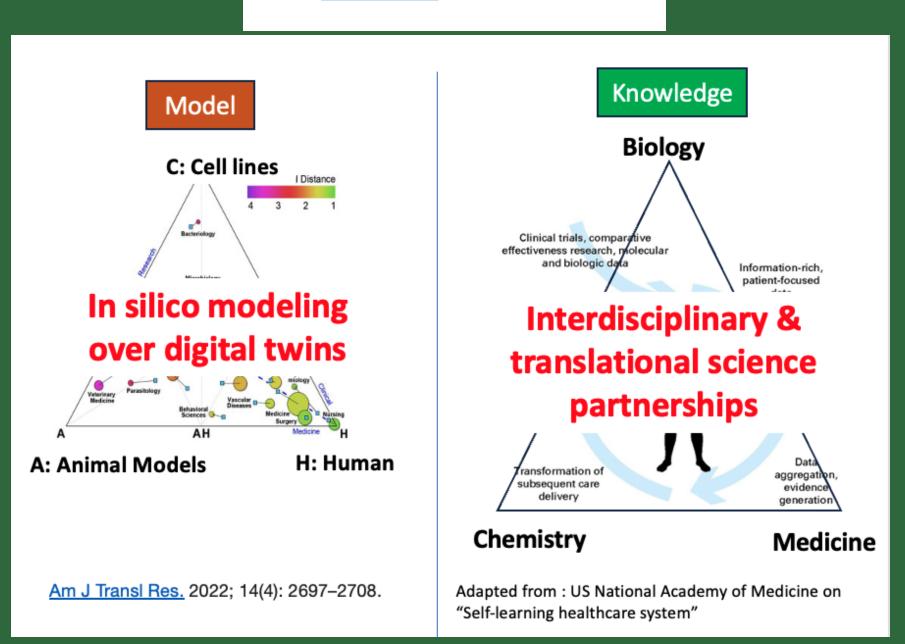


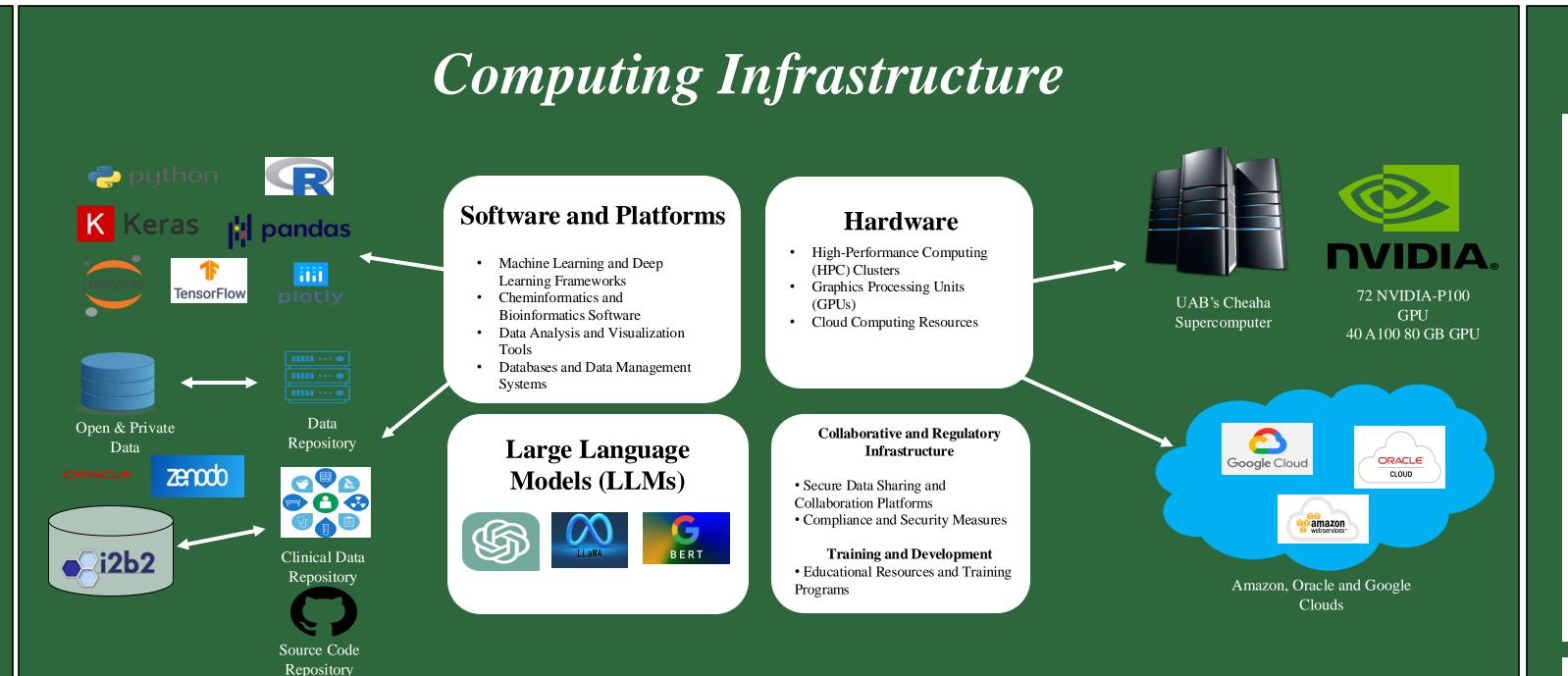
In Biotechnology and Bioprocess Engineering 25: 895-930 (2020) DOI 10.1007/s12257-020-0049-y

New Disruptive Drug Discovery Pathways Required

Data + Model + Knowledge + Regulatory transformation











PAGER Web App: An Interactive, Online **Gene Set and Network Interpretation Tool** for Functional Genomics https://doi.org/10.3389/fgene.2022.820361

Top 30 enriched P-type PAGs in PAGER

BEERE: a web server for biomedical entity expansion,

ranking and explorations

GeneTerrain

A) Creating four GeneTerrians based on four samples, a gene expression

biology, which will be calculated through PPI to find the positions of genes

in an x-y coordinate. Finally, a 2-D Gaussian distribution will be mapped

matrix including four samples and five genes. Creating a PPI network

B) Demonstrating of the effects of Sigma on the GeneTerrian, which

generates five genes, with the values of 0.05, 0.1, and 0.2 selected

through the expression level of the gene and their position.

Digital twin simulations of cancer growth and

if (x_i, y_i) is cancer and $P_{mutation}$

 P_{growth} — and $P_{metastasis}$ +

A Framework of the Geneterrain Method.

metastasis

FIN FYN or Tyrosine-protein kinase

https://doi.org/10.1093/nar/gkz428

and prioritization https://pubmed.ncbi.nlm.nih.gov/36407327/ CAMPA FACIN WINTER WINT APO DVL1 DVL2 WNT1 WNT5B WNT8A WNT8B WNT3A WNT5A WNT11 WNT4 LRP5 LRP6 KEAP1 ASSO

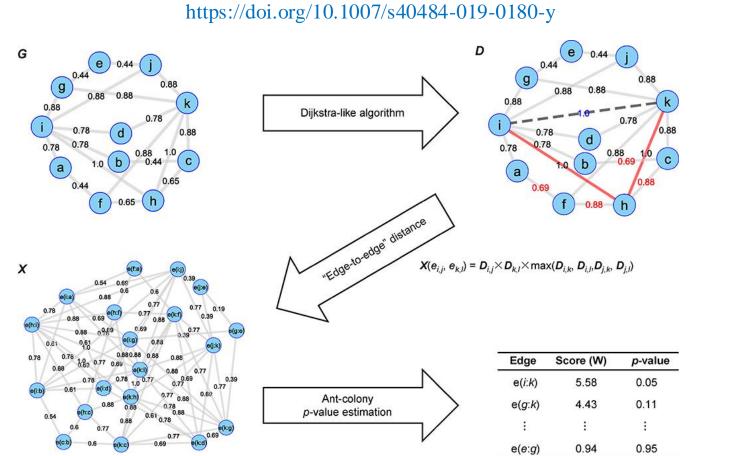
DOBS MICES AKTS AKTS AKTS KPCB MKOS MKOS PSS MYC NESS ACTS ACTS KSM82 ACLEA HANDKT TOPA BAX DISTAZ ELAT POLK KOD2 PORON TOES JAKS JAKS JAKS ILZRB ILZ (IS I GRB2 MET SOS1 PK3CA PK3CD TGFB1 AXIN1 DVL3 1000 CTNB1 GSK3B SHC1 GAB1 9000 SMAD3 EGFR PLCG

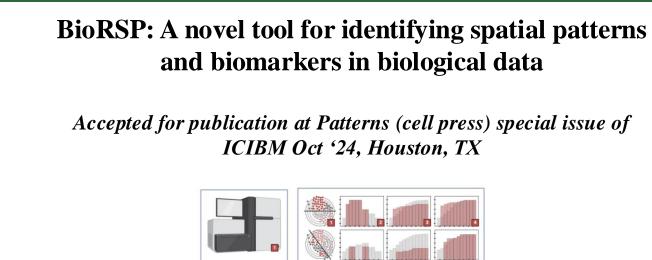
Weighted In-Network Node Expansion and Ranking

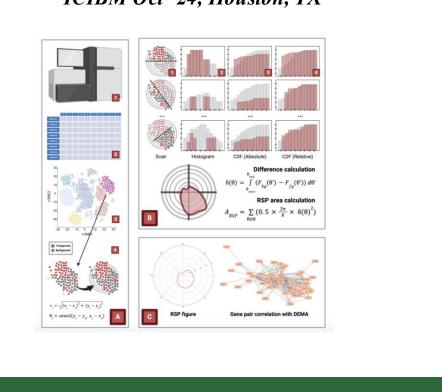
(WINNER):

A network biology tool for biomolecular characterization

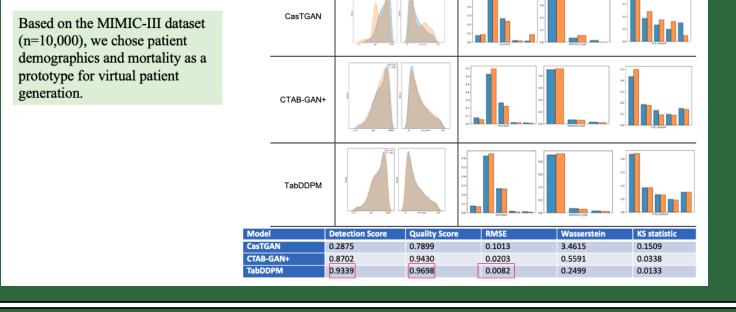
WIPER: Weighted in-Path Edge Ranking for biomolecular association networks



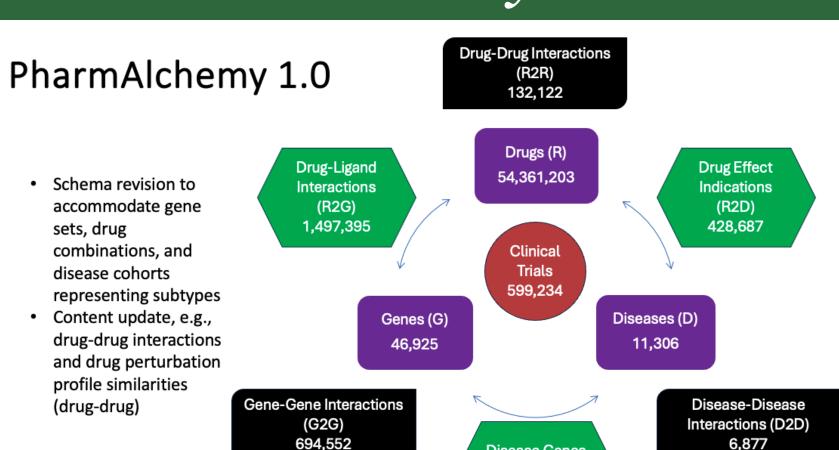




Virtual patients using generative AI NCI Supplement proposal submitted in Aug '24



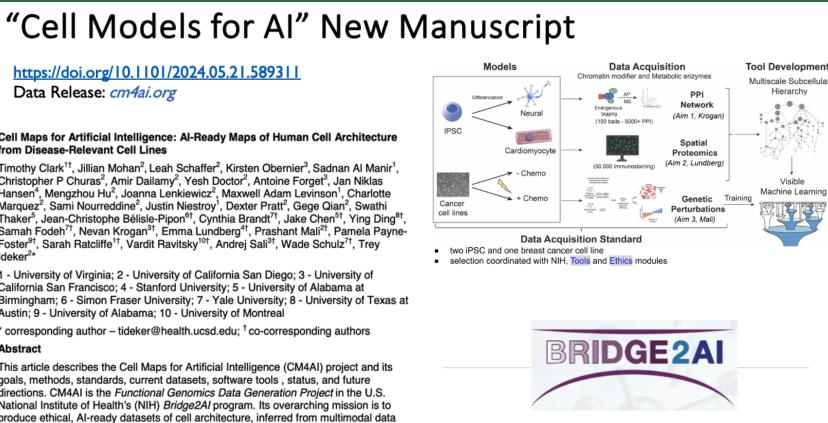
Data Layer

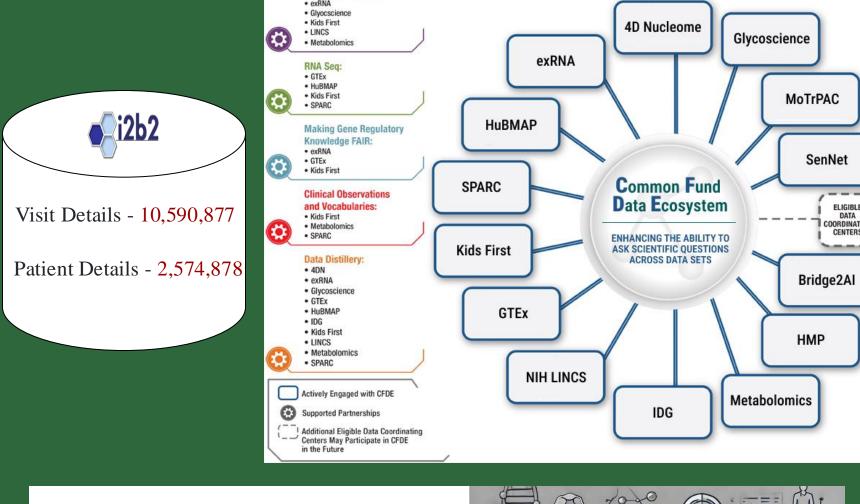


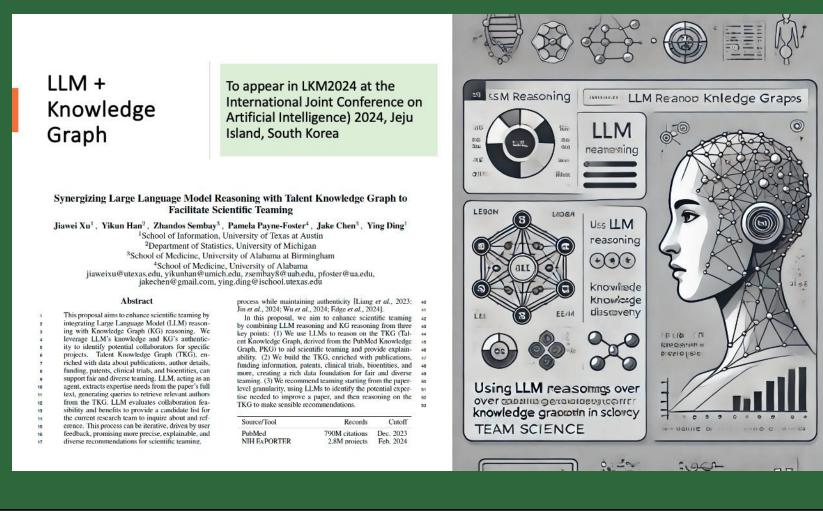
(D2G)

84.038

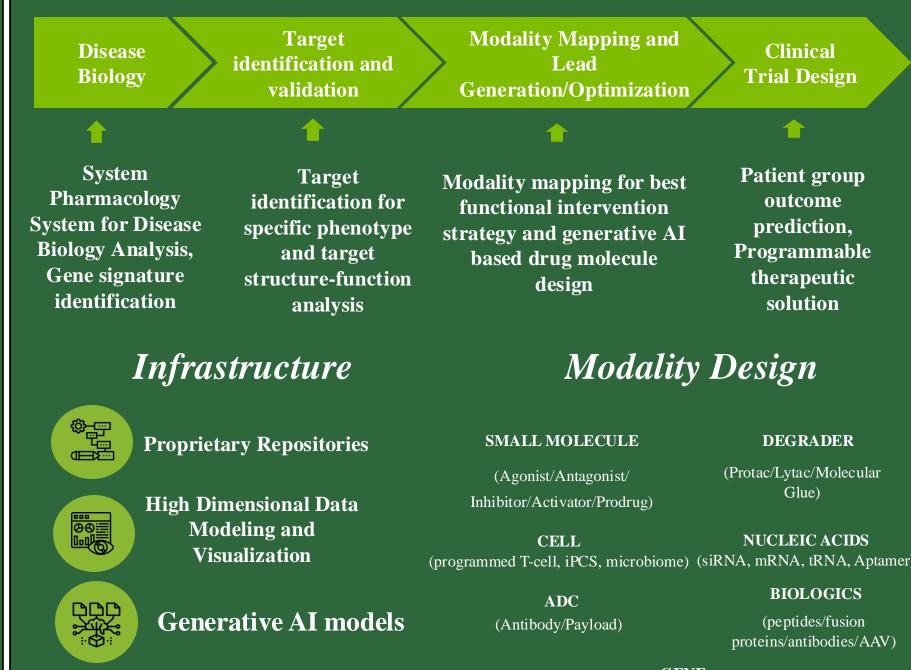












(CRISPR-Cas editing, functional Gene delivery)

Selected Drug Discovery Meetings Participated/Organized in 2024













HEALTHDAY

" Building Health Al

Ecosystem:

From Data

Harmonization to

Knowledge Discovery

Tuesday, August 27th, 2024

Convention Center Plaça de Willy Brandt, 11-14

Sant Martí, 08019 Barcelona

International Barcelona