

Mehdi Yazdani-Jahromi

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Summary

AI research scientist and PhD in Computer Science specializing in representation learning, with primary applications in computational biology and drug discovery. Industry experience at Johnson and Johnson and Microsoft, where I developed language models for mRNA and machine learning systems for large scale biomedical data. Strong programming background in Python, and JavaScript, with deep experience in PyTorch, Transformers and graph neural networks. Published in NeurIPS, ICLR and Briefings in Bioinformatics, with work spanning mRNA modeling, drug target interaction prediction and algorithmic fairness.

Skills

Programming Languages: Python, Dart, JavaScript, C++, C#, SQL, Solidity, MATLAB

Python Packages: Pytorch, Numpy, Pandas, Keras, TensorFlow, Scikit-learn, Scipy, Networkx, iGraph, dgl, matplotlib, seaborn

JavaScript Frameworks: Vue.js, Electron, NativeScript

Tools: Kubernetes, Docker, Git, HTML, CSS, LATEX, AWS, Azure, Slurm

Quantitative Research: Machine Learning Methods, Artificial Intelligence, Large Language Models, Generative AI, Genomics Sequence Modeling, Drug Discovery, Graph Neural Networks, Transformers, Recurrent Neural Networks, LSTMs, Mathematical Optimization, Mathematical Modeling

Soft Skills: Research team coordination, Student mentoring, Technical writing, Conference presentations, Resource allocation, Strategic planning, Analytical thinking, Complex problem resolution, International team collaboration, Multi-cultural communication, Self-directed learning, Professional networking

Education

PhD	University of Central Florida , Computer Science	Jan 2021 – Oct 2025
	• Thesis: Advancing Drug Discovery with Structural and Representation Learning of Biological Systems Link	
	• GPA: 3.91/4.0	
MS	University of Central Florida , Computer Science	Jan 2021 – Aug 2023
	• GPA: 3.9/4.0	
MS	Sharif University of Technology , Industrial Engineering	Sept 2017 – Dec 2019
	• GPA: 3.88/4.0	

Research Experience

University of Central Florida, Graduate Research Assistant

Orlando, FL

Aug 2021 – Dec 2025

- Conducted advanced research in computational drug discovery, focusing on drug-target interaction and algorithmic fairness.
- Developed and implemented machine learning models, including Transformers and Graph Neural Networks, for computer vision applications.
- Collaborated on multiple interdisciplinary projects, contributing to the advancement of AI methodologies in drug discovery.
- Published research findings in reputable journals and presented them at international conferences.
- Utilized tools such as Pytorch, TensorFlow, and Scikit-learn to develop and evaluate innovative algorithms.
- Engaged in data analysis and model optimization to enhance prediction accuracy and computational efficiency.
- Assisted in mentoring undergraduate and graduate students and contributed to the academic community through collaborative efforts and knowledge-sharing sessions.

Microsoft, Research Intern, Bio-LLMs

Redmond, WA

June 2025 – Sept 2025

- Built a novel agent orchestration system for planning, tool use, and judgment.
- Explored agentic AI on deidentified clinical datasets with end to end experiments and evaluation.
- Developed a RAG retrieval stack with hybrid dense and sparse search, reranking, and citation attribution.
- Contributed to the development of evaluation metrics and benchmarks for agentic AI systems.

Johnson & Johnson (Janssen R&D), Data Science Intern, AI/ML for Drug Discovery

May 2024 – Nov 2024

- Developed and trained HELM (Hierarchical Encoding for mRNA Language Modeling), the first mRNA antibody language model, achieving up to 8% increase in prediction accuracy and enabling the generation of more diverse and biologically plausible sequences.
- Explored alternative attention architectures such as Mamba and Hyena, enhancing transformer model efficiency and effectiveness in processing mRNA sequences.
- Implemented large-scale distributed training on Kubernetes, resulting in significant reduction in training time for LLMs on extensive datasets, optimizing resource utilization and scalability.
- Collaborated with cross-functional teams to refine and deploy advanced LLM architectures, enhancing model accuracy and efficiency for large-scale data processing tasks.

Selected Publications

Equi-mRNA: Protein Translation Equivariant Encoding for mRNA Language Models

2025

Mehdi Yazdani-Jahromi, Ali Khodabandeh Yalabadi, Ozlem Ozmen Garibay

[10.48550/arXiv.2508.15103](https://arxiv.org/abs/2508.15103) (Neurips 2025)

BoKDiff: Best-of-K Diffusion Alignment for Target-Specific 3D Molecule Generation

2025

Ali Khodabandeh Yalabadi, **Mehdi Yazdani-Jahromi**, Ozlem Ozmen Garibay

[10.48550/arXiv.2501.15631](https://arxiv.org/abs/2501.15631) (Advances in Bioinformatics)

HELM: Hierarchical Encoding for mRNA Language Modeling

2024

Mehdi Yazdani-Jahromi, Mangal Prakash, Tommaso Mansi, Artem Moskalev, Rui Liao

[10.48550/arXiv.2410.12459](https://arxiv.org/abs/2410.12459) (ICLR 2025, Neurips 2024 Workshop on AI for New Drug Modalities)

Fair Bilevel Neural Network (FairBiNN): On Balancing fairness and accuracy via Stackelberg Equilibrium	Jan 2024
Mehdi Yazdani-Jahromi , Ali Khodabandeh Yalabadi, AmirArsalan Rajabi, Aida Tayebi, Ivan Garibay, Ozlem Ozmen Garibay 10.48550/arXiv.2410.16432 (Neurips 2024)	
Learning Fair Representations: Mitigating Statistical Dependencies	Jan 2024
Aida Tayebi, Mehdi Yazdani-Jahromi , Ali Khodabandeh Yalabadi, Niloofar Yousefi, Ozlem Ozmen Garibay 10.1007/978-3-031-60611-3_8 (HCII conference 2023 Oral Presentation)	
FragXsiteDTI: an interpretable transformer-based model for drug-target interaction prediction	Jan 2024
Ali Khodabandeh Yalabadi, Mehdi Yazdani-Jahromi , Niloofar Yousefi, Aida Tayebi, Sina Abdidizaji, Ozlem Ozmen Garibay 10.1007/978-1-0716-3989-4_5 (Recomb 2024 (Oral), Neurips 2023 Workshop on New Frontiers of AI for Drug Discovery and Development)	
Controlling the misinformation diffusion in social media by the effect of different classes of agents	Jan 2023
Ali Khodabandeh Yalabadi, Mehdi Yazdani-Jahromi, Sina Abdidizaji, Ivan Garibay, Ozlem Ozmen Garibay 10.48550/arXiv.2401.11524 (The Computational Social Science Society of the Americas Annual Conference)	
Agent-Based Modeling of C. Difficile Spread in Hospitals: Assessing Contribution of High-Touch vs. Low-Touch Surfaces and Inoculations' Containment Impact	Jan 2023
Sina Abdidizaji, Ali Khodabandeh Yalabadi, Mehdi Yazdani-Jahromi, Ozlem Ozmen Garibay, Ivan Garibay 10.48550/arXiv.2401.11656 (The Computational Social Science Society of the Americas Annual Conference)	
Through a fair looking-glass: on mitigating bias in image datasets	Jan 2023
Amirarsalan Rajabi, Mehdi Yazdani-Jahromi , Ozlem Ozmen Garibay, Gita Sukthankar 10.1007/978-3-031-35891-3_27 (HCII conference 2023 (Oral), AAAI 2023 Workshop on Representation Learning for Responsible Human-Centric AI)	
BindingSiteAugmented DTA to enable A Next-Generation Pipeline for Interpretable Prediction Models in Drug-Repurposing	Jan 2023
Niloofar Yousefi, Mehdi Yazdani-Jahromi , Aida Tayebi, Elayaraja Kolanthai, Craig J Neal, Tanumoy Banerjee, Agnivo Gosai, Ganesh Balasubramanian, Sudipta Seal, Ozlem Ozmen Garibay 10.1093/bib/bbad136 (Briefings in Bioinformatics)	
AttentionSiteDTI: Attention Based Model for Predicting Drug-Target Interaction Using 3D Structure of Protein Binding Sites	Jan 2022
Mehdi Yazdani-Jahromi , Niloofar Yousefi, Aida Tayebi, Elayaraja Kolanthai, Craig J Neal, Sudipta Seal, Ozlem Ozmen Garibay 10.1093/bbac272 (Briefings in Bioinformatics)	
UnbiasedDTI: Mitigating Real-World Bias of Drug-Target Interaction Prediction	Jan 2022
Aida Tayebi, Niloofar Yousefi, Mehdi Yazdani-Jahromi , Elayaraja Kolanthai, Craig J Neal, Sudipta Seal, Ozlem Ozmen Garibay 10.3390/molecules27092980 (MDPI Molecules)	

Academic Services

Reviewer of Journal *Briefings in Bioinformatics*

Reviewer of *Computational and Structural Biotechnology Journal*

Reviewer of *IEEE Transactions on Neural Networks and Learning Systems Journal*

Program Committee Member of *AAAI Artificial Intelligence for Social Impact 2025 & 2026.*

Reviewer of *Neurips 2025 (The Thirty-Ninth Annual Conference on Neural Information Processing Systems)*