

# Darren Liu

Biophysics Ph.D. Graduate Student | University of Chicago

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## Summary

- Molecular biologist with expertise in high-throughput assay development, laboratory automation, and chemical biology.
- Biophysicist with expertise in analyzing high-dimensional biological datasets, protein language models, and building biologically informed computational models.
- Interdisciplinary collaborator with deep knowledge in many fields such as molecular evolution, dimensionality reduction, and DNA sequencing.

## Education

### University of Chicago

*Ph.D. Biophysical Sciences. PI: Aaron Dinner and Arvind Murugan*

2022-present

- Research Interests: protein evolution, automated liquid handling (robots, acoustic), dimensionality reduction, protein language models, chemical biology, machine learning

### University of Chicago

*M.S. Biophysical Sciences*

2022-2023

### Boston University

*B.A. in Chemistry: Biochemistry. Double minors in Biology and Computer Science*

2018-2022

## Research and Work Experience

### University of Chicago

*Graduate Student. PI: Aaron Dinner, Arvind Murugan, Department of Biophysical Sciences*

2022-present

- Developed a high-throughput automated plaque assay workflow integrating viral molecular evolution, liquid handling, and image segmentation to characterize protein evolvability.
- Built several families of Ising models to identify key parameters governing evolvability

### Boston University School of Medicine

*Undergraduate Researcher. PI: William Lehman, Department of Pharmacology, Physiology & Biophysics*

2022

- Simulated tropomyosin point mutations with molecular dynamics (NAMD)
- Connected molecular changes to disease phenotypes (hypertrophic cardiomyopathy)

### Stanford University

Summers of 2018-2022

*Undergraduate Researcher. PI: Julia Salzman, Department of Biochemistry*

- Implemented classical and machine learning image segmentation methods for spatial transcriptomics.
- Mapped relationships between gene expression and viral load in COVID-positive patients
- Developed and evaluated models to extract RNA isoform information from MERFISH image data

## Teaching and Service

### Center for Living Systems Chalk Talk Organizer

2025-present

*Organizer for Chalk Talk seminar series*

### University of Chicago Teaching Assistant

2024

*Created assignments, graded coursework, and lectured for "Computational Biology in Microbial Ecosystems" with PI Seppe Kuehn*

### UCOMBO

2022-present

*Volunteer. Taught computational biology (Python, DNA) in elementary, middle, and high schools*

### GirlsWhoCode

2022

*Volunteer. Taught fundamentals of coding to girls in elementary and middle school*

<b>@rtifice</b>	2022-2023
<i>Volunteer. Taught middle school kids basics of circuits and coding</i>	
<b>Boston University Chemia</b>	2020-2022
<i>Volunteer. Tutored undergraduates in organic and physical chemistry</i>	
<b>MVLA Aspire</b>	2017
<i>Volunteer. Provided free ACT tutoring for students</i>	

## Skills

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- Programming: Python (NumPy, PyTorch, SciPy, Biopython, Jupyter, Sci-kit Learn), High-Performance Computing (SLURM, bash), Git, LaTeX
- Computation: Dimensionality reduction, clustering, regression, protein language models (ESM2), molecular dynamics (NAMD), AlphaFold, VMD/PyMOL, image segmentation (ImageJ, OpenCV, scikit-image, PyTorch), sequencing analysis
- Wet Lab: Assay development, automated liquid handling (Tecan Fluent, Echo Acoustic Liquid Handler), plate readers (fluorescence and absorbance assays), directed molecular evolution, cell culture and cloning, DNA sequencing

## Publications

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I. Firlar, M. Rynkiewicz, J. Creso, **D. Liu**, J. Moore, W. Lehman, S. Campbell. Predictive genotype-phenotype correlation of four tropomyosin-1 variants of unknown significance. (2023) Biophysical Journal, Volume 122, Issue 3, 121a

## Presentations and Posters

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*Mapping the evolvability of molecular specificity across a protein family.* **Poster, Gordon Research Conference on Molecular Mechanisms in Evolution** (June 2025)