

Spencer Guo

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SUMMARY

- Biomolecular simulation and sampling expert with 6+ years of experience in molecular modeling
- Interdisciplinary collaborator with broad knowledge in structural and machine learning methods
- Scientific communicator and author of peer-reviewed publications in *Nature* and *PNAS*

EDUCATION

Ph.D. Chemistry, **The University of Chicago** 9/2020 – 6/2025 (expected)
Supervisors: Profs. Aaron Dinner and Benoît Roux
Thesis: Long-timescale kinetics and mechanisms of protein conformational change
Research interests: molecular dynamics simulations, protein kinetics, membrane proteins, machine-learning methods for statistical mechanics, integration of experimental biophysics data (cryo-EM, NMR, electrophysiology)

M.S. Chemistry, **The University of Chicago** 9/2020 – 9/2021

B.S. Biological Chemistry, **Stanford University**, *Minor in Computer Science* 9/2016 – 6/2020

RESEARCH + WORK EXPERIENCE

The University of Chicago Graduate Research Assistant
Department of Chemistry 9/2020 – present

- Applied dynamical analysis and enhanced sampling to elucidate membrane protein mechanisms
- Benchmarked a non-Markovian algorithm for computing kinetic statistics with short trajectories
- Developed an method leveraging pretrained GNNs to automatically featurize molecular structures
- Collaborated closely with experimentalists to integrate NMR and HDX data
- Wrote a successful grant application to obtain Anton supercomputer time for simulations

Stanford University Undergraduate Research Assistant
Department of Chemistry, Markland Lab 9/2018 – 6/2020

- Simulated IR spectra of liquid water using *ab initio* molecular dynamics
- Adapted a neural network method to calculate molecular dipoles

Schrödinger Python Development Intern
New York, NY 6/2019 – 9/2019

- Developed tool to identify critical residue/ligand interactions for drug development
- Extended multiple sequence viewer (MSV) to analyze similarity at binding sites

Genentech Protein Engineering Intern
South San Francisco, CA 6/2018 – 9/2018

- Synthesized novel peptide library for PROTAC optimization using rational macrocycle design
- Analyzed instrumental purity and spectral data (LC-MS, HPLC, NMR)

SERVICE EXPERIENCE

Science One-Pager Initiative
Illinois Science Policy Consortium 3/2024 – present

- Authored a one-page policy brief about therapeutic use of psychedelics for mental health disorders
- Summarize and synthesize scientific studies to support evidence-based policy in Illinois legislature

Modern Materials and Technology Volunteer
The University of Chicago 9/2021 – present

- Teach a weekly colloquium to ~15 high school students in Chicago
- Create lesson plans demonstrating applications of chemistry and physics to modern technology

Department of Chemistry

The University of Chicago

Director of Graduate Student Initiatives

7/2022 – 6/2023

- Served as liaison between graduate students and faculty to increase open communication
- Developed policies to alleviate student concerns and help their transition to graduate school
- Designed policies at cross-department weekly meetings and organize feedback from students

SKILLS

Technical	structural modeling, molecular dynamics simulations, enhanced sampling techniques, graph neural networks, Monte Carlo methods
Programming	(<i>experienced</i>) Python (NumPy/SciPy), jax, PyTorch, bash, git, L ^A T _E X (<i>familiar</i>) C/C++, Matlab
Software	(<i>experienced</i>) Amber, OpenMM, VMD, MDAnalysis, Alphafold (<i>familiar</i>) GROMACS, BioPython, PyMOL, CP2K, DFTB+

PUBLICATIONS + PREPRINTS

* Denotes equal contribution.

1. K. Jeong, S. Guo, S. Allaw, & A. R. Dinner. Analysis of the dynamics of a complex, multipathway reaction: Insulin dimer dissociation. *J. Phys. Chem. B*, ASAP (2024).
2. N. Zhang*, D. Sood*, S. Guo*, N. Chen, *et al.* Temperature-Dependent Fold-Switching Mechanism of the Circadian Clock Protein KaiB. *Proc. Natl Acad. Sci. USA*, **121**, e2412327121 (2024).
3. Z. Pengmei, C. Lorpaiboon, S. Guo, J. Weare, & A. R. Dinner. geom2vec: pretrained GNNs as geometric featurizers for conformational dynamics. arXiv:2409.19838 (2024).
4. S. Guo, R. Shen, B. Roux, & A. R. Dinner. Dynamics of activation in the voltage-sensing domain of *Ciona intestinalis* phosphatase Ci-VSP. *Nat. Commun.* **15**, 1408 (2024).
5. C. Lorpaiboon, S. Guo, J. Strahan, J. Weare, & A. R. Dinner. Accurate estimates of dynamical statistics using memory. *J. Chem. Phys.* **160**, 084108 (2024).
6. J. Strahan, S. Guo, A. R. Dinner, & J. Weare. Inexact iterative numerical linear algebra for neural network-based spectral estimation and rare-event prediction. *J. Chem. Phys.* **159**, 014110 (2023).

PRESENTATIONS

- Talk at Center for Learning Systems Cell State Transitions Workshop, Chicago, IL** 5/2024
– Learning dynamical statistics from short-trajectory data
- Poster at IMSI “Learning Collective Variables and Coarse Grained Models” Workshop** 2/2024
– Inexact subspace iteration for neural network-based rare-event prediction
- Invited lecture at Associated International Laboratory (CNRS/UIUC), Hauteluce, France** 1/2024
– Complex kinetics and mechanism of voltage activation in the voltage-sensing domain of Ci-VSP
- Poster at AI & Science Summer School 2023, Chicago, IL** 7/2023
– Inexact iterative numerical linear algebra for neural network-based spectral estimation and rare-event prediction
- Poster at Biophysical Society Annual Meeting, San Francisco, CA** 2/2022
– Dynamical analysis of voltage-dependent activation in Ci-VSP

AWARDS + GRANTS

St. Jude National Graduate Student Symposium, St. Jude Children’s Hospital	2024
Anton 2 Biomolecular Simulation Grant, Pittsburgh Supercomputing Center	2021
NSF Graduate Research Fellowship, National Science Foundation	2020
Eckhardt Fellowship, Physical Sciences Division, The University of Chicago	2020

Last updated December 21, 2024.