Erik Nordquist

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Education

2023 **Ph.D.** Chemistry, University of Massachusetts Amherst.

2018 **B.S.** Chemistry and Physics, The College of Idaho.

Work Experience

2018–23 **Graduate Research Assistant**, University of Massachusetts, Amherst, Department of Chemistry.

- Advisor: Jianhan Chen, Ph.D.
- Thesis: Understanding Protein Structure-Function Relationships Using Simulations and Machine Learning
- Focus: Computational biophysics, Molecular dynamics, Machine learning
- 2023 Postdoctoral Fellow, University of Maryland Baltimore, Department of Pharmaceutical Sciences.
 - Advisor: Alexander D. MacKerell, Jr., Ph.D.
 - Focus: Computer-aided drug design (CADD), Computational methods development, Molecular dynamics

Appointments

- 2025– **Adjunct Professor**, Stevenson University, Department of Chemistry.
- 2023– **Postdoctoral Fellow**, University of Maryland, Baltimore, Department of Pharmaceutical Sciences.

Fellowships and Awards

- 2024 **Best of Biophysical Journal 2023,** publication titled "Inner pore hydration free energy..." **Talk award**, Research Symposium, UM Greenebaum Comprehensive Cancer Center
- 2023– NIH T32 Postdoctoral Fellowship, University of Maryland, Baltimore
- 2022 **Paul H. Terry Endowment Award**, Chemistry Dept., University of Massachusetts Amherst **Teaching Fellowship**, College of Natural Sciences, University of Massachusetts Amherst
- 2020–22 NIH T32 Predoctoral Fellowship, University of Massachusetts Amherst
- 2020 William E. McEwen Poster Award, Chemistry Dept., University of Massachusetts Amherst

Peer-Reviewed Publications

- Nordquist E, Zhao M, Yu W, MacKerell A. Computational Modeling of PROTAC Ternary Complexes as Ensembles Using SILCS-PROTAC. J. Chem. Inf. Model. (submitted) DOI: 10.26434/chemrxiv-2025-jk9sn
- Patel R, **Nordquist E**, Polli J. Prediction of surfactant-mediated dissolution of poorly soluble drugs from drug particles. **Eur. J. Pharm. Sci.** 2025, 107052. DOI: 10.1016/j.ejps.2025.107052
- Nordquist E[#], Zhao M[#], Kumar A, MacKerell A. Physics- and Machine-Learning Based Method to Identify Druggable Binding Sites Using SILCS-Hotspots. J. Chem. Inf. Model. 2024, 64, 19, 7743-7757. DOI: 10.1021/acs.jcim.4c01189 *Contributed equally.
- Nordquist E, Jia Z, Chen J. Small Molecule NS11021 Promotes BK Channel Activation by Increasing Inner Pore Hydration. J. Chem. Inf. Model. 2024, 64, 19, 7616-7625. DOI: 10.1021/acs.jcim.4c01012

- Nordquist E*, Zhang G*, Barethiya S, Ji N, White K, Han L, Jia Z, Shi J, Cui J, and Chen J. Incorporating physics to overcome data scarcity in predictive modeling of protein function: a case study of BK channels. PLOS Comput. Biol. 2023 19(9): e1011460. DOI: 10.1371/journal.pcbi.1011460 *Contributed equally.
- Zhang L, Barethiya S, Nordquist E, Chen J. Machine Learning Generation of Dynamic Protein Conformational Ensembles. Molecules 2023, 28(10), 4047. DOI: 10.3390/molecules28104047
- Nordquist E, Zhiguang J, Chen J. Inner pore hydration free energy controls activations of the big potassium channel and its mutants. Biophys. J. 2023, 122, 1158-1167. DOI: 10.1016/j.bpj.2023.02.005 Selected for Best of 2023 Edition.
- Nordquist E#, Schultz S#, and Chen J. Using Metadynamics To Explore the Free Energy of
 Dewetting in Biologically Relevant Nanopores. J. Phys. Chem. B 2022, 126, 34, 6428-6437 DOI:
 10.1021/acs.jpcb.2c04157 #Contributed equally. Note: S. Schultz was an undergraduate I mentored
 during this project.
- Nordquist E, Clerico E, Chen J, Gierasch L. Computational Modeling of Hsp70-Client Interactions: Past, Present, and Future. J. Phys. Chem. B 2022, 126, 36, 6780–6791 DOI: 10.1021/acs.jpcb.2c03806
- Nordquist E, English C, Clerico E, Sherman W, Gierasch L, Chen J. Physics-based modeling provides predictive understanding of selectively promiscuous substrate binding by Hsp70 chaperones. PLOS Comput. Biol. 2021, 17 (11): e1009567. DOI: 10.1371/journal.pcbi.1009567
- Gong X, Chiricotto M, Liu X, **Nordquist E**, Feig M, Brooks CL, Chen J. Accelerating the generalized born with molecular volume and solvent accessible surface area implicit solvent model using graphics processing units. **J. Comput. Chem.** 2020, 41, 830–838. DOI: 10.1002/jcc.26133

Teaching and Mentoring

- 2025 **Instructor of record,** General Chemistry Lab, Stevenson University. Full teaching responsibility for one section of the lab class, including grading and office hours.
- Instructor of record, First-Year Seminar, University of Massachusetts Amherst. Title: "Reconciling Atomic Chaos and Human Order". I designed this course through the College of Natural Science Teaching Fellowship, including a two-day "teaching bootcamp." (link)
 - **Course goals**: develop skills in scientific communication, and a perform 4-week mini research project. Assessments were primarily formative, with self-feedback to foster growth mindsets.

2020-24 Guest lectures:

Computer-aided Drug Design in a graduate Cancer Biology course, UMB (2x); Discussion seminar moderator on AlphaFold2 in an undergrad Biophysics course, Amherst College; Lecture on molecular mechanics and additive force fields in a graduate Stat. Mech. Course, UMass

2018–19 Lab TA, General Chemistry I Lab, University of Massachusetts Amherst

Mentoring Experience:

Dami Showunmi (undergrad., UMBC, 2025–); Anthony O'Donnell (grad., JHU, 2024–); Samantha Schultz (undergrad., UMass, 2020–2021, **resulted in publication**); Callie Jillson (undergrad., UMass, 2019–2020)

Presentations

Talk, Frontiers at the Chemistry-Biology Interface Symposium, Johns Hopkins University, "Computational modeling of ternary complexes as ensembles using SILCS-PROTAC," Baltimore, MD. Talk, Chemistry Department Seminar, George Washington University, "Computational design of PROTACs", Washington D.C.

- Talk, Institute for Bioscience and Biotechnology Research, University of Maryland Baltimore, Early-Career Research Symposium, "Computational design of PROTACs", Rockville, MD.
 - **Talk**, University of Maryland Greenbaum Comprehensive Cancer Center Research Symposium, "Physics- and machine-learning-based method for identifying druggable binding sites with SILCS-Hotspots." (Talk award) Baltimore, MD.
 - **Talk**, The College of Idaho Natural Science Symposium, "Computer simulations of proteins help understand their function." Caldwell, ID.
 - **Poster**, Biophysical Society Annual Meeting, "Computational mapping of allosteric modulators of the BK channel." Philadelphia, PA.
- 2023 **Poster,** Biophysical Society Annual Meeting, "A predictive model of voltage gating of BK channels via physical modeling and machine learning." San Diego, CA.
- Talk, University of Massachusetts Amherst ResearchFest (PH Terry award): "Predicting protein function with physics, experiments and machine learning." Amherst, MA.
 - **Poster,** Biophysical Society Annual Meeting, "Free energy of hydrophobic dewetting in gating of BK channels." San Francisco, CA.
- Talk, Northeastern Structural Symposium, "Physical origins of selective promiscuity to Hsp70s revealed through physics-based modeling." Virtual.
 - **Poster,** University of Massachusetts Amherst ResearchFest (WE McEwen Award): "Physical origins of selective promiscuity to Hsp70s revealed through physics-based modeling." Amherst, MA.
- **Talk,** Biophysics at University of Massachusetts Amherst, "Understanding the origins of DnaK's selective promiscuity with physics-based modeling"
 - **Poster**, Molecular Biophysics in the Northeast, "Understanding the origins of DnaK's selective promiscuity with physics-based modeling." Boston, MA.

Service

- Facilitator, Skills for Faculty Positions panel, University of Maryland Office of Postdoctoral Scholars Moderator, Career Panel Discussion, University of Maryland, Baltimore Cancer Retreat Poster judge, Frontiers at the Chemistry-Biology Interface Symposium, Johns Hopkins University
- 2023–24 Facilitator, Responsible Conduct of Research NIH training, University of Maryland, Baltimore
- 2024– Organization committee, Annual Cancer Research Retreat, University of Maryland, Baltimore (2x)
- 2020–20 **Organization committee, Alumni Networking Symposium**, Chemistry-Biology Interface program, University of Massachusetts Amherst (2x)
- 2020 **Search committee,** Grad. Program Manager, Chemistry Dept. University of Massachusetts Amherst
- 2019–21 **ResearchFest committee** for Chemistry Dept., University of Massachusetts Amherst (3x)

Journals Refereed for:

Comm. Chem., Biophys. J., J. Phys. Chem., ACS Omega, J. Chem. Theory Comput., J. Chem. Inf. Model., Drug Discov. Today

Outreach

2023-25 **Volunteer instructor,** Baltimore Underground Science Space (BUGSS) (link), STEM outreach for Baltimore community members in community-accessible lab space

Volunteer instructor, RAMP Program for STEM activity for high-schoolers in Baltimore (link) (2x) **Poster judge**, Mount Royal Middle School Science Fair, Baltimore (2x)

Volunteer instructor, CURE Program University of Maryland, Baltimore, STEM outreach for middleand high-schoolers in Baltimore (link) (3x)

- **Volunteer,** Halloween STEM outreach event at Community Engagement Center, University of Maryland, Baltimore
- 2020–23 Reviewer for Journal of Emerging Investigators, 25 articles by middle- / high-school students (link)
- 2022 Lab for girls summer science camp, Eureka! at University of Massachusetts Amherst (link)

Professional Development

2025	Professional Society Memberships:
	Biophysical Society (2022–), American Chemical Society (2025–)
2024	Writing an Effective Teaching Philosophy Statement, online CIRTL workshop
	Teaching Biophysics at a PUI workshop, Biophysical Society Annual Meeting
2024	Safety preparedness trainings, CPR/AED, Stop the Bleed, Civilian Active Shooter Events, Fire
	Extinguishers, and Naloxone Training, University of Maryland, Baltimore (certificates)
2022	CITRL associate certification, University of Massachusetts Amherst, achieved through training on
	evidence-based and inclusive teaching practices (link)
2021	Evidence-based Undergraduate STEM Teaching, online course (link)
	Inclusive STEM Teaching, online course (link)

Book Chapters

• **Nordquist E**, Horrigan F, MacKerell A. Computational ligand binding site prediction. **Springer Nature.** (accepted) 2025.