

NABIN GIRI

Berkeley, CA

☎ +1 573-810-3959 🌐 nabingiri.io ✉ ngiri@lbl.gov 🔗 linkedin.com/nabin-giri 🎓 [Google Scholar](#)

Revised : August 2, 2025

EDUCATION

University of Missouri

January 2021 – May 2025

Ph.D. in Computer Science, Minor in Statistics

Columbia, MO

Emphasis: Deep Learning and Structural Biology

Dissertation: Deep Learning for Modeling Protein Atomic Structures from cryo-EM Density Maps

Summer Schools: Oxford Machine Learning & London Geometry and Machine Learning

2022, 2023

Seminar: Preparing Future Faculty Seminar (Spanned one academic semesters)

2024

University of Central Missouri

May 2018 – May 2020

Masters of Science in Computer Science - Thesis Track

Warrensburg, MO

Workshop: Graduate Research Workshop Series (Spanned one academic semester)

Thesis: Recommendation System Using Factorization Model and MapReduce Framework

Bangalore University

May 2011 – May 2014

Bachelors of Computer Application - First Class Honors

Bangalore, India

Semester Projects: Supermarket Management System & Online Admission System

EXPERIENCE

Lawrence Berkeley National Laboratory

June 2025 – Present

Machine Learning Postdoctoral Scholar

Berkeley, CA

University of Missouri

January 2021 – May 2025

Graduate Researcher at Bioinformatics and Machine Learning Lab

Columbia, MO

Amazon.com Inc.

May 2024 – August 2024

Applied Scientist II Intern

Seattle, WA

BIOVIA - Dassault Systèmes

May 2023 – August 2023

Scientific Software Development Intern

San Diego, CA

University of Central Missouri

May 2018 – May 2020

Graduate Student Worker (GSW) - James C Kirkpatrick Library

Warrensburg, MO

Max International

March 2017 – April 2018

System Engineer, Head of Department

Kathmandu, Nepal

Capgemini

October 2014 – October 2016

Software Engineer

Bangalore, India

TEACHING

Machine Learning for Biomedical Informatics

Spring 2024

Course Level: CMP_SC 8180 (INFOINST 8880)

University of Missouri

PUBLICATIONS (First-author papers **numbered in red.**)

- **[13] A labeled dataset for AI-based cryo-EM map enhancement.** Nabin Giri, Xiao Chen, Ligu Wang, and Jianlin Cheng. *Computational and Structural Biotechnology Journal*. 2025.
- **[12] Atomic Protein Structure Modeling from Cryo-EM Using Multi-Modal Deep Learning and AlphaFold3.** Nabin Giri and Jianlin Cheng. *bioRxiv* 2025.
- **[11] Protein-Ligand Structure and Affinity Prediction in CASP16 Using a Geometric Deep Learning Ensemble and Flow Matching.** Alex Morehead, Jian Liu, Pawan Neupane, Nabin Giri and Jianlin Cheng. *Proteins: Structure, Function, and Bioinformatics*, 2025.
- **[10] De novo atomic protein structure modeling for cryoEM density maps using 3D transformer and HMM.** Nabin Giri and Jianlin Cheng. *Nature Communication*, 2024.
- **[9] Cryo2StructData: A Large Labeled Cryo-EM Density Map Dataset for AI-based Modeling of Protein Structures.** Nabin Giri, Ligu Wang, and Jianlin Cheng. *Nature Scientific Data*, 2024. Dataset available in *Harvard Dataverse*.
- **[8] Outcomes of the EMDataResource cryo-EM Ligand Modeling Challenge.** Catherine L. Lawson, Andriy Kryshchak, Grigore D. Pintilie, Stephen K. Burley, . . . , Nabin Giri, . . . , Helen M. Berman, and Wah Chiu. *Nature Methods*, 2024.
- **[7] Deep Learning for Protein-Ligand Docking: Are We There Yet?** Alex Morehead, Nabin Giri, Jian Liu and Jianlin Cheng. *ICML AI4Science*, 2024.
- **[6] Impact of AlphaFold on Structure Prediction of Protein Complexes: The CASP15-CAPRI Experiment** Marc F. Lensink, Guillaume Brysbaert, Nessim Raouraoua, . . . , Nabin Giri, . . . , Sameer Velankar and Shoshana J. Wodak. *PROTEINS: Structure, Function, and Bioinformatics*, 2023.
- **[5] Combining pairwise structural similarity and deep learning interface contact prediction to estimate protein complex model accuracy in CASP15** Raj S. Roy, Jian Liu, Nabin Giri, Zhiye Guo and Jianlin Cheng. *PROTEINS: Structure, Function, and Bioinformatics*, 2023. *[Ranked First in CASP15]*
- **[4] Deep learning for reconstructing protein structures from cryo-EM density maps: recent advances and future directions** Nabin Giri, Raj S. Roy, and Jianlin Cheng. *Current Opinion in Structural Biology*, 2023.
- **[3] Improving Protein-Ligand Interaction Modeling with cryo-EM Data, Templates, and Deep Learning in 2021 Ligand Model Challenge** Nabin Giri and Jianlin Cheng. *Biomolecules*, 2023.
- **[2] DRLComplex: Reconstruction of protein quaternary structures using deep reinforcement learning** Elham Soltanikazemi, Raj S. Roy, Farhan Quadir, Nabin Giri, Alex Morehead, and Jianlin Cheng. *The International Conference on Intelligent Biology and Medicine (ICIBM)*, 2022.
- **[1] High-Performance Deep Learning Toolbox for Genome-Scale Prediction of Protein Structure and Function** Mu Gao, Peik Lund-Andersen, . . . , Nabin Giri, . . . , Jianlin Cheng and Ada Sedova. *IEEE/ACM Workshop on Machine Learning in High Performance Computing Environments (MLHPC)*, 2021.

AWARDS

Outstanding PhD Student Award <i>College of Engineering</i>	2025 <i>University of Missouri</i>
Graduate Research Award <i>Graduate Education and Research</i>	2020 <i>University of Central Missouri</i>
Graduate Housing Scholarship <i>Webb Housing</i>	2020 <i>University of Central Missouri</i>
Conference Scholarship - Gold Pass <i>Strata Data Conference</i>	2019 <i>O'Reilly Media, Inc</i>
Microsoft Partner Meet <i>Ranked 2nd out of 15 participants in the competition</i>	2018 <i>Microsoft Partner Meet Nepal</i>

SERVICE

The Protein Journal

Manuscript Reviewer

2025 - Present

GigaScience

Manuscript Reviewer

2025 - Present

International Journal of Biological Macromolecules

Manuscript Reviewer

2024 - Present

Neural Information Processing Systems (NeurIPS)

Manuscript Reviewer

2023 - Present

IEEE International Conference on Bioinformatics and Biomedicine (BIBM)

Program Committee and Manuscript Reviewer

2023 - 2025

Briefings in Bioinformatics

Manuscript Reviewer

2022 - Present

IEEE/ACM Transactions on Computational Biology and Bioinformatics

Manuscript Reviewer

2022 - Present

Undergraduate Research Forum

Research Abstract Reviewer

University of Missouri (MU)

2022

Upsilon Pi Epsilon-Gamma Chapter

Member

University of Missouri (MU)

2021 - Present

MU Nepalese Student Association

Member

University of Missouri (MU)

2021 - Present

Nepalese UCM Association

President

University of Central Missouri (UCM)

2018 - 2020

Association for Computing Machinery (ACM)

Member

University of Central Missouri (UCM)

2018 - 2019

International Student Services

International Students Ambassador

University of Central Missouri (UCM)

2018 - 2019

INVITED TALKS AND PRESENTATIONS

Modeling Atomic Structures from Cryo-EM Maps

Phenix Group, Lawrence Berkeley National Laboratory

2025

Virtual

Modeling Atomic Structures from Cryo-EM Maps

Dana-Farber Cancer Institute, Harvard Medical School

2025

Virtual

Modeling Atomic Structures from Cryo-EM Maps

Computational Biosciences Group, Lawrence Berkeley National Laboratory

2025

Berkeley, CA

Modeling Atomic Structures from Cryo-EM Maps

Artificial Intelligence Department, Brookhaven National Laboratory

2025

Upton, NY

De novo Protein Structure Modeling for CryoEM using 3D Transformer & HMM

Intelligent Systems For Molecular Biology (ISMB). Presented by Dr. Jianlin Cheng

2024

Montreal, Canada

Deep Learning for Modeling Protein Atomic Structures from Cryo-EM Maps

Department of Electrical Engineering and Computer Science

2024

University of Missouri

Mizzou Tigers Summer Internship Experiences

Electrical Engineering and Computer Science Graduate Student Association

2024

University of Missouri

Cryo2Struct: AI-Based Modeling of 3D Atomic Structures from Cryo-EM Maps

CryoEM Super Group, Department of Biochemistry

2024

University of Missouri

Improving Relevance Prediction with Multimodal Models <i>Amazon Science</i>	2024 <i>Seattle, WA</i>
De Novo Protein Structure Modeling for Cryo-EM Maps Using 3D Transformer and HMM <i>BIOVIA</i>	2024 <i>Virtual</i>
Conformational Sampling with AlphaFold and Diffusion-Based Protein-Ligand Modeling <i>BIOVIA</i>	2024 <i>Virtual</i>
Cryo2StructData: Large Labeled Cryo-EM Dataset for AI-Based Protein Structure Modeling <i>BIOVIA</i>	2023 <i>Virtual</i>
Modeling Protein-Ligand interaction from cryo-EM data, Templates and Deep Learning <i>Organized by EMDDataResource, presented the work developed for the 2021 Ligand Model Challenge</i>	2021 <i>Virtual</i>
Three Minute Thesis (3MT) <i>Presented Master's Thesis Research</i>	2020 <i>University of Central Missouri</i>

PRESS

[Paving the way for new treatments](#)

University of Missouri, College of Engineering. September, 2024

[De novo atomic protein structure modeling for cryoEM density maps - Behind the Paper](#)
Nature Portfolio Communities. July, 2024

[Journey into Nabin Giri's BIOVIA Internship: Conformational Sampling with AlphaFold](#)
BIOVIA. September, 2023

[Mizzou team ranks first in category at CASP15 protein prediction competition](#) University of Missouri, College of Engineering. January, 2023