

# ASHLIN JAMES PORUTHOOR



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## EDUCATION

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<b>Doctor of Philosophy, Biophysics</b> , University of Rochester (Anticipated)	2018 - 2023
<b>Master of Science, Biophysics</b> , University of Rochester	2018 - 2021
<b>Master of Science, Physics</b> , NIT Calicut	2016 - 2018
<b>Bachelor of Science, Physics (Honors)</b> , St.Stephen's College, University of Delhi	2013 - 2016

## SKILLS

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### Computational Biology and Structural Biology:

• All-Atom and Coarse-Grained Molecular Dynamics Simulations : (MD Engines) ◇ GROMACS ◇ OpenMM ◇ NAMD (MD Visualization) ◇ VMD ◇ PyMOL (Other MD Tools) ◇ MARTINI ◇ CHARMM-GUI • Free Energy Calculations using Enhanced Sampling : (Methods) ◇ Weighted Ensemble ◇ Metadynamics ◇ Umbrella Sampling ◇ Thermodynamic Integration (Tools) ◇ WESTPA ◇ PLUMED • Alchemical and Geometrical Binding Free Energy Calculations : ◇ BFEE2 • Docking : ◇ Schrödinger Maestro Glide • Quantum Mechanics Calculations : ◇ VASP ◇ Gaussian • Membrane, Membrane - GPCR, Membrane - RNA modeling • Small Molecule Parameterization • Clustering • SVMs

### Technical Expertise:

• Programming: Python, MATLAB, Fortran, C++, Bash • Others: Git - GitHub, Bitbucket, Jupyter Lab, AWS, MS Office Suite • System Biology Tools – Flux analysis and metabolic network reconstruction: COBRA, TIGER

## RESEARCH EXPERIENCE

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<b>Grad Research Student</b> University of Rochester Medical Center, Grossfield Lab	June 2019 - Present <i>Rochester, NY</i>
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- Spearheaded a method development project in a field new to the lab. Created new directions for the project that demanded mentoring, collaborating, and training of grad and undergrad lab members
- Developed a computational pipeline to estimate the free energy landscape of phase separation in the lipid bilayers ◇ Analyzed > 40 TB data ◇ Coarse-Grained Modeling ◇ GROMACS ◇ Thermodynamics ◇ Enhanced Sampling
- Characterizing the effects of collective variable decisions on enhanced sampling outcomes by creating a protocol that gauges collective variables before, after, and during the enhanced runs ◇ Benchmarking ◇ Optimization
- Characterizing the effects of box size in MD simulations on the thermodynamics of phase-separating systems
- Equilibrium partitioning of lipopeptides between two co-existing phases. ◇ Peptide - Membrane modeling
- Using Machine Learning methods in the pipeline to effectively search and rank a better set of collective variables
- Improving the efficiency of free energy calculation by coupling replica-exchange to weighted ensemble dynamics
- Tracking Ultrafast dynamics of Rhodopsin upon light activation to interpret the XFEL experiments ◇ Analyzed > 10000 short All-Atom simulations ◇ NAMD ◇ Cross-functional collaboration ◇ Membrane - GPCR modeling

<b>Computational Sciences Co-op</b> Moderna, Molecular Engineering and Modeling	Jan 2022 - June 2022 <i>Cambridge, MA</i>
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- Studied RNA – lipid bilayer interactions with NAMD All-Atom Molecular Dynamics simulations in AWS EC2
- Studied RNA - small molecule interaction via docking and Molecular Dynamics. Proposed and implemented a python analysis suite for initial validation of binding pocket interactions and ligand conformational dynamics
- Investigated and benchmarked alchemical and geometrical routes for RNA - small molecule binding free energy calculations. Implemented a pipeline for test molecules that can be extended for a large ligand library screening
- Proposed and initiated an internal pilot project that involved cross-functional teams

**Summer Research Fellow**  
JNCASR, Theoretical Sciences Unit

April 2018 - June 2018  
Bangalore, India

- Studied the sensitivity of population dynamics of bacteria towards the nutrient environment using modeling and simulation. Modeled quorum sensing and chemotaxis behaviors and emulated racing conditions between them

**Summer Research Fellow**  
JNCASR, Theoretical Sciences Unit

May 2017 - July 2017  
Bangalore, India

- Conducted metabolic network reconstructions and flux balance analyses on constrained-based *in silico* yeast models. Studied the effect of various metabolites on the yeast growth rates upon network perturbations
- Integrated transcriptomic data into *in silico* models of *Plasmodium Falciparum* to emulate multiple malaria variants in southeast Asia to understand the rise of drug resistance using cross-functional efforts

## OTHER RELEVANT EXPERIENCE

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**Moderna – Carnegie Mellon University, Artificial Intelligence (AI) – Academy Certification (2022):** Internal training to educate and empower Moderna employees to integrate AI and ML solutions into their workflow

**URBEST Trainee (2021 - Present)** University of Rochester initiative to Broaden Experiences in Scientific Training for early career scientists. Training on leadership and management skills for scientists via coursework, panel discussions, informational interviews, career stories, mentoring, and other personalized programs

**International Students and Scholars Advisory Board (ISSAB), University of Rochester (2022 - Present)** Represent the international grad student community in the School of Medicine and Dentistry at UR Medical Center

**Graduate Student Society, International student liaison (2022 - Present)** Formed an internal student committee and conducting focus groups and panel discussions to enhance the international grad student experience

**UR2 mentorship program (2020 - 2022)** A program run by grad students to mentor and train first-generation undergrads and those from less privileged backgrounds on tools and resources for a research career

**Teaching Assistant, BPH509 - Molecular Biophysics (2020)** Instructed students in theoretical, experimental, and computational methods to study macromolecules. Topics include statistical mechanics, optical melting experiments, dynamic programming algorithms, molecular dynamics, protein folding, and isothermal titration calorimetry

**Open source contributions (2020 - 2022)** **LOOS** : A lightweight object-oriented structure analysis library for MD simulations. **WESTPA** : The Weighted Ensemble Simulation Toolkit with Parallelization and Analysis

**General Secretary, Choreography Society, St.Stephen's College (2015 - 2016)** Headed a 60+ dance crew with Indian classical and fusion choreography and production. One major and minor production each semester

## SCHOLASTIC ACHIEVEMENTS

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**Biophysics Student Seminar Award, 2021** : For best student seminar based on popular votes

**Graduate Women In Science Mentor-Up Winner, 2021** : University challenge to set and achieve multiple short-term personal and professional goals through mentoring-up and training

**Neuman Travel Award, 2020, 2021** : To share and present the thesis research at academic conferences

**Dr. K. Swaminathan Memorial Award, 2016** : For an excellent academic record and overall achievements in cultural activities during undergrad level, St. Stephen's College, University of Delhi, India

**Top 1%** of all the students who appeared at the Higher Secondary Examination, March **2013**. Nominated by Kerala State Higher Secondary Board for the Central Government Scholarship for higher education

**INSPIRE SHE, 2013 -2018** : Five-year scholarship by the Department of Science and Technology, Government of India, based on an excellent performance at the higher secondary and undergrad level

**INSPIRE AWARD, 2011** : For the outstanding academic performance at the high school level, from the Department of Science and Technology, Government of India