Shantanu Khatri, Sr. Research Fellow

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Professional Summary

Computational Biologist with extensive experience in molecular dynamics simulations, protein structural modeling, and **high-performance computing**. Proficient in **multi-omics data analysis** and adept at designing and formulating **interdisciplinary** *in silico* **workflows**.

Education

Ph.D., Computational Biology

Council of Scientific and Industrial Research–Institute of Genomics and Integrative Biology, New Delhi, India (2020–Present)

Thesis: Understanding Conjugation Machinery in Human Autophagic System

Advisor: Dr. Lipi Thukral

Master of Science (M.Sc.), Biological Science (2013 - 2015)
 Hemvati Nandan Bahuguna Garhwal University, Uttarakhand, India

Technical Expertise

- Building and optimizing pipelines for multi-omics data integration and analysis, including proteomics, metabolomics, clinical data, epigenetics and genomics.
- Database management and web resource development for bioinformatics resources (RAPSAP).
- Scripting and automating computational workflows with Python, Unix, and Bash, utilizing libraries like
 Pandas, Scikit-learn, NumPy, Biopython and MDAnalysis.
- Large-scale compute with high-performance computing tools like SLURM and PBS.
- Containerization tools, Docker and Singularity on CPU/GPU (NVIDIA DGX Platform), executing scientific
 on AWS.
- Protein modeling with AI based pipelines AlphaFold, ProteinMPNN and RFdiffusion in HPC and performing protein-ligand docking with HADDOCK and AutoDock.
- Multiscale protein modeling and analyzing protein dynamics and interactions of high-order protein assemblies and protein-membrane systems using molecular dynamics simulation, GROMACS, CHARMM.
- Visualizing and presenting molecular structures using ChimeraX, PyMOL, VMD, and Blender.
- Free Energy Calculations (MM-PBSA, Poisson—Boltzmann Surface Area) and analyzing protein structural networks in high-order oligomers.
- Experimental design and validation techniques, including ITC and protein purification, to support and corroborate computational findings.
- Application of AI/ML and deep learning approaches for bioinformatics and biological data analysis.
- Proficient in ML workflows (PyTorch)

Research Projects

 Leveraged AlphaFold2-Multimer to predict high-order oligomeric assemblies regulating conjugation machinery in human autophagy and performed microsecond molecular simulations to analyse protein structural dynamics.

- Investigated structural differences in various functional forms of E1 enzyme ATG7 by leveraging AlphaFold to generate assemblies and performing molecular dynamics simulations to map critical interactions with key proteins such as LC3B, ATG12,ATG3 and ATG10. Validated these interactions through in-silico mutagenesis and invitro biophysical experiments (ITC).
- Predicted the bipartite membrane protein ATG2A, revealing its architecture, and identified dynamic transitions of cavity-lining residues through simulations, suggesting their critical role in regulating lipid transfer.
- Developed web-resource <u>RAPSAP</u> (Resource of AlphaFold2 Predicted Structures of Autophagy Pathway), a curated database providing comprehensive structural information on the complete human autophagy protein interactome.
- Conducted serological testing on voluntary participants as part of the Phenome India Cohort Project, utilizing Elecsys Anti-SARS-CoV-2 electrochemiluminescence immunoassay (ECLIA) on Cobas Immunoassay analysers for qualitative antibody detection followed by comprehensive data analysis and interpretation.

Publications

- Malhotra Nidhi*, Shantanu Khatri*, Ajit Kumar, Akanksha Arun, Purba Daripa, Saman Fatihi, Sureshkumar Venkadesan, Niyati Jain, and Lipi Thukral. "Al-based AlphaFold2 significantly expands the structural space of the autophagy pathway." Autophagy 19, no. 12 (2023): 3201-3220. *equal authors.
- Shantanu Khatri, Ekansh Jadone, Lipi Thukral, Deciphering ATG7's multifaceted structural landscape in human autophagic conjugation system, 2025. (In preparation)
- Mapping the functional terrain of E1-like ATG7: Insights into cross-functional roles, Shantanu Khatri,
 Lipi Thukral 2025. (In preparation)

Awards & Recognitions

- Council of Scientific & Industrial Research-National Eligibility Test: Junior Research Fellowship (June 2019)
- Council of Scientific & Industrial Research-National Eligibility Test: Lectureship (December 2018)
- Selected as Springer Nature Student Ambassador (2023)
- Recognition By Springer Nature Group for contribution in the 'Her Research, Our Future' Forum: 21
 March 2024

Conferences & Workshops

- Poster: 'Al-based AlphaFold2 significantly expands the structural space of the autophagy pathway'
 EMBO Workshop on Computational Structural Biology, EMBL, Heidelberg, Germany (Dec 2023)
- Poster: 'Autophagy pathway' 63rd Research Council Meeting, CSIR-IGIB, (Nov 2023)
- Delegate: Mini symposium on 'Latest in Autophagy and Lysosome Biology' at CSIR-IGIB: 12 January 2023
- Delegate: Symposium on 'Data Driven Approaches to Understand Biological Systems at Bioinformatics Centre', CSIR-IGIB, New Delhi: 29 April 2023

Courses/ Certificates

- Fundamentals of AI/ML in Precision Medicine: Course by the Department of Genetics, Stanford Medicine, covering topics such as machine learning algorithms, multiomics data integration, predictive modeling, and AI applications in healthcare.
- Fundamentals of Data Science in Precision Medicine and Cloud Computing: Course by the Department of Genetics, Stanford Medicine, focused on multi-omics data analysis, cloud-based workflows, computational pipelines, and data-driven insights in precision medicine.
- Deep learning with Pytorch , python-first framework.

Volunteer Work & Leadership

- Nodal Officer: AWS Workshop-Biology on Cloud at CSIR-IGIB, New Delhi, 6 November 2023.
- Co-Organizer: Science Entrepreneurship Competition (IGIB BIG Ideas), November 2023
- Project Coordinator: 'Bringing Genomics Closer to Society' an audiovisual project as a part of One Week, One Lab at CSIR-IGIB.
- Organizer: Interactive Discussion on 'Women in STEM' as part of Springer Nature IWD 2024
- Volunteered in International Workshop," Applications of AI and Data-Driven Approaches in Structural Biology 2024, CSIR-IGIB.

Extra-Curricular Activities

- Active participation in institutional-level tournaments for table tennis, cricket, and badminton, with awards including Runner-up in National CSIR-SSBMT Table Tennis Tournament
- Fitness Drive 2022 Runner-up & Fitness Drive 2024 Winner in Table Tennis Tournament, CSIR-IGIB.

Languages

- English (Fluent/Professional Proficiency)
- Hindi (Native Proficiency)