

Parth Bibekar

Curriculum Vitae

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Education

- 2024–Present **Institute of Bioengineering, SV, EPFL, Switzerland,**
PhD in Computational and Quantitative Biology.
Advisor: Prof. Matteo Dal Peraro
- 2019–2024 **Indian Institute of Science Education and Research, Kolkata, India,**
BSMS Dual Degree, Major in Biology & minor in Computational and Data Sciences.
CGPA – 8.21/10

Research Interests

Research Int. Bioinformatics, Molecular modeling, Computational Biophysics

Current Position

- Sept 2024–
Ongoing I am an incoming PhD student at the [Laboratory for Biomolecular modeling](#) at EPFL, Switzerland, under the supervision of Prof. Matteo Dal Peraro. Developing ML based techniques for various applications in biomolecular modeling.

Publications

- Mar 2024 Bibekar et al., PeSTo-Carbs: geometric deep learning for prediction of protein-carbohydrate binding interfaces. *Journal of Chemical Theory and Computation* (2024), [10.1021/acs.jctc.3c01145](https://doi.org/10.1021/acs.jctc.3c01145)
- Feb 2023 Dutta et al., Conformational ensemble of the NSP1 CTD in SARS-CoV-2: Perspectives from the free energy landscape, *Biophysical Journal* (2023), [10.1016/j.bpj.2023.02.010](https://doi.org/10.1016/j.bpj.2023.02.010)

Projects

- Sept 2024 –
Ongoing **Ribonucleic acid Sequence design from TerTiary structure (RISoTTto), EPFL Lausanne, Switzerland.**
- Developed RISoTTto, a novel model capable of predicting RNA sequences from their backbone scaffolds.
 - RISoTTto was the primary focus of my MS thesis, pioneering context-aware RNA sequence design by considering surrounding proteins, ligands, or ions.
- June 2022 –
Sept 2024 **Predicting Protein - Carbohydrate binding interfaces using Deep Learning, Prof. Matteo Dal Peraro, EPFL Lausanne, Switzerland.**
- Predicting binding interfaces between Proteins and Carbohydrates using deep geometric transformers.
 - Using transfer learning to specialize on carbohydrates with great pharmaceutical importance.

- Nov 2021–Oct 2022 **Exploring the the Free Energy Landscape of IDRs using Machine Learning**, Prof. Neelanjana Sengupta, IISER Kolkata, India. DOI:[10.1016/j.bpj.2023.02.010](https://doi.org/10.1016/j.bpj.2023.02.010).
- Study the Conformational Landscape of two SARS-CoV-2 IDRs using Molecular Dynamics Simulation and fitting FEL using GMM based models.
 - Sample the intermediate rare Conformations using the data driven latent space

- June 2021 – Sept 2021 **Exploring the Free Energy Landscape of amyloid assembly**, Prof. Neelanjana Sengupta, IISER Kolkata, India.
- Study the Free Energy Landscape of hIAPP fibril dissociation using Molecular Dynamics Simulations and enhance sampling methods.

- Apr 2021 **WeLearn Bot [Code]**, Team Project.
- I created and maintain a cli application for interacting with WeLearn in the IISER-Kolkata domain.
 - The application is widely used by many students in IISER Kolkata and has been released as a Python package.

Find my other projects here [Github: ParthBibekar](#).

Achievements

- Aug 2023 **E3 Program Fellow**.
- I have been awarded the EPFL Excellence in Engineering Program fellowship, for a summer project in the Laboratory for Biomolecular Modeling at EPFL, Switzerland.
- Jun 2022 **ThinkSwiss Research Scholarship**.
- I was awarded the ThinkSwiss Research Scholarship of for a period of three months for a research project in École polytechnique fédérale de Lausanne, Switzerland.
 - I worked with Prof. Matteo Dal Peraro on a Biomolecular modeling project to predict Protein - Carbohydrate interfaces.
- Dec 2021 **National HPC Hackathon**.
- Part of the winning team in the National HPC Hackathon organised by Amazon Web Services and Intel in association with the Office of Principal Scientific Advisor to the Government of India.

Skills

- Languages Python, C, MATLAB, Bash
- Frameworks TensorFlow, Keras, PyTorch, PyTorch Geometric
- Molecular Dynamics GROMACS, AMBER, PLUMED
- HPC AWS, GCC, etc.
- Data analysis Gnuplot, OriginLab, Grace
- Utilities Conda/Anaconda, Git/Github, VScode, Jupyter Notebook
- Web Dev HTML/CSS
- OS Linux, Windows