

# SUMIT TARAFDER

✉ [sumittarafder@vt.edu](mailto:sumittarafder@vt.edu)

🔗 [sumittarafder.github.io](https://sumittarafder.github.io)

🌐 [github.com/SumitTarafter](https://github.com/SumitTarafter)

## Research Interests

---

**Topics** 3D structure prediction, refinement or quality estimation of protein, RNA and associated complexes  
**Areas** Structural bioinformatics, deep learning

## Education

---

### Virginia Tech

*Ph.D. student in Computer Science*

**Aug. 2022 – Present**

*Blacksburg, VA*

### Bangladesh University of Engineering & Technology (BUET)

*M.Sc. in Computer Science and Engineering*

**Jan. 2022**

*Dhaka, Bangladesh*

### Bangladesh University of Engineering & Technology (BUET)

*B.Sc. in Computer Science and Engineering*

**Feb. 2017**

*Dhaka, Bangladesh*

## Experience

---

### Virginia Tech

*Graduate Research Assistant, Bhattacharya Lab*

**Aug. 2022 – Present**

*Blacksburg, VA*

- Focused on projects related to the prediction and quality estimation of RNA 3D structures and protein-RNA complexes.
- Recently implemented research projects are available in *GitHub*.

### Virginia Tech

*Graduate Teaching Assistant, CS 5914 -TS: Warehouse Scale Computing*

**Aug. 2022 – Dec. 2022**

*Blacksburg, VA*

- Graded multiple paper review and summary writings for individual students and conducted office hours.

### United International University

*Lecturer, Department of Computer Science and Engineering*

**May 2017 – Oct. 2021**

*Dhaka, Bangladesh*

- I was the instructor and coordinator for a wide range of undergraduate-level theory and practical courses.

## Publications

---

- **Sumit Tarafder**, Debswapna Bhattacharya. “*RNAbpFlow: Base pair-augmented SE(3)-flow matching for conditional RNA 3D structure generation*”, **2025**. [Under Review]
- **Sumit Tarafder**, Debswapna Bhattacharya. “*lociPARSE: a locality-aware invariant point attention model for scoring RNA 3D structures*”, Journal of Chemical Information and Modeling, Volume 64, Issue 22, Pages 8655–8664, **November 2024**. Impact Factor: 5.7
- **Sumit Tarafder**, Rahmatullah Roche, Debswapna Bhattacharya. “*The landscape of RNA 3D structure modeling with transformer networks*”, Biology Methods and Protocols, Volume 9, Page 47, Issue 1, **July 2024**. Impact Factor: 2.5
- **Sumit Tarafder**, Mazharul Islam, Swakkhar Shatabda, Atif Rahman. “*Figbird: A probabilistic method for filling gaps in genome assemblies*”, Bioinformatics, Volume 38, **August 2022**, Pages 3717–3724, Issue 15. Impact factor: 6.9
- **Sumit Tarafder**, Md. Toukir Ahmed, Sumaiya Iqbal, Md Tamjidul Hoque, M. Sohel Rahman. “*RBSURFpred : Modeling Protein Accessible Surface Area in Real and Binary Space using Regularized and Optimized Regression*”, Journal of Theoretical Biology, Volume 441, **January 2018**, Pages 44 - 57. Impact factor: 2.0
- Rahmatullah Roche, **Sumit Tarafder**, Debswapna Bhattacharya. “*Single-sequence protein-RNA complex structure prediction by geometric attention-enabled pairing of biological language models*”, **2024**. [Under Review]
- Rahmatullah Roche, Bernard Moussad, Md Hossain Shuvo, **Sumit Tarafder**, Debswapna Bhattacharya. “*EquiPNAS: improved protein-nucleic acid binding site prediction using protein-language-model-informed equivariant deep graph neural networks*”, NAR, volume 52, Page e27, Issue 5, **January 2024**. Impact Factor: 14.9

## Talks

---

- Presented our work “*lociPARSE*” as a poster at *ISMB '24* (Montreal, CA)
- Going to present “*lociPARSE*” as a short talk at the upcoming *GLBIO conference* (Minnetota, USA)

## Awards

---

<b>Travel Grant</b>	Awarded by Department of Computer Science, Virginia Tech for <i>ISMB '24</i>
<b>Travel Grant</b>	Awarded by ISCB and GLBC to present my paper at <i>GLBIO '25</i>
<b>Fellowship</b>	Pratt Fellowship from Department of Computer Science, Virginia Tech in <i>'24</i>
<b>Dean's List Award</b>	University Merit Scholarship & Dean's List from <i>BUET</i>

## Services

---

- Reviewer at IEEE/ACM Transactions on Computational Biology and Bioinformatics
- Sub-reviewer at ISMB 2025, IEE BIBM 2024, BIOKDD 2024

## Skills

---

**Languages:** Python, Java, C, C++, HTML/CSS

**Tools:** Git, Docker, Chimera, PyMOL, VS Code

**Frameworks:** PyTorch, PyTorch Lightning, Matplotlib, Scikit-learn, LangChain