

EDUCATION	<b>The Pennsylvania State University</b> <i>Ph.D. Candidate, Computer Science and Engineering</i> <ul style="list-style-type: none"> <li>• Advisor: Dr. Mingfu Shao</li> <li>• Research area: My research interests are in Bioinformatics and Computational Biology. Specifically, I work with short- and long-read RNA-seq data to improve the accuracy and resolution of circular RNA assembly.</li> </ul>	Pennsylvania, USA Aug 2021 - Present
	<b>Bangladesh University of Engineering and Technology (BUET)</b> <i>BS.c. in Computer Science and Engineering</i> <ul style="list-style-type: none"> <li>• CGPA: 3.90/4.00</li> </ul>	Dhaka, Bangladesh Feb 2015 - Apr 2019
WORK EXPERIENCE	<b>The Pennsylvania State University</b> <i>Graduate Research/Teaching Assistant</i>	Pennsylvania, USA Aug 2021 - Present
	<b>Ahsanullah University of Science and Technology (AUST)</b> <i>On study leave, Department of Computer Science and Engineering</i>	Dhaka, Bangladesh July 2019 - July 2021
	<b>Dhanmondi Tutorial</b> <i>Teacher</i>	Dhaka, Bangladesh Jan 2019 - April 2019
PUBLICATIONS	<ol style="list-style-type: none"> <li>1. <b>Tasfia Zahin</b>, Irtesam Mahmud Khan, Mingfu Shao. Accurate Reconstruction of Circular RNAs from Complex Rolling Circular Long Reads with CircPlex. <i>bioRxiv</i>, Nov 2025. <a href="https://doi.org/10.1101/2025.11.21.689841">https://doi.org/10.1101/2025.11.21.689841</a></li> <li>2. Zhezheng Song*, <b>Tasfia Zahin*</b>, Xiang Li, Mingfu Shao. Accurate Detection of Tandem Repeats from Error-Prone Sequences with EquiRep. <i>Genome Research</i>, December 2025 35: 2714-2721.</li> <li>3. Zhezheng Song*, <b>Tasfia Zahin*</b>, Xiang Li, Mingfu Shao. Accurate Detection of Tandem Repeats from Error-Prone Sequences with EquiRep. <i>Proceedings of 29th International Conference on Computational Molecular Biology (RECOMB 2025). Lecture Notes in Computer Science</i>, vol 15647, pp. 390–394. Springer, Cham, 2025.</li> <li>4. <b>Tasfia Zahin</b>, Qian Shi, Carl Zang, Mingfu Shao. Accurate Assembly of Circular RNAs with TERRACE. <i>Genome Research</i>, September 2024 34: 1365-1370.</li> <li>5. <b>Tasfia Zahin</b>, Qian Shi, Carl Zang, Mingfu Shao. Accurate Assembly of Circular RNAs with TERRACE. <i>Proceedings of 28th International Conference on Computational Molecular Biology (RECOMB 2024). Lecture Notes in Computer Science</i>, vol 14758, pp. 444–447. Springer, Cham, 2024.</li> <li>6. Xiaofei Carl Zang, <b>Tasfia Zahin</b>, Irtesam Mahmud Khan, Qian Shi, Yi Xing, Mingfu Shao. Amaranth: Enhanced Single-Cell Transcript Assembly via Discriminative Modeling of UMI Reads and Internal Reads. <i>bioRxiv</i>, Nov 2025. <a href="https://doi.org/10.1101/2025.11.24.690228">https://doi.org/10.1101/2025.11.24.690228</a></li> <li>7. Irtesam Mahmud Khan, Xiaofei Carl Zang, Ange Teng, <b>Tasfia Zahin</b>, Mingfu Shao. Boosting Transcript Assembly via Delineating Transcript Start and End Sites. <i>bioRxiv</i>, Oct 2025. <a href="https://doi.org/10.1101/2025.10.13.682211">https://doi.org/10.1101/2025.10.13.682211</a></li> <li>8. <b>Tasfia Zahin*</b>, Md Hasin Abrar*, Mizanur Rahman Jewel, Tahrina Tasnim, Md Shamsuzzoha Bayzid, Atif Rahman. An alignment-free method for phylogeny estimation using maximum likelihood. <i>BMC bioinformatics</i>, 2025 Mar 7;26(1):77.</li> </ol>	
ACADEMIC SERVICES	<ul style="list-style-type: none"> <li>• Sub-reviewer of <i>ISMB 2026</i>, <i>RECOMB 2026</i>, <i>RECOMB 2025</i>, <i>WABI 2025</i>, <i>RECOMB 2024</i>, <i>ISMB 2024</i>, <i>WABI 2024</i>, <i>WABI 2023</i>, <i>RECOMB 2022</i>, <i>ACM-BCB 2022</i>.</li> <li>• Volunteer at the 2024 Workshop on Emerging Methods for Sequence Analysis (WEMSA 2024) organized by The Pennsylvania State University.</li> </ul>	

CONFERENCE PARTICIPATION	<ul style="list-style-type: none"> <li>• Research and poster presenter at RECOMB 2025, Seoul, South Korea.</li> <li>• Research presenter at RECOMB 2024, Boston, Massachusetts, USA.</li> <li>• Research presenter at the 2023 Workshop on Emerging Methods for Sequence Analysis (WEMSA 2023) organized by The Pennsylvania State University.</li> <li>• Attendee of ISMB 2022, Madison, Wisconsin, USA.</li> <li>• Attendee of RECOMB 2020 Virtual Conference.</li> </ul>
TEACHING EXPERIENCES	<ul style="list-style-type: none"> <li>• Teaching assistant for CMPSC 465 Data Structures and Algorithms, Fall 2021, The Pennsylvania State University.</li> <li>• Teaching assistant for CMPSC 465 Data Structures and Algorithms, Fall 2024, The Pennsylvania State University.</li> <li>• Instructor for CSE 2186 Basic Programming Techniques, Fall 2020, Ahsanullah University of Science and Technology.</li> <li>• Instructor for CSE 1287 and CSE 1288 Computer Programming, CSE 2146 Introduction to Computer Systems, Spring 2020, Ahsanullah University of Science and Technology.</li> <li>• Instructor for CSE 3107 and CSE 3108 Microprocessors, Spring 2020, Ahsanullah University of Science and Technology.</li> <li>• Instructor for CSE 3107 and CSE 3108 Microprocessors, CSE 2186 Basic Programming Techniques Fall 2019, Ahsanullah University of Science and Technology.</li> <li>• Instructor for CSE 3107 and CSE 3108 Microprocessors, Spring 2019, Ahsanullah University of Science and Technology.</li> </ul>
AWARDS AND HONORS	<ul style="list-style-type: none"> <li>• Winner of Huawei Seeds for the Future, 2019, Travel Grant by Huawei Technologies.</li> <li>• Dean's List Honor, 2015-2019, Bangladesh University of Engineering and Technology (BUET).</li> <li>• Daily Star O and A Level Award of Excellence, 2012 and 2014, The Daily Star Newspaper, Bangladesh.</li> </ul>
REFERENCE	<p><b>Dr. Mingfu Shao</b>  <i>Associate Professor</i>  Department of Computer Science and Engineering  The Pennsylvania State University  W205A Westgate Building  University Park, PA 16802  Email: mxs2589@psu.edu</p> <p><b>Dr. Atif Hasan Rahman</b>  <i>Associate Professor</i>  Department of Computer Science and Engineering  Bangladesh University of Engineering and Technology (BUET)  ECE Building, West Palashi, Dhaka-1000, Bangladesh  Email: atif@cse.buet.ac.bd</p>