

EDUCATION	<p><b>The Pennsylvania State University</b> Pennsylvania, USA  <i>Ph.D. Candidate, Computer Science and Engineering</i> Aug 2021 - Present</p> <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> <p><b>Bangladesh University of Engineering and Technology (BUET)</b> Dhaka, Bangladesh  <i>BS.c. in Computer Science and Engineering</i> Feb 2015 - Apr 2019</p> <ul style="list-style-type: none"> <li>• CGPA: 3.90/4.00</li> </ul>
WORK EXPERIENCE	<p><b>The Pennsylvania State University</b> Pennsylvania, USA  <i>Graduate Research/Teaching Assistant</i> Aug 2021 - Present</p> <p><b>Ahsanullah University of Science and Technology (AUST)</b> Dhaka, Bangladesh  <i>On study leave, Department of Computer Science and Engineering</i> July 2019 - July 2021</p> <p><b>Dhanmondi Tutorial</b> Dhaka, Bangladesh  <i>Teacher</i> Jan 2019 - April 2019</p>
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. Proceedings of <i>29th International Conference on Computational Molecular Biology (RECOMB 2025)</i>. <i>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. Proceedings of <i>28th International Conference on Computational Molecular Biology (RECOMB 2024)</i>. <i>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 - ISMB 2026, RECOMB 2026, RECOMB 2025, WABI 2025, RECOMB 2024, ISMB 2024, WABI 2024, WABI 2023, RECOMB 2022, ACM-BCB 2022.</li> <li>• Volunteer - The 2024 Workshop on Emerging Methods for Sequence Analysis (WEMSA 2024), The Pennsylvania State University, Pennsylvania, USA.</li> </ul>

- CONFERENCE PARTICIPATION
- Research and poster presenter - RECOMB 2025, Seoul, South Korea.
  - Research presenter - RECOMB 2024, Boston, Massachusetts, USA.
  - Research presenter - The 2023 Workshop on Emerging Methods for Sequence Analysis (WEMSA 2023), The Pennsylvania State University, Pennsylvania, USA.
  - Attendee - ISMB 2022, Madison, Wisconsin, USA.
  - Attendee - RECOMB 2020 Virtual Conference.
- TEACHING EXPERIENCE
- Teaching assistant - The Pennsylvania State University, Pennsylvania, USA.
    - CMPSC 465 Data Structures and Algorithms, Fall 2021, Fall 2024.
  - Instructor - Ahsanullah University of Science and Technology, Dhaka, Bangladesh.
    - CSE 3107 and CSE 3108 Microprocessors, Spring 2019, Fall 2019.
    - CSE 2186 Basic Programming Techniques, Fall 2020, Fall 2019.
    - CSE 1287 and CSE 1288 Computer Programming, Spring 2020.
    - CSE 2146 Introduction to Computer Systems, Spring 2020.
- PROFESSIONAL AND STUDENT ENGAGEMENT
- IT support sub-committee member - The 23rd International Conference on Computer and Information Technology (ICCIT) 2020, Ahsanullah University of Science and Technology.
  - Executive committee member - Robotics Club, Ahsanullah University of Science and Technology, 2020-2021.
  - Mentor - Bangladeshi Women in Computer Science and Engineering (BWCSE) Mentor-mentee program, Bangladesh University of Engineering and Technology, 2020-2021.
  - Media and publication secretary - English Medium Buetians' Association (EMBA), Bangladesh University of Engineering and Technology, 2018-2019.
  - Adjudicator - BRAC Inter-University Debate Competition 2016.
  - Delegate - Bangladesh Youth Leadership Summit 2016.
  - Member - Bangladeshi Women in Computer Science and Engineering (BWCSE), Bangladesh University of Engineering and Technology, 2015-2019.
  - Member - Debating Club, Bangladesh University of Engineering and Technology, 2015-2017.
- AWARDS AND HONORS
- Winner, Huawei Seeds for the Future 2019, Travel Grant by Huawei Technologies.
  - Dean's List Honor, 2015-2019, Bangladesh University of Engineering and Technology.
  - Daily Star O and A Level Award of Excellence, 2012 and 2014, The Daily Star Newspaper, Bangladesh.
- REFERENCE
- Dr. Mingfu Shao**  
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