

# Tamjeed Azad

<https://tamjeedazad.com> | [github.com/tamjazad](https://github.com/tamjazad) | [linkedin.com/in/tamjazad](https://linkedin.com/in/tamjazad)

Email: tamjeeda@princeton.edu

Last Updated February 2026

## Education

---

- **Princeton University** Princeton, NJ  
PhD in Computer Science (Anticipated). August 2022 - Present
  - Advised by Prof. Yuri Pritykin.
  - Incidental MA in Computer Science conferred May 2024.
- **Columbia University** New York, NY  
BS in Computer Science, Minor in Economics. GPA: 3.99 August 2018 - May 2022
  - Degree conferred Magna Cum Laude.
  - Advised by Prof. Elham Azizi.

## Research Interests

---

- single-cell transcriptomics, cell-cell interactions, spatial transcriptomics, immunology

## Honors

---

- Cummins Merit Fellowship Recipient, Princeton University, 2022-2023.
- Allen School CSE PhD Research Fellowship, University of Washington, 2022-2023 (declined).
- Tau Beta Pi Honor Society Member since November 2020.
- NSF REU Participant, University of Tennessee at Chattanooga, iCompBio 2020.
- Columbia Summer Undergraduate Research Fellowship (SURF) 2019.
- CP Davis Scholar, Columbia University Scholars Program (CUSP), 2018-2022.
- Columbia University Dean's List all eligible semesters, 2018-2022.
- National Merit Scholar (Tennessee), 2018.

## Publications & Preprints

---

- **Azad, T.**, Walker, S.K., Victora, G.D., Pritykin, Y. (2026). Predictive modeling of molecular activity underlying physical cell-cell interactions. *Cell Reports Methods*.  
<https://doi.org/10.1016/j.crmeth.2026.101301>
- Sun, A., **Azad, T.**, Brown, C., Pritykin, Y. (2026). Benchmarking cell type deconvolution in spatial transcriptomics and application to cancer immunotherapy. *BioRxiv*.  
<https://doi.org/10.64898/2026.01.13.699379>
- Shi, L., Uzuni, A., Wang, X.K., . . . , **Azad, T.**, . . . , Azizi, E., Reshef, R. (2025). Spatiotemporal Single-Cell Analysis Reveals T Cell Clonal Dynamics and Phenotypic Plasticity in Human Graft-versus-Host Disease. *BioRxiv*. <https://doi.org/10.1101/2025.05.24.655962>
- Camara, A., Taneja, N., **Azad, T.**, Allaway, E., Zemel, R. (2022). Mapping the Multilingual Margins: Intersectional Biases of Sentiment Analysis Systems in English, Spanish, and Arabic. *Proceedings of the Second Workshop on Language Technology for Equality, Diversity and Inclusion*, 90–106, Association for Computational Linguistics. Oral Presentation.  
<http://dx.doi.org/10.18653/v1/2022.ltedi-1.11>

- **Azad, T.**, Alonzo, S. H., Bonsall, M. B., Klug, H. (2022). Life history, mating dynamics and the origin of parental care. *Journal of Evolutionary Biology*, 35, 379–390.  
<https://doi.org/10.1111/jeb.13959>
- Chien, T., Harimoto, T., Kepcs, B., Gray, K., Coker, C., Hou, N., Pu, K., **Azad, T.**, Nolasco, A., Pavlicova, M., Danino, T. (2022). Enhancing the tropism of bacteria via genetically programmed biosensors. *Nature Biomedical Engineering*. 6, 94–104.  
<https://doi.org/10.1038/s41551-021-00772-3>

## Presentations

---

- (Oral Presentation) **Azad, T.**, Qin, H. (2020) Recurrent neural networks for COVID-19 prediction: A cross-study with historical flu data and temperature data. *12th Annual National Institute for Mathematical and Biological Synthesis Undergraduate Research Conference*.

## Professional Experience

---

- **Software Development Engineer Intern** Amazon  
Items and Offers Platform, Amazon Selection and Catalog Systems.      *Summer 2022, Summer 2021*
  - Added internal authentication + user analytics to Offer Service Buyability Analyzer Tool.
  - Used the AWS CDK to create an end to end prototype log monitoring application for OfferService.
- **Research Assistant, Azizi Lab** Columbia University  
Research in single-cell transcriptomics.      *February 2020 - May 2022*
  - Investigated generative modeling for scRNA-seq data using ensembles of variational autoencoders.
  - Analyzed immune cell dynamics in acute lymphoblastic leukemias using scRNA-seq data.
  - Analyzed patterns of gene expression across T cell lineages in Growth v Host Disorder patients using paired scRNA-seq/scTCR-seq data.
- **Summer Research Intern, Qin Lab** University of Tennessee at Chattanooga  
Interned through iCompBio REU 2020.      *May 2020 - July 2020*
  - Constructed RNN-based models for regional COVID-19 case prediction using historical flu and temperature data.
- **Research Assistant, Synthetic Biological Systems Lab** Columbia University  
Investigated engineering bacterial biosensors.      *October 2018 - August 2019*
  - Engineered bacteria that could selectively grew and fluoresced in a low pH environment.
- **Research Assistant, Klug Lab** University of Tennessee at Chattanooga  
Computational research in evolutionary ecology.      *November 2016 - August 2018*
  - Investigated link between life history traits, mating dynamics, and parental care's evolution through exploratory computational analyses.

## Teaching Experience

---

- **Graduate Teaching Fellow** Princeton University  
At the McGraw Center for Teaching and Learning.      *Since August 2024*
  - Help lead Assistant in Instruction (AI) Orientation for new AIs at Princeton.
  - Help enrich and support teaching initiatives on campus.

- **Graduate Teaching Assistant** Princeton University  
TA in the computer science department. September 2023 - May 2024
  - (Spring 2024) TA for Artificial Intelligence for Precision Health.
  - (Fall 2023) TA for Introduction to Computational Molecular Biology.
- **Undergraduate Teaching Assistant** Columbia University  
TA in the computer science and electrical engineering departments. September 2020 - May 2022
  - (Spring 2022, Fall 2021, Spring 2021) Course Assistant for Fundamentals of Computer Systems.
  - (Spring 2021, Fall 2020) Lab Assistant for Intro to Electrical Engineering.

## Other Projects

---

- **PolyWiz** (Spring 2021) Programming language designed for a 5-person team project in Programming Languages and Translators class. Language uses C-like syntax and has extensive native support for polynomial computing and plotting. Translator architecture written using OCaml and C. Source code and project report accessible on GitHub profile.

## Technical Skills

---

- **Languages:** Python, R, Java, MATLAB, C/C++, OCaml, TypeScript, JavaScript

## Service

---

- Reviewer: Journal of Computational Biology and ACM BCB (2025)  
Subreviewer: ISMB (2024, 2025) and RECOMB (2024 - 2026)

## Community Activities

---

- **Writer, Princeton Insights** Princeton University  
Highlighting Princeton research in accessible and short reviews. February 2023 - August 2023
  - Article: *Belayer: Modeling Layered Tissues in Spatial Transcriptomics*.  
<https://insights.princeton.edu/2023/08/modeling-spatial-transcriptomics/>
- **Student Ambassador, Princeton University Concerts** Princeton University  
Help organize/advertise invited concert performers on campus. January 2024 - August 2024
- **Senior Advisor, Club Zamana** Columbia University  
Umbrella organization for Columbia's South Asian community. 2021-2022 School Year
  - Media Chair, 2019-2020 School Year; Organizational Committee Member, 2018-2019 School Year.
- **E-Board Member, Columbia Science Review** Columbia University  
Hosts science-focused events and publishes a science-focused magazine. 2019 Spring Semester