

EEMAN ABBASI

☎ (413) 313-8879 ✉ eabbasi@sas.upenn.edu 🌐 <https://eemanabbasi.github.io/profile/>

Education

University of Pennsylvania

Ph.D. Candidate, Biological Sciences, GPA: 3.98

expected Nov 2024

Philadelphia, PA

Mount Holyoke College

B.A., magna cum laude, Computer Science and Biology, GPA: 3.90

Sep 2015 – May 2019

South Hadley, MA

Recipient of DAAD Rise Fellowship

Research Experience

Data Science Intern | [Concerto Biosciences](#)

April 2024 - Present

- Analyzed **fluorescence-tagged image-based** datasets to uncover biological signatures and assess microbial synergy through **Python**-based data analysis, leveraged **AWS** for data processing and storage.
- Conducted **bioinformatic analysis** on **WGS** and **16s** microbial data, and identified key microbial candidate combinations for disease indications. Analyses contributed to product development initiatives, and presented key insights to company leadership.
- Collaborated with the Senior Data Scientist and the experimental discovery team to enhance data filtering methods and develop new **machine learning** approaches for optimizing the data discovery process.

Ph.D. Candidate, [Akçay Lab](#) | UPenn

Sep 2019 – Present

- Developed mathematical models and applied statistical techniques to investigate microbial interactions and the role of the immune response in shaping the microbiome. Key research contributions include:
 - * Microbiome-Immune ODE Model: Developed an ODE model in **Python** to simulate within-microbiome interactions and the impact of the host immune response. Insights from this work were selected for presentation at the [Young AMICI Symposium](#).
 - * Multi-omics Analysis of Cancer Microbiome Data: Applied advanced statistical techniques and bioinformatics tools in **Python** and **R** for **predictive modeling**, and **metabolic modeling**. Utilized large scale **TCGA** cancer microbiome datasets (both **16S** and **WGS**) and immune **RNA-seq** expression data to explore the impact of microbial features on patient outcomes. Identified microbial richness and abundance as significant predictors of patient survival and prognosis.
 - * Agent-Based Simulation of Microbiome Exchange across social networks: Developed a **Python-based agent-based model** simulating microbiome exchanges between individuals in environments with varying pathogen pressures. This model examined how social interactions and pathogen exposure influence microbiome composition and social network modularity.

Research Intern | [NIMBioS](#)

May 2018 - Dec 2020

- Developed a **discrete time population dynamics model** in **R** for simulating plant-seed disperser dynamics in hunting and harvesting regimes
- Performed sensitivity analysis using real-world parameter ranges to identify the most critical parameter of plant and seed disperser growth for targeted conservation efforts
- Collaborated and co-authored a publication with five researchers and presented findings to over 150 attendees at the Doctoral Consortium on Computational Sustainability

Women Summit Fellow, Data Science for all | [Correlation One](#)

Aug 2020 - Oct 2020

- Partnered with four women summit fellows to quantify the impact of public sentiment in New York and California on COVID-19 case counts and mobility patterns

Leadership, and Community Engagement

Probono Consultant, Penn Biotech Group Healthcare Consulting | Wharton

Aug 2022 – May 2024

- Collaborated with a team of nine professionals to perform market analysis and recommend 100+ potential clients to a health-tech company. Conducted patient journey analysis for a gene therapy client to identify treatment experiences
- Collaborated with a team of four professionals as part of the mini capstone MBA project to conduct epidemiology analysis, asset evaluation, and an overview of the competitive landscape, to evaluate the potential acquisition of a biotech company.

Career and Wellness Coordinator, Biology Graduate Group | UPenn

Aug 2022 - May 2023

- Planned and executed a department-wide career seminar series and two wellness workshops, catering to more than 25 individuals in the biology graduate community

Graduate Resident Advisor | UPenn

Aug 2022 - Present

- Led emergency responses and provided support to 24 students. Coordinated a social programming committee for the entire hall and executed 12 successful events for 500 house residents

Teaching Assistant | UPenn

Aug 2020 - May 2023

- Designed and implemented course curriculum and provided mentorship for six core courses in ecology, environmental science, and evolution. Led weekly recitation sessions, and taught microbial ecology as a guest speaker

Additional Information

Languages and Frameworks: Java, Julia, Python (including packages such as pandas, numpy, scipy, biopython, scikit-learn, statsmodels, matplotlib, seaborn, plotly, scikit-bio, and scikit-learn), R, Unix, Linux, AWS, Slurm Scheduler

Interests: Running, hiking, zumba, and interested in increasing participation and retention in Environmental Data Science community

Publications

- **Abbasi, E.**, Akcay, E. Host control and species interactions jointly determine microbiome community structure. (2024) . *Theoretical Population Biology*, <https://doi.org/10.1016/j.tpb.2024.06.006>
- **Abbasi, E.**, The Role of Competition and Mutualism in Microbiome and Host Immune Interactions.(2024) Manuscript in prep.
- **Abbasi, E.**, The Impact of Social Microbial Transmission on Host Immunity and Network Structure. (2024) Manuscript in prep.
- Fong, C., Galaz, C., **Abbasi, E.**, Gubbins, N., and Jouzi, Z. Broadening participation in environmental data science: insights from practitioners. (2024). *Environmental Data Science*, Accepted, waiting to be published
- Kevin De Angeli, **Eeman Abbasi**, Alan Gan, Daniel J. Ingram, Xingli Giam, Charlotte H. Chang. (2021). Modeling the impact of wild harvest on plant–disperser mutualisms: Plant and disperser co-harvest model. *Ecological Modelling*, 439. <https://doi.org/10.1016/j.ecolmodel.2020.109328>