

CONTACT INFORMATION	Department of Statistics and Data Science Carnegie Mellon University Pittsburgh, PA USA	website: aerosengart.github.io email: arosenga@andrew.cmu.edu
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SUMMARY I am a Ph.D. student in the Department of Statistics & Data Science at Carnegie Mellon University. I primarily do work related to analysis of biomedical data. Currently, I am focused on statistical methods for wastewater-based epidemiology.

EDUCATION	<p>Current Ph.D. Statistics, Carnegie Mellon University, Pittsburgh, PA USA <i>Advised by F. William Townes</i></p> <p>2024 M.S. Statistics, Carnegie Mellon University, Pittsburgh, PA, USA</p> <p>2021 B.S. Data Science, University of Michigan, Ann Arbor, MI USA</p> <p>2021 B.F.A. Jazz Studies, University of Michigan, Ann Arbor, MI USA</p>
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RESEARCH

Preprints

- [1] **Rosengart AL**, Bidwell AL, Wolfe MK, Boehm AB, Townes FW. Spatio-temporal variability of the Pepper Mild Mottle Virus biomarker in wastewater (2024). <https://arxiv.org/abs/2408.12012>.

Peer-Reviewed Journal Articles

- [1] Rahimikollu J, Xiao H, **Rosengart A**, Rosen ABI, Tabib T, Zdinak PM, He K, Bing X, Bunea F, Wegkamp M, Poholek A, Joglekar A, Lafyatis R, Das J. SLIDE: Significant Latent Factor Interaction Discovery and Exploration across biological domains. *Nature Methods* (2024). doi:10.1038/s41592-024-02175-z.
- [2] Wheeler J, **Rosengart A**, Jiang Z, Tan K, Treutle N, Ionides EL. Informing policy via dynamic models: Cholera in Haiti. *PLOS Computational Biology* (2024). doi:10.1371/journal.pcbi.1012032.
- [3] Li SR, Moheimani H, Herzig B, Kail M, Krishnamoorthi N, Wu J, Abdelhamid S, Scioscia J, Sung E, **Rosengart A**, Bonaroti J, Johansson PI, Stensballe J, Neal MD, Das J, Kar U, Sperry J, Billiar TR. High-dimensional proteomics identifies organ injury patterns associated with outcomes in human trauma. *Journal of Trauma and Acute Care Surgery* (2023). doi:10.1097/ta.0000000000003880.
- [4] Abdelhamid SS, Scioscia J, Vodovotz Y, Wu J, **Rosengart A**, Sung E, Rahman S, Voinchet R, Bonaroti J, Li S, Darby JL, Kar U, Neal MD, Sperry J, Das J, Billiar TR. Multi-Omic Admission-Based Prognostic Biomarkers Identified by Machine Learning Algorithms Predict Patient Recovery and 30-Day Survival in Trauma Patients. *Metabolites* (2022). doi:10.3390/metabo12090774.

Class Projects and Other Unpublished Work

- [1] Shen A, Rosengart A. Can ChatGPT Predict the Weather? A Study of the Universality of the Self-Attention Mechanism. Carnegie Mellon 10-716 (Advanced Machine Learning). Spring 2024.
- [2] Rosengart A, Scharfstein K, Thivierge G. Variational and Proximal Causal Effect Estimation in the Presence of Unmeasured Confounding. Carnegie Mellon University 10-708 (Probabilistic Graphical Models). Winter 2023.

TALKS

Invited

- [1] Significant Latent Factor Interaction Discovery & Exploration. CSI Retreat, Center for Systems Immunology, University of Pittsburgh. October 2022.

Formal Presentations

- [1] Wastewater-based Epidemiology. CDC Site Visit, Delphi Research Group, Carnegie Mellon University. June 2023.
- [2] Characterizing Variability in Pepper Mild Mottle Virus for Wastewater Epidemiology. CDC Center for Forecasting and Analytics Site Visit, Delphi Research Group, Carnegie Mellon University. June 2024.

AWARDS

- Phi Beta Kappa. 2021.

WORK EXPERIENCE

2023 Summer

Laboratory Researcher

University of Pittsburgh, Center for Systems Immunology

Applied large language modeling tools to prediction tasks in protein binding. Led code composition and documentation for construction, training, and use of protein binding Transformer.

2022 Spring-Summer

Laboratory Researcher

University of Pittsburgh, Center for Systems Immunology

Developed methodology for identification of putative mechanisms of disease from multi-omic biomedical data. Composed and documented code.

2021 Summer

Trauma ATLAS Intern

Pittsburgh Trauma Research Center, University of Pittsburgh

Analyzed scale, multi-omics data from experimental trials in trauma patient treatments. Composed sample code scripts for data cleaning, visualization, and gene set enrichment analysis to fellow researchers. Implemented consensus-based feature selection for dimension reduction.

2021 Spring

Undergraduate Research Trainee

Department of Statistics, University of Michigan

Trained in performing time series analysis and statistical modeling of stochastic processes for epidemiological study using partially observed Markov processes.

2021 Fall

Laboratory Researcher

Center for Biologic Imaging, University of Pittsburgh

Worked on the development of efficient and accurate computational methodologies for analysis of large-scale experimental biomedical imaging data. Adapted preexisting software and the development of a software pipeline for cleaning, manipulating, and mapping terabyte-scale mouse brain imaging slices to the Allen Mouse Brain Atlas.

Technical Skills Programming: R, Python, L^AT_EX, Shell