

Aaron Colon

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Summary

Undergraduate researcher in Biostatistics & Health Data Science with experience in computational epidemiology, bioinformatics, and DNA computation. I am interested in data-driven public health forecasting and outbreak simulation.

Education

Lehigh University, Bethlehem, PA	Graduation: May 2027
Biostatistics & Health Data Science, Pre-medicine GPA: 3.77	
Easton High School, Easton, PA	Graduation: June 2023 Honors:
AP Scholar with Honors, High Honor Roll, Top 1% Graduating Class	

Technical Skills:

- **Programming & Data Science:** Python, R, GitHub
- **Modeling & Analytics:** Epidemiological modeling (SIR), Simulation design, Feature engineering
- **Bioinformatics Tools:** HHPred, Phamerator, DNA Master, PhagesDB, GeneMarkS, DeepTMHMM, SOSUI
- **Data Visualization:** Matplotlib, ggplot2

Research Involvement

Forecasting Disease Outbreaks using Temporal Networks in Congregate Settings	2024-Current
<ul style="list-style-type: none">• Hypothesis: Epidemic outbreaks in congregate settings can be predicted and mitigated through network-based modeling of temporal interactions.• Results: Built temporal contact network models using real-world conference data; simulations showed that targeted interventions at high-degree or high-betweenness nodes significantly reduced outbreak magnitude.• Skills learned and applied: Network analysis (Python: pandas, networkx, matplotlib), SIR simulation modeling, data visualization, public health informatics, and mathematical modeling of infection dynamics.	
DNA Sticker Computation via Electrowetting	2023-Current
<ul style="list-style-type: none">• Hypothesis: DNA strands can be used as programmable “stickers” in a microfluidic environment to perform combinatorial computations through electrowetting-on-dielectric (EWOD) control.• Results: Demonstrated droplet-based manipulation of DNA “sticker” sequences on EWOD chips, validating the feasibility of bio-inspired logical operations and contributing to an upcoming manuscript.• Skills learned and applied: EWOD operation, DNA computation principles, lab instrumentation, and research writing	
Sea-Phages/Sea-Genes	2023-2024
<ul style="list-style-type: none">• Hypothesis: The genomic structure and coding potential of Mycobacterium smegmatis phage <i>Hanako</i> could reveal unique gene functions contributing to host-phage interactions.• Results: Successfully annotated and submitted <i>Hanako</i>’s genome to NCBI GenBank, identifying hypothetical proteins and refining gene start/stop predictions through comparative analysis using Starterator, GeneMarkS, and BLASTp.• Skills learned and applied: Genome annotation, bioinformatics tools (DNA Master, PhagesDB, HHPred, DeepTMHMM, SOSUI, Phamerator), comparative genomics, GenBank formatting and submission.	

Presentations:

- **Aaron Colon**, Cheyenne Desmond, Thomas McAndrew, Forecasting Disease Outbreaks using Temporal Networks in Congregate Settings
 - Lehigh University, Bethlehem, PA. *Lehigh College of Health Research Expo*, 2025.
 - Lehigh University, Bethlehem, PA. *Lehigh Summer Research Expo*, 2025.
- **Aaron Colon**. et. al, DNA Sticker Computation via Electrowetting
 - Lehigh University, Bethlehem, PA. *Lehigh Summer Research Expo*, 2024.
 - John Hopkins University, Baltimore, MD. *30th International Conference on DNA Computing and Molecular Programming (DNA30)*, 2024.

Awards

- Dean’s List (4x)
- Best Research Award (DNA Sticker Computation via Electrowetting)
- Tri-Alpha Inductee
- Lehigh Trustee Scholar