# **ZORIAN THORNTON**

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### **EDUCATION**

University of Washington - Seattle, WA

Sep. 2019 - Present

Ph.D Student in Genome Sciences

Virginia Tech - Blacksburg, VA

Aug. 2015 - May 2019

**B.Sc Statistics** 

Virginia Tech - Blacksburg, VA

Aug. 2015 - May 2019

B.Sc Computational Modeling and Data Analytics

Minors in Mathematics and Computer Science

### RESEARCH EXPERIENCE

### Fred Hutchison Cancer Research Center

Seattle, WA

Doctoral Candidate, advised by Frederick Matsen IV

July 2021 - Present

• Developing torchdms: a flexible modeling framework for fitting neural networks with PyTorch to deep mutational scanning experiments. The goal of torchdms is to infer a *fitness landscape* that accurately predicts quantitative phenotypes of unseen protein variants while preserving biophysically interpretable parameters.

### Fred Hutchison Cancer Research Center

Seattle, WA

Doctoral Student, advised by Frederick Matsen IV

June 2020 - July 2021

• Developing and fitting neural network models to predict protein variant phenotypes from deep mutational scanning experiments to accurately predict fitness of unseen variants and infer the shape of global epistasis.

### University of Washington Department of Genome Sciences

Seattle.WA

Lab Rotation with Brian Beliveau

March 2020 - June 2020

• Implemented a bioinformatic pipeline for the design of split-oligo probes for fluorescence in situ hybridization (FISH) experiments to enable fast and affordable design of highly-specific RNA FISH probes.

## Fred Hutchison Cancer Research Center

Seattle, WA

Lab Rotation with Frederick Matsen IV

Jan. 2020 - March 2020

• See related entry above

### University of Washington Department of Genome Sciences

Seattle, WA

Lab Rotation with William S. Noble

Sept. 2019 - Dec. 2019

• Implemented a novel method for systematically finding potential functional inter-chromosomal contacts in Hi-C data.

### Virginia Tech Department of Statistics

Blacksburg, VA

Undergraduate Research with Allison Tegge

Aug. 2018 - May 2019

 Implemented Self-Organizing Maps and conducted survival analysis to identify genes and pathways associated with progression of colorectal cancer from patients included in The Cancer Genome Atlas Program.

# Virginia Tech Department of Statistics

Undergraduate Research with Leah Johnson

Blacksburg, VA Aug. 2017 - May 2019

• Developed a new formula for the disease basic reproductive number,  $R_0$ , to include temperature sensitive midge life history traits to predict potential regions for the spread of Bluetongue viral disease.

# University of Washington Department of Genome Sciences

Seattle, WA

Undergraduate Research with Jim Bruce

June. 2017 - Aug. 2017

• Visualized inter-surface regions of cross-linked proteins and built probabilistic models to quantify competition between proteins for linkage with other proteins to help scientists characterize protein function, discover mutations, and discover protein-protein interactions to tackle molecular challenges such as cancer

#### **PUBLICATIONS**

- Yu, T., **Thornton, Z.**, Hannon, W., DeWitt, W.S., Radford, C., Matsen IV, F.A. and Bloom, J.D. (2022). A biophyscial model of viral escape from polyclonal antibodies *Virus Evolution*, 8(2), veac110.
- El Moustaid, F., **Thornton, Z.**, Slamani, H., Ryan, S. and Johnson, LR (2021). Predicting temperature-dependent transmission suitability of bluetongue virus in livestock. *Parasites Vectors*, 14(1), pp.1-14..
- Keller, A., Chavez, J.D., Eng, J.K., **Thornton, Z.** and Bruce, J.E. (2018). Tools for 3D Interactome Visualization. *Journal of proteome research*, 18(2), pp.753-758.

#### RESEARCH PRESENTATIONS

- Zorian Thornton, Predicting Disease Progression of Colorectal Cancer via Self-Organizing Maps, RECOMB 2019, George Washington University, Washington DC, May 2019
- Zorian Thornton, Modeling Bluetongue Virus via Markov Chain Monte Carlo Methods, Student Experiential Learning Conference, Virginia Tech, Blacksburg, VA, Apr. 2018
- Zorian Thornton, Viewing Molecular Interaction Interfaces Through Directed Computational Methods, Department of Genome Sciences Research Symposium, University of Washington, Seattle, WA, Aug. 2017

### WORK EXPERIENCE

### Adaptive Biotechnologies

Seattle, WA

Machine Learning Computational Biology Intern

June 2022 - Sept. 2022

• Conducted benchmarking of semi-supervised learning methods for predicting T-cell receptor specificity from high-throughput sequencing data.

# Virginia Tech Statistical Applications and Innovations Group

Blacksburg, VA

Associate Collaborator

June 2019 - Sept. 2019

• Assistant statistics consultant to Virginia Tech graduate students and faculty.

# Virginia Tech Department of Statistics

Blacksburg, VA

Course Development Contributor for Dr. Christian Lucero

May. 2019 - June 2019

• Assisted in writing of lectures, and homework assignments for an introductory machine learning class: Introduction to Data Analysis and Visualization (CMDA 3654)

### Virginia Tech Department of Statistics

Grader for Dr. Jane Robertson Evia

Blacksburg, VA Aug. 2018 - May 2019

• Statistics for Social Science (STAT 3604)

Nielsen

Chicago, IL

Data Science Intern

June 2018 - Aug. 2018

• Implemented statistical framework to identify possible errors in scanned receipt data and implemented pipeline to attempt to correct errors

### Virginia Tech Math Emporium

Blacksburg, VA

Instructional Assistant

Jan. 2017 - Aug. 2018

• Teaching assistant for introductory math courses including differential and integral calculus, graph theory, differential equations, and linear algebra

### **HONORS**

# National Science Foundation Graduate Research Fellowship Program

Awarded 2020

• Honorable mention

### Mu Sigma Rho, National Honor Society

Inducted April 2019

• National honorary society for statistics whose purpose is the promotion and encouragement of scholarly activity in statistics, and the recognition of outstanding achievement among the students and instructional staff in eligible academic institutions.

### College of Science Dean's Roundtable Scholarship, Virginia Tech

Aug. 2018 - May 2019

 Merit-based scholarship awarded annually to four students who are and will continue to be excellent representatives of the College of Science

# Luther and Alice Hamlett Undergraduate Research Grant

Aug. 2018 - May 2019

• Selective research grant awarded to Virginia Tech undergraduates conducting research in the Academy of Integrated Sciences

#### Fralin Undergraduate Research Fellowship

Aug. 2017 - May 2018

• Selective fellowship awarded to Virginia Tech undergraduates conducting research involving the life sciences

# Eckert Statistics Scholar, Virginia Tech

Jan. 2016 - May 2019

• Three-year scholarship awarded annually to two Statistics majors for academic excellence

### PROGRAMMING LANGUAGES, TOOLS, AND CONCEPTS

**Preferred Programming Languages** 

Python, R, C/C++

**HPC Tools** 

CUDA, MPI, openMP, SLURM

Dev Tools & Environments

Git, VSCode, Jupyter

Other Programming Languages

Java, MATLAB, SQL

### PROFESSIONAL ORGANIZATIONS

- The American Association of Immunologists member since 2020
- National Society of Blacks in Computing member since 2017
- American Statistical Association member since 2016
- National Society of Black Engineers member since 2016