

# Zach Eliason

*Bioinformatician with strong computational skills and expertise in genomics and cancer immunology research. Enjoys deep learning approaches and coding solutions to real-world problems.*

## EDUCATION

**BRIGHAM YOUNG UNIVERSITY, COLLEGE OF LIFE SCIENCES** | Provo, UT

**B.S. Bioinformatics, Minor in Computer Science** Dec. 2023

GPA: 3.99/4.0, Major GPA: 4.0/4.0

- + 2022 Amgen Scholar, Washington University in St. Louis
- + Recipient of 4 year full-tuition university scholarship
- + College of Life Sciences Dean's List (top 5%)

## BIOINFORMATICS WORK EXPERIENCE

**BIOINFORMATICS INTERN** | *May 2023—Aug. 2023*

**Zayt Bioscience** Berlin, Germany

- + Developed and assessed 100+ novel synthetic promoters using state-of-the-art deep learning to enhance protein production in gene expression experiments
- + Built custom BLAST-guided DNA alignment pipeline to facilitate comparative analysis of target genes across over 100 fungal species

**BIOINFORMATICS RESEARCH ASSISTANT** | *Jun. 2022—Aug. 2022*

**Zhang Lab, Washington University** St. Louis, MO.

- + Led team of 3 on HLA genotyping research project for improving cancer outcomes
- + Designed and implemented novel genotyping tool achieving over 90% accuracy
- + Conducted literature review in order to compile paper for publication

**DATA MANAGER & RESEARCH ASSISTANT** | *Jan. 2021—Apr. 2022*

**Abbott Lab, BYU** Provo, UT.

- + Analyzed hydrology time series spanning 3+ years to advance conservation efforts
- + Oversaw and trained 10+ computational assistants to maintain database integrity
- + Orchestrated bi-monthly lab meetings to ensure communication and organization
- + Designed data cleaning and correction tools for large data sets in use by 12+

**BIOINFORMATICS RESEARCH ASSISTANT** | *Apr. 2021—Apr. 2022*

**Piccolo Lab, BYU** Provo, UT.

- + Led team of 10 in study of publication trends to advance pedagogical methods
- + Created data visualizations, ran analyses, and oversaw manuscript for publication
- + Designed, built, and tested web tool for teaching and evaluating bioinformatics students used by 4 professors and 100+ students

## CONTACT

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[eliasonzach@gmail.com](mailto:eliasonzach@gmail.com)

[github.com/zacheliason](https://github.com/zacheliason)

[linkedin.com/in/zacheliason](https://linkedin.com/in/zacheliason)

## SKILLS

### PROGRAMMING

- + **Languages:** Python, R, C++, SQL & NOSQL databases, Bash (Unix Shell), web programming
- + **Version Control:** Git

### DEEP LEARNING

- + **Frameworks:** PyTorch
- + **Architectures:** CNN, RNN, VAE, GAN, transformer

### DOMAIN KNOWLEDGE

- + Genetics, genomics, biology, statistics
- + BLAST, de novo and reference genome alignment

### OTHER

- + Adobe Creative Suite, proficiency in Russian language, initiative, teamwork

*Last updated 07-22-2023.*

*View this resume online at*

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