# Gavin Coladonato

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#### **EDUCATION**

### University of California, Santa Cruz

Sep. 2021 – Jun. 2025

Bachelor's of Science in Biomolecular Engineering and Bioinformatics

- Relevant coursework: Applied Machine Learning, Research Programming in Life Sciences, Bioinformatics Models
  and Algorithms, Computational Genomics, Data Structures and Algorithms, Python Programming, Probability and
  Statistical Inference, Database Management Systems, Genetics and Genomics, Cell and Molecular Biology.
- Grade Point Average: 3.6

#### EXPERIENCE

#### Undergraduate Research Intern

Jun. 2024 – Aug. 2024

Stanford Medicine, Radiological Sciences Laboratory

Stanford, CA

- Designed and implemented a novel machine learning approach for reconstructing high-resolution neural shape models from Diffusion Tensor MRI scans, enhancing anatomical fidelity.
- Built an end-to-end preprocessing and modeling pipeline in PyTorch and NumPy, including segmentation and normalization.
- Worked independently in collaboration with a Stanford Medicine postdoctoral researcher to deliver reproducible, production-ready code.

## Student Outreach Program Coordinator

Sep. 2022 – Jun. 2023

UCSC Engaging Education

Santa Cruz, CA

- Coordinated outreach to Bay Area minority students by organizing campus tours, guest speaker events, and informational sessions highlighting college opportunities.
- Led guided campus tours and presented university resources and support services to prospective students and families.
- Managed event logistics and conducted direct phone outreach to engage participants and answer questions.

### RESEARCH

## Deep Learning Research Assistant

Jun. 2023 - Dec. 2023

Neuromorphic Computing Group

- Created a Python tutorial notebook introducing Spiking Neural Networks (SNNs) and demonstrating their benefits for reducing model storage and energy usage.
- Collaborated with researchers developing open-source tools for SNN model conversion and optimization in PyTorch.

#### Projects

#### Computational Analysis of Polycythemia Vera Driver Mutation

Mar. 2024 - Mar. 2025

 $Independent\ project\ and\ course\ research$ 

- Developed a computational pipeline to confirm JAK2 V617F as the primary driver mutation in Polycythemia Vera, integrating gene expression analysis, pathway enrichment, and machine learning classification.
- Conducted differential gene expression and network modeling using publicly available genomic datasets (GEO, TCGA, COSMIC).

#### LEADERSHIP & SERVICE

# Eagle Scout (Highest Rank)

Awarded Nov. 2020

Scouts BSA

Mountain View, CA

- Designed and led a COVID-19 relief project supporting the Stanford Medicine Van, a mobile medical unit providing free healthcare to uninsured students across the SF Bay Area.
- Coordinated a food and school-supply drive in the local community, organizing over 100 volunteer hours.
- Collected and distributed 100 backpacks with school supplies, 100 meal bags, 120 blankets, and 50+ gift cards for families in need.

#### SKILLS

- Python (5+ yrs), PyTorch, NumPy, Pandas, scikit-learn
- Machine Learning, Bioinformatics Analysis
- Bash (2+ yrs), Git, Jupyter Notebooks, Linux
- Strong written and verbal communication