Shawn Schulz

shawnschulz@gmail.com • Cell: (510)-334-8000 • https://github.com/shawnschulz

EDUCATION

University of California, Berkeley, Berkeley, CA

2024—

Masters of Science, Molecular Science and Software Engineering

University of California, Berkeley, Berkeley, CA

2018 - 2022

Bachelor of Arts, Major in Molecular and Cell Biology

GPA: 3.9

WORK EXPERIENCE

Data Engineer, Stanford Medicine, Kuo and Gawad Labs

June 2022-Nov. 2023

- Developed enterprise user focused HPC pipeline in **python**, **R**, **C++**, obtaining \$1**M**+
- Improved machine learning analysis reducing errors by 500% using a gaussian process model.
- Developed and maintained a database serving **800TB** of genomic data, saving over **400 days** of user runtime via improved concurrency with **bash**, and **linux server environment shell**.
- Automated building and testing, improved error handling and compatibility serving 30+ enterprise users using R, docker, nextflow, github actions, git cli.
- Performed high impact data analysis on large raw datasets for \$100,000+ grant using PCA and k-means clustering, preventing overfitting and improving performance via numpy and pytorch.
- Solved data science challenges in cancer detection and presented to audiences of 50+ peers.

Machine Learning Researcher, Berkeley Artificial Intelligence Research (BAIR) Nov. 2021—May 2022

- Developed, tested, evaluated and benchmarked SCVI **variational autoencoder deep learning** model serving thousands of users. Solved sophisticated machine learning problems.
- Produced high impact results and figures for deep learning models in a \$100,000+ grant.
- Improved model performance by 200% using sci-kit learn, numpy and pytorch libraries to improve processing throughput and streamline batch correction.
- Modeled using **PCA** and visualized using **matplotlib** to find SARS-CoV2 infection pathology.
- Used gaussian mixture model infection level classifier to find novel mechanisms of COVID-19

PROJECTS

gpt-flow

- Created AI workflow visualization tool for text **LLM** data inputs and outputs using **javascript/typescript**, **nodejs**, and **React** framework.
- Developed web server backend serving 100 users using flask.

transcriptome-transformer

• Developing multi-ome classifier and protein solver using C++ and CUDA.

Robotic liquid handler

• Embedded systems programming of robotic pipette attachment for 3d printers using C++.

OTHER EXPERIENCE

Undergraduate Researcher, UCSF Department of Surgery, Wang Lab

Nov. 2020-Aug. 2021

• Used a computer vision package in **Matlab** to analyze immunofluorescence microscopy data to find diameter size of blood vessels and flow rate.

Ashland Free Medical Clinic, Clinic Coordinator

Dec. 2020—

• Managed a team of **nine** volunteers and managed web EHR for over **200 patients**.

PUBLICATIONS

- NF-κB inhibitor alpha has a cross-variant role during SARS-CoV-2 infection in ACE2-overexpressing human airway organoid
- Cellular and molecular characterization of peripheral glia in the lung and other organs

SKILLS and COURSEWORK

- C++, C#, Python, Javascript/Typescript, R, Rust
- git, AWS, OpenAI API, react, SQL, Excel, sci-kit learn, jupyter, conda, flask, html, css
- Biotech Software, Machine Learning, High Performance Computing, Full Stack Development