Joshua P. Schaaf

610-550-6432 • Davis, CA • joshuaschaaf@gmail.com • https://github.com/viabard

SKILLS

Python | R | Git | SQL | C | JavaScript | Linux | Excel | PowerPoint | Pandas | NumPy | scikit-learn | TensorFlow Keras | PyTorch | Matplotlib | PCR | Methylation | FASTA/Q | ChIP-Seq | Github Actions | CI/CD | AWS

EXPERIENCE

Bioinformatics Programmer Analyst II, HJF WHIRC GYN-COE, Remote *January 2022-Present*

- Analyze, design, develop, modify, test, and implement software tools and scripts supporting bioinformatic analysis of multi-omic data including clinical, proteomic, and genomic
- Recommend enhancements or changes to current system configuration to maintain optimum system performance and utility
- Create detailed test cases and document all programming and system changes

Data Analyst, FOXO Technologies, Remote *July 2021-November 2022*

- Analyzed 'wide' epigenetic data (~1 million features) using Python (pandas, numpy) and R to provide accurate predictors for clinical targets using automated machine learning (ML) platform DataRobot. Analysis using working knowledge of *scikit-learn* and other ML/data mining algorithms
- Created easy to understand figures/interactive dashboards with data visualization tools in Python (plotly, matplotlib, seaborn), R (ggplot), and lucidchart
- Performed Quality Control and Validation analyses on big data, utilizing Python in AWS Sagemaker Processes, Sagemaker Notebooks, Cloudwatch, storing data on AWS S3
- Developed RAPA (Robust Automated Parsimony Analysis), an open-source Python package for performing recursive feature reduction of ML models created with DataRobot
- Created unit tests and update/maintain documentation for Python packages supporting the company
- Utilized GitHub for software development, version tracking, and data analysis collaboration
- Utilized AWS Sagemaker and Cloudwatch to analyze/create ML models with data stored in S3
- Presented analytical findings to groups of 5-10 people in an understandable and concise manner
- Quickly switched between projects, adapting to different techniques in the start-up environment

Genetic Research Experience, Balciunas Lab, Temple University, PA *December 2018-May 2021*

- Effectively collaborated with a team to engineer a fully 'floxed' *tcf21* allele for the study of post-embryonic regenerative abilities of zebrafish hearts, and of the role *tcf21* plays in heart regeneration
- Created a pipeline for transforming large, raw ChIP-seq fastq datasets with GalaxyHub, bowtie 2, samtools, R and Python scripts, and MEME-ChIP with statistical programming practices
- Experienced with Gel Electrophoresis, PCR, Zebrafish Maintenance/Rearing, Survival Surgeries (*Danio rerio*), Conditional Mutagenesis, Cre-loxP, CRISPR/Cas9, Fluorescent Microscopy
- **Poster**: Conditional Mutagenesis of Zebrafish *tcf21* at 2019 Undergraduate Research Symposium
- Presented at weekly lab meetings with in-depth analysis and discussion of obtained data, as well as peer-reviewed journal presentations
- Accepted to Research Experiences for Undergraduates REU, CSUSM: Cancelled, COVID-19 Pandemic

CODING PROJECTS

Imputation Analysis with High Dimensional Data: Using many techniques described in literature, including HoloClean in PostgreSQL, I analyzed different imputation methods for highly dimensional data.
Using Machine Learning for Species Recognition: Developed Python scripts to retrieve over 300,000 images of PA moths from the iNaturalist API, then a TensorFlow deep learning model for species recognition.
Using Machine Learning for Chest X-Ray Abnormality Detection: Created a machine learning model with Python scripts for CXR preprocessing with scikit-image and object detection with YOLOv5 in PyTorch.

Joshua P. Schaaf

610-550-6432 • Davis, CA • joshuaschaaf@gmail.com • https://github.com/viabard

EDUCATION

Temple University, College of Science and Technology — Philadelphia, PA
Professional Science Masters (PSM), Bioinformatics — *May 2022 - Cumulative GPA: 4.0/4.0*
Bachelor of Science, Biochemistry, Minor in Computer Science — *May 2021 - Cumulative GPA: 3.9/4.0*

Honors and Distinctions: ***Magna Cum Laude***, Distinction in Major, Dean's List, **Temple Honors Program**, **Natan Luehrmann-Cowen Memorial Award** (College-Wide Academic Excellence, Musical Instrument)
Scholarships: **NSF REU** at CSUSM with Dr. Arun Sethuraman (Cancelled due to SARS-CoV-2), **Science Scholars Program** (SSP), Temple Academic Scholarship, TUDMB Scholarship

RELEVANT COURSEWORK

Mathematics/CS:	Coding in C; Biological Models in Python; Program Design/Abstraction (Java); Data Structures (Java); Calculus I, II, Honors III; **Machine Learning**; **Biostatistics**
Physical Sciences:	Honors Organic Chemistry I, II; Physical Chemistry of Biomolecules; Physics I, II
Biological Sciences:	Genomics (R, Python); Genetics; Cell Structure and Function; Biochemistry I, II; Fundamentals of Genomic Evolutionary Medicine; Ethics in Biotechnology

LEADERSHIP AND ACTIVITIES

1st Flute, Video Game Orchestra at UC Davis — *2022*
Volunteer Belayer, Reach Climbing & Fitness — *2022*
Principle 2nd Violinist, Lower Merion Symphony — *2021-2022*
1st Flute/Piccolo Player, Temple University Diamond Marching Band/Collegiate Band — *2017- 2018*