Joshua P. Schaaf
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EDUCATION

Temple University, College of Science and Technology — Philadelphia, PA
Candidate for Accelerated Professional Science Masters (PSM) — *May 2022*
Bioinformatics — *Cumulative GPA: 4.0/4.0*

Bachelor of Science — *May 2021*
Biochemistry, Minor in Computer Science — *Cumulative GPA: 3.88/4.0*

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

EXPERIENCE

Data Analyst, FOXO Technologies, Minneapolis, MN — *July 2021-Present*

- Analyze 'wide' epigenetic data (~1 million features) using Python (pandas, numpy) and R to provide accurate predictors for clinical targets using automated machine learning platform DataRobot. Analysis using working knowledge of *scikit-learn* and other machine learning/data mining algorithms
- Develop RAPA (Robust Automated Parsimony Analysis), an open-source Python package for performing recursive feature reduction of ML models created with DataRobot
- Create unit tests and update/maintain documentation for RAPA
- Utilize GitHub for software development, version tracking, and data analysis collaboration
- Utilize AWS Sagemaker and Cloudwatch to analyze/create ML models with data stored in S3
- Present analytical findings to groups of 5-10 people in a clear, understandable, and concise manner
- Create easy to understand figures/interactive dashboards with data visualization tools in Python (plotly, matplotlib, seaborn), R (ggplot), and lucidchart

Assistant Research Scientist, TheWell Bioscience, North Bruswick, NJ — *May2021-June 2021*

- Created a chatbot with Zoho's Costomer Relationship Management with a workflow meant to help customers with specific questions at the landing page
- Reported on performance metrics for consumer engagement and retention

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

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- 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

Chemical Biology Lab Intern, Wang Group Lab, Temple University, PA *May 2018-December 2018*

- Contributed to the development of a homogeneous Drug-Antibody Conjugate linker
- Gained skills in NMR, Mass Spectrometry, HPLC, Ultra-Violet Spectroscopy, Thin-Layer Chromatography, Column Chromatography, and Solid Phase Peptide Synthesis

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.

LEADERSHIP AND ACTIVITIES

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*