

Annika Jorgensen

Email: aljorge2@asu.edu
Phone Number: 385-315-4242
Website: annikajorgensen.com

EDUCATION

- | | |
|---|------------------------|
| Arizona State University
MS Biology: Computational Life Sciences, GPA: 3.97 | Tempe, AZ
2023–2024 |
| Arizona State University
Magna Cum Laude Honors BS Biophysics, GPA: 3.60 | Tempe, AZ
2019–2023 |

TECHNICAL SKILLS

- **R and RStudio:** Proficient in statistical programming, specializing in bioinformatics and biostatistics
- **Python and MATLAB:** Skilled in use of IDE and writing code for scientific computing
- **Bash:** Strong command of shell scripts for system automation and administration with SAM/BAM files
- **Git:** Extensive experience in version control and collaborative software development workflows
- **SQL:** Familiar with database management, query optimization, and data manipulation
- **Biostatistics:** Adept at utilizing statistical tests to analyze and interpret complex biological datasets
- **Quality Assessment:** Trained in ensuring data integrity and reliability through rigorous quality control and preprocessing techniques using FASTQ
- **Data Visualization:** Experienced in creating statistical visualizations that are easily understood by nonscientists
- **Sequence alignment and Variant Calling:** Strong background in analyzing high-throughput sequencing data to identify genetic variations and make biological insights.
- **Molecular Biotechnology:** Familiar with laboratory techniques including NGS, PCR, and genetic engineering

WORK EXPERIENCE

- | | |
|---|---|
| Bioinformatics Programmer–Dr. Xin Sun & Dr. Allen Wang
Xin Sun Lab & Center for Epigenomics | July 2024- Present
San Diego, CA |
| <ul style="list-style-type: none">– Employ software tools to interpret biological data– Analyze RNAseq and ATACseq data– Conduct statistical methods on biological data | <ul style="list-style-type: none">– Debug computer programs– Maintain webages– Documents changes to operational systems |
| Graduate Research Assistant–Dr. Melissa Wilson
Sex Chromosome Lab | November 2021-May 2024
Tempe, AZ |
| <ul style="list-style-type: none">– Statistically analyzed sex-differential expression of genes with RNA-seq data– Implemented biostatistical methods in R including multivariate regression, principal component analysis, and hypothesis testing | <ul style="list-style-type: none">– Developed computational biology curricula for an undergraduate class– Queried the Reactome database and performed hypergeometric overrepresentation analyses |
| Commercial Intern
Civica Rx | Summer 2020-Spring 2021
Lehi, UT |
| <ul style="list-style-type: none">– Designed and Maintained drug sales databases– Queried large databases with Tableau | <ul style="list-style-type: none">– Created government official datasheets in Excel– Drafted sales agreements for hospital partners |