# Claire Hsieh

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#### **EDUCATION**

Bachelor of Science in **Genetics and Genomics** at University of California, Davis

Expected June 2024

Minor: Quantitative Biology and Bioinformatics, Statistics

GPA: 3.736

Relevant coursework:

- Statistical Data Science
- Regression Analysis
- Probability & Stochastic Processes with Applications to Biology
- Linear Algebra
- Mathematical Statistics

- Comparative Genomics
- Functional Genomics
- Genes and Gene Expression
- Human Genetics and Genomics
- Data Structures
- Theory & Practice of Bioinformatics

#### **TECHNICAL SKILLS**

Programming Languages: Python (numpy, pandas, matplotlib), R, Perl, BLAST, MATLAB

Software: YASARA

Operating System / Scripting Language: Windows, Linux, Unix, Bash

Version Control: Git

#### **RESEARCH SKILLS**

- General Laboratory
- Bacteriology
- Gel Electrophoresis
- qPCR

- Grid Preparation (Electron Microscopy)
- Plaque Assay
- Miniprep (DNA extraction)

#### **RESEARCH INTERESTS**

- Statistical Genomics
- Cancer and aging-related research
- Metabolomics
- Machine learning with clinical applications
- Markov Models

#### **A**WARDS

• 2023 College of Biological Sciences Summer Undergraduate Research Award for \$7000 (declined)

#### RESEARCH EXPERIENCE

## Research Assistant | Topological Molecular Biology Lab | Apr 2022 - Present

- Analyzed viral datasets to extract and annotate proteins using sequence and structural similarity
- Utilized profile HMMs in predicting SARS COV-2 mutation probabilities weighted by country
- Cleaned and aligned genomic datasets using MAFFT and Clustal Omega
- Grew and collected microbial cultures
- Set up and performed gel electrophoresis
- Prepared frozen samples using GP2 and imaged using Cryo-EM
- Designed primers and performed qPCR to quantify DNA
- Extracted DNA using miniprep or chloroform

## Research Assistant | Korf Lab | Dec 2022 - Present

- Developed Python programs that use intron position to predict gene enhancement
- Examined orthologous genes to identify introns with potential for enhancing genes
- Created algorithms to identify motifs using discretized position weight matrices, regular expressions, and k-mers
- Implemented a Stochastic Viterbi and forward-backwards algorithm in Python to generate random, probable isoforms

# Irvine Summer Institute in Biostatistics and Undergraduate Data Science | UC Irvine | Jul 2023 – Aug 2023

- Learned fundamentals of probability, Monte Carlo methods, linear models, generalized linear models, causal inference, and Bayesian statistics
- Used R and machine learning to predict presence of colorectal cancer using metabolomics

#### **WORK EXPERIENCE**

#### Accounting Assistant | Quicklinks Advisor 365 | Aug 2020 - Dec 2022

- Create and update financial statements
- Reconcile transactions and identify discrepancies

## **C**LUBS

## Davis Data Science Club | Social Media Officer | Sep 2022 - Present

• Designed and distributed posters to promote club meetings and events

### **COMMUNITY SERVICE**

#### Priceless Pets | Jan. 2019 - Feb. 2020

- Walked dogs
- Trained new volunteers
- Cleaned kennels
- Fed and medicated animals

#### **LANGUAGES**

**English**: Native Language

Mandarin: Intermediate Listener, Intermediate Speaker, Beginner reading and writing

#### **PROFESSIONAL TRAINING**

## US Cyber Challenge | Jul 2022

• Coursework: Memory Forensics, Network Forensics, Linux

## Scientific Computing with Python | freeCodeCamp | Mar 2022

• <a href="https://www.freecodecamp.org/certification/Claire Hsieh/scientific-computing-with-python-v7">https://www.freecodecamp.org/certification/Claire Hsieh/scientific-computing-with-python-v7</a>

# Data Analysis with Python | freeCodeCamp | Jan 2023

• https://www.freecodecamp.org/certification/Claire Hsieh/data-analysis-with-python-v7

## **O**THER

Hobbies: Reading, Playing guitar, Running