JIACHEN AI

jiachenai@g.ucla.edu | +1 424-293-9716

EDUCATION

Sept. 2023 – Jun. 2025 University of California, Los Angeles, Fielding School of Public Health, USA

GPA: 4.00/4.00 (four quarters) **Major:** Biostatistics (Master of Public Health)

Steve Wallace Policy Fellowship

APEx Funding at Department of Human Genetics May 2024

Twice UCLA Fielding School of Public Health Dean's Leadership Grant

Apr. 2024 & May 2023

Sept. 2018 – June 2022

June 2022

Jun. 2024

Skills:

Biostatistics Methods, Statistical Learning, Regression Analysis, Probability Theo and Statistics, Linear Algebra

R, Python (NumPy, Pandas, Matplotlib), SAS, STATA, SQL, LaTeX

Linux scripting, Git/GitHub

Container technology: Docker

Data wrangling (Tidyverse), Data visualization (ggplot2)

Databases: OLAP (DuckDB), Google BigQuery

Frontend development (shiny, web app)

Cloud computing (GCP, AWS, Azure, OCI)

High-performance computing (HPC) on cluster (Hoffman2)

Deep learning with Keras + TensorFlow/PyTorch

Southwest University (SWU), School of Life Sciences, China

Outstanding Graduate, Outstanding Intern, Outstanding Undergraduate Graduation Thesis

Major: Biological Sciences (Bachelor of Science) **GPA:** 3.84/4.00 (2nd/65)

International or Overseas Research Award

Dec. 2021 Twice Third-Class Scholarships for Academic Excellence Dec. 2021 & Dec. 2020

Second-Class Scholarship for Academic Excellence, Merit Student Award

Dec. 2019

Skills:

Bioinformatics, Biochemistry, Cell Biology, Molecular Biology, Genetics, Microbiology, Physiology, Neurology

PCR, qPCR, DNA/RNA extraction and quantification, CRISPR/Cas9, cloning, electrophoresis, next-generation sequencing library preparation

RNA-seq, ChIP-seq, single-cell RNA-seq, Bisulfite sequencing, GWAS (Genome-Wide Association Studies)

Western blotting, ELISA, mass spectrometry, protein-DNA interaction assays, chromatin immunoprecipitation (ChIP)

Cell culture, transfection, microscopy (confocal, fluorescence), immunofluorescence, flow cytometry

RESEARCH

Clustering on Mice DNA Methylation Aging Pattern

Los Angeles, USA

Graduate Student Researcher (supervised by Prof. Feng Gao)

Sept. 2024 – Current

K-means based unsupervised hierarchical clustering of methylation levels, investigating the characteristics of methylation profile in mice aging stages — young, midlife, and late-life.

Longitudinal Study on Flu Vaccination Impact: DNA Methylation Influence the Regulation of **RIG-I Signaling Within UGA6 Cohort**

Los Angeles, USA

Volunteer Graduate Student Researcher (supervised by Prof. Matteo Pellegrin)

Mar. 2024 – Current

Multivariate linear regression with LOOCV to establish prediction models for associations between DNA methylation CpG sites and cellular signaling pathway with sero-protection after influenza injection at day0 and day28, with considering epigenetic confounders, including age, gender, BMI, and cell types.

Statistical Data Analysis for Biology and Biomedicine Research

Undergraduate Research Assistant

Chongqing, China Jun. 2021 – Jan. 2023

By multivariate regression in R and found significant correlation between rice yield and irrigation amount in 5 growing stages.

- Via R, performed logistic regression with backward selection and found 3 categorical variables that correlated with semen quality.
- Through deduction and proofs with Markov chains, multivariable calculus and matrices, discussed the recurrence and transience of random walks in three dimensions.

Lymphocyte Regeneration after Thymocyte Damage in Zebrafish

Chongqing, China

Volunteer (supervised by Prof. Li Li)

Oct. 2019 – June 2020

- Discovered that the Cxcl12b/Cxcr4b chemokine signaling pathway plays a key role in the repair of zebrafish T-lymphocytes by real-time PCR (RT-qPCR) and then demonstrated the finding using a receptor antagonist, WZ811.
- Detected apoptosis and a significant reduction of thymus lymphocytes through whole mount in situ hybridization (WISH) and confocal microscopy, in zebrafish treated with 2 mM Metronidazole for 12 hours in vivo.

PUBLICATION

Ai, J. (2021) Lifestyle does affect semen quality: factor determination in a statistical way. Lecture Notes in Medicine and Life Sciences, 4, 163–171. https://www.clausiuspress.com/conference/article/artId/7114.html.

TEACHING

Graduate Teaching Assistant at UCLA

MCD BIO 187AL: Research Immersion Laboratory in Genomic Biology

Los Angeles, USA

June 2024 – July 2024

• Lead lab session and TA session to demonstrate the usage of bioinformatics tools, including NCBI Blast, InterProScan, JPRED4, PSIPRED, DeepTMHMM, WoLF PSORT, SignalP, AlphaFold3 server, ChimeraX

EXPERIENCE

Computational Genomics Summer Institute (UCLA)

Warren Alpert Computational Biology and AI Network Fellow

Los Angeles, USA July 2024 - Current

- Attended workshops in Biobanks, machine learning in biomedical research.
- Cooperating with Warren Alpert faculty in analysing epigenetics data.

Higher Education in Public Health (UCLA)

Los Angeles, USA

Panel Member

Apr. 2024

Presented the FSPH master's program of biostatistics for incoming undergraduate students and answered questions.

SWU Life Science Administrative Office

Chongqing, China

Office Assistant (6 semesters)

Oct. 2018 – Jun. 2021

- Held 5 meetings per semester. Liaised between administrative teachers and the whole class (65 schoolmates), including giving feedback to inquiries from students.
- Provided guidance to approximately 40 freshmen regarding their academic performance, elective courses, clubs, research mentors, etc.

SWU Central Library

Chongqing, China

Volunteer (40 hours)

Oct. 2019 – Jan. 2020

• Organized a charity evening for over 200 people sponsored by the SWU Library Volunteer Association, including being responsible for venue reservation and decoration.

SWU Social Study

Chongqing, China

Team Leader in a project—Ecological Civilization Construction in Jinyun Mountain

Nov. 2019 – Dec. 2019

• Led a team of 5 whose influence reached the top 10/150 with over 1000 votes from strangers to do a field study, interviewed 3 pundits and gathered 226 effective questionnaires about ecosystem preservation and tourism development.

INTERNSHIP

Western Biomedical Technology Inc.

Chongqing, China

Intern, Immunization Team of IT Department

Mar. 2021 – June 2021

- Applied Python and convolution method, developed a light-band recognition program, and improved the precision in western-blot by 20%.
- Devised a program of calculating standard concentration curves in ELISA utilizing R, increasing the calculation speed by 60%.
- Presentation every two weeks. Responsible for experiments and reports in Immunology, Histopathology, and Molecular Biology.
- Promoted the work efficiency of the whole team by virtue of liaising with customers and colleagues in different departments.