## Xiangyu (Leo) Shi

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#### **EDUCATION**

Carnegie Mellon University, Silicon Valley Master of Science, Software Engineering

September 2025 - December 2026

## University of California San Diego **Bachelor of Science, Mathematics - Computer Science**

September 2021 - March 2025

• Overall GPA: 3.9, Provost Honors

#### RESEARCH EXPERIENCE

## Saier Lab, UC San Diego

July 2024 - Present

## Research Assistant

- Designed and implemented a data processing pipeline with a full command-line interface to analyze protein family architectures in the Transporter Classification Database (>23000 protein systems).
- Devised algorithms to parse CDD and Pfam output, detect gaps between conserved domains via adjacent domain pair analysis, and identify optimal domain pairs for classification.
- Created family-level visualization tools to generate domain architecture plots, highlighting both general and characteristic domains, as well as architecture summary plots for comparative analysis.
- Co-author on manuscript in preparation describing a novel transporter classification algorithm and computational tool.

# J Craig. Venter Institute

**June 2025 – August 2025** 

### **Viroinformatics Intern**

- Developed deep learning models to predict antibody titer values and binding properties of COVID-19 variant proteins, supporting rational vaccine design.
- Automated large-scale data transformation and processing pipelines with **Python** and **Bash**.
- Deployed ML workflows on **Docker + AWS EC2**, ensuring scalability for collaborative research use.
- Presented research findings in an institute-wide seminar at JCVI, demonstrating ability to communicate complex computational biology results to interdisciplinary audiences.

# **Ehlers Lab, The Scripps Research Institute**

September 2024 – April 2025

#### **Research Assistant**

- Optimized bi-clustering algorithms for SNP epistasis analysis (GWES), reducing average runtime from 16 hours to 10 minutes, enabling large-scale genetic studies previously infeasible.
- Performed performance profiling and algorithmic tuning to streamline genetic data analysis pipelines, improving efficiency and reproducibility in genomic research workflows.
- Co-author (second author) on manuscript in preparation on GWES data analysis, contributing to algorithm design, implementation, and experimental validation.

#### TECHNICAL EXPERIENCE

# **Yunming Technology**

**January 2025 – June 2025** 

## **Software Engineer**

- Engineered an AI-powered Retrieval-Augmented Generation (RAG) chatbot system for enterprise CRM research applications.
- Implemented query reconstruction, live search, knowledge base filtering, and Server-Sent Events (SSE) integration for scalable deployment.