Willow Ross Carretero Chavez



Education

Work

Experience

- Massachusetts Institute of Technology Expected Graduation May 2024 Cambridge, MA
 - Cumulative GPA 4.6
 - Candidate for BS in Computer Science and Molecular Biology with Minor in Chemistry
 - Relevant Coursework: Fundamentals of Programming, Organic Chemistry I, Math for Computer Science, Multivariable Calculus, Intro to Biological Chemistry, Genetics
- Mater Dei Catholic High School Graduated: May 2019 GPA: 4.63 Chula Vista, CA
- Wayfair LLC Boston, MA

Software Engineering Co-Op: January 2022 — August 2022

- Learned diverse enterprise software tools such as Docker, Kafka, Google BigQuery, Kubernetes, SQL, Git, Pytest, Buildkite, DataDog
- Participated in weekly book clubs, learning functional programming concepts in Haskell as well as a variety of software architectural patterns in Python
- Created multiple microservice APIs using Python, FastAPI, and PostgreSQL
- Rapidly delivered results by merging 30 PRs within the first month, and later 14 within a particularly productive week
- D.E. Shaw Research New York, NY

Early College Intern: May 2021 – September 2021

- Executed free energy perturbation (FEP) molecular dynamics simulations of ligand-protein complex systems
- Implemented several methods of charge neutralization for charge-changing FEP simulations
- Created novel method of FEP network generation using integer linear programming
- Sinskey Lab @ Massachusetts Institute of Technology Cambridge, MA

UROP Student Intern: February 2021-May 2021

- Performed cell counts and metabolite analysis of samples from Erbi and miniBio bioreactors
- Carried out nanodrop, ddPCR, qPCR, BCA, and ELISA assays for nucleic acid and protein quantification
- Mathnasium of Mission Gorge Santee, CA

Instructor: August 2019-April 2021

- Taught students (2nd-12th grade) topics ranging from addition to single-variable calculus
- Effectively leveraged online teaching platforms to perform engaging remote instruction
- Kufareva Lab @ UCSD Skaggs School of Pharmacy and Pharmaceutical Sciences La Jolla, CA Staff Research Associate: October 2019–December 2020

Volunteer: July 2018-August 2018, February 2019-October 2019

- Analyzed protein structure using the MolSoft ICM protein modeling suite, implementing a new method of protein binding pocket similarity scoring in ICMScript
- Authored toolkit to validate, process, and visualize Boolean models of cell signaling networks using Python and R; manuscript currently in preparation
- Performed analysis and visualization of TMT MS² phosphoproteomic data using R
- Executed mammalian tissue culture, PCR, and Gibson cloning protocols

Programming Languages — Python (NumPy), R, ICMScript, Haskell, C

- Familiar with *nix and bash environments (Arch, Ubuntu/Debian)
- Bilingual Native speaker of English and Spanish, conversational French knowledge
- Hobbies: Hiking, backpacking, biking whatever gets me out into nature!

Interests and Skills