

Joel G. Yancey

j.y@ucla.edu

(916) 524-1990

<https://joelyancey.io>

CORE  
COMPETENCIES

**scientific programming, system administration, Linux, shell scripting, OOP, Qt (GUI), UX, C++, Python, R, MATLAB, git, CI/CD, ML, HPC, computational neurobiology, bioinformatics algorithms, RNA-seq, GWAS, HTML, CSS, SQL, Django, JavaScript, TypeScript**

CERTIFICATIONS

**UNIX & Linux System Administration** UCSD Extension

San Diego, CA

11/2020

EDUCATION

**Neuroscience, Bachelor of Science** UCLA

Los Angeles, CA

**Bioinformatics Minor** UCLA School of Engineering and Applied Science

09/2017

WORK EXPERIENCE

**Web Developer**  
**Self-employed**

San Diego, CA

12/2023 – present

- Developed custom websites using JavaScript and Python-based web development frameworks; React, Django, CSS, HTML
- Designed backend web apps employing cloud-based HPC frameworks via AWS Developer Tools

**Scientific Programmer I (Sejnowski Lab)**

San Diego, CA

**Computational Neurobiology Laboratory @ Salk Institute for Biological Studies**

11/2022 – 12/2023

- Developed and maintained AlignEM-SWiFT, a popular open-source, cross-platform Qt/GUI application for EM image registration (or "alignment"). Deployed the application for high-volume scientific use on the high performance computer (HPC) Lonestar6 at the Texas Advanced Computing Center (TACC). It has been cited in published and ongoing research.
- Developed a Python wrapper (API) for a suite of C++ image registration/image processing programs (SWIFT)
- Designed deployment strategies for compiled and interpreted software applications to run with multiprocessing on HPC
- Debugged, tracked issues, and implemented feature requests for a C++ molecular kinetics simulation software (MCell4)

**Jr. Systems Administrator (Sejnowski Lab)**

San Diego, CA

**Computational Neurobiology Laboratory @ Salk Institute for Biological Studies**

10/2019 – 11/2022

- Maintained essential IT infrastructure used in computational neurobiology research, including operating systems, security tools, applications, servers, email systems, laptops, desktops, software, and hardware
- Installed, configured, maintained, upgraded workstations and experiment rigs running Linux or Windows
- Wrote bash scripts, created custom software images, and security tested Debian-based computing machines
- Configured virtual machines for security testing, system monitoring, and performance testing
- Assembled compute workstations to specifications of Salk Institute researchers and staff in the computational laboratories

**Research Assistant I/Lab Manager (Xu & Huang Lab)**

San Diego, CA

**Sanford Burnham Prebys Medical Discovery Institute**

07/2018 – 10/2019

- Developed a novel phagocytosis quantification assay used in published research (Journal of Experimental Medicine).
- Wrote Python scripts and Excel VBA macros for analysis of genetic and other bioinformatics data.
- Analyzed batches of Next-Gen Sequencing (NGS) data, such as RNA-seq data to investigate differential gene expression
- Managerial responsibilities included purchasing, training coworkers in the use of scientific instruments, acquisition and installation of laboratory equipment, primary contact for all oversight compliance, i.e. AAALAC and OSHA

**Laboratory Assistant**

Los Angeles, CA

**Microscopic Tech & Electron Microscopy Core @ UCLA Brain Research Institute**

06/2014 – 09/2014

- Developed film and facilitated access to UCLA's common equipment JEOL 100CX transmission electron microscope.
- Performed histology including cryosectioning, mounting, and staining of brain and other tissues

- Developed a graphical user interface (GUI) in MATLAB for conducting a novel experiment according to protocol
- Used machine learning algorithms (SVM) to decode neural population responses. Developed MATLAB scripts for publication-quality figures i.e. rasters, PSTHs, and neural activity plots.
- Led a 3-year in vivo electrophysiology experiment investigating how the brain encodes spatiotemporal events

## PUBLICATIONS

- MCell4 with BioNetGen: A Monte Carlo Simulator of Rule-Based Reaction-Diffusion Systems with Python Interface** PLOS Computational Biology 04/2024  
Adam Husar, Mariam Ordyan, Guadalupe Garcia, Joel Yancey,..., Thomas Bartol, Terrence Sejnowski.  
Contribution: Conceptualization, Methodology, Software, Validation, Visualization, Writing  
DOI : <https://dx.plos.org/10.1371/journal.pcbi.1011800>
- Multimic landscape and functional analysis of Alzheimer's disease-associated gene variants in human ESC-derived microglia** Journal of Experimental Medicine 12/2020  
Liu T., Zhu B., Liu Y., Zhang X., Yin J., Li X., Hodges A., Zhou L., Yancey J.,...,Huang T., Tanzi R., Xu H.  
Contribution: data curation, formal analysis, software  
DOI : <https://doi.org/10.1084/jem.20200474>
- Role of Rab GTPases in Alzheimer's Disease** ACS Chemical Neuroscience 02/2019  
Zhang X., Huang T., Yancey J., Luo H., and Zhang Y.  
Contribution: writing, editing  
DOI : <https://doi.org/10.1021/acschemneuro.8b00387>

## POSTER PRESENTATIONS & SOFTWARE DEMONSTRATIONS

- AlignEM-SWiFT: Open-source Software for Aligning Electron Micrographs using Signal Whitening Fourier Transforms** San Diego, CA 2022  
Society for Neuroscience 2022  
Source code/PDF: [github.com/mcellteam/swift-ir/tree/joel\\_develop](https://github.com/mcellteam/swift-ir/tree/joel_develop) / [mcell.org/sfn22\\_poster.pdf](https://mcell.org/sfn22_poster.pdf)  
Demos: [youtube.com/playlist?list=PLmDQKF70E5LzYo7MG2nAJcTQjKaXI-Ldy](https://youtube.com/playlist?list=PLmDQKF70E5LzYo7MG2nAJcTQjKaXI-Ldy)
- AlignEM-SWiFT: Graphical Interface for Aligning Electron Micrographs using Signal Whitening Fourier Transforms** Austin, TX 2022  
2022 TACCSTER Symposium @ Texas Advanced Computing Center
- Decoding Stimulus Features From Cortical Population Responses** Los Angeles, CA 2015  
2015 UCLA Neuroscience Undergraduate Poster Fair  
PDF / source: [github.com/joelyancey/SRP199-Poster-UCLA](https://github.com/joelyancey/SRP199-Poster-UCLA)
- Neuroscience Model Builder** C++/Qt tool for diagraming neural circuits. Source code can be downloaded from GitHub and compiled. Tested on Debian GNU/Linux 10 and macOS 14.5 Los Angeles, CA 2014  
Source / Demo: [github.com/joelyancey/finalProject\\_neuralNetwork](https://github.com/joelyancey/finalProject_neuralNetwork) / [youtu.be/AZcQfiHGj8g](https://youtu.be/AZcQfiHGj8g)

## COMPUTER SCIENCE COURSEWORK

<b>C++</b>	Advanced Programming (PIC 10C, UCLA) Intermediate Programming (PIC 10B, UCLA) Introduction to Programming (PIC 10A, UCLA)	<b>Python</b>	Algorithms in Bioinformatics (CS CM 122, UCLA) Computational Genetics (CS CM 124, UCLA) Computational and Systems Biology (CS M184, UCLA)
<b>UNIX &amp; Shell</b>	Shell Programming (CSE-40079, UCSD Ext) UNIX & Linux Security (CSE-41272, UCSD Ext) UNIX & Linux Sys Admin. I (CSE-41269, UCSD Ext) UNIX & Linux Sys Admin. II (CSE-41270, UCSD Ext)	<b>R</b>	Probability Theory (STATS 100A, UCLA) Statistical Methods for Life Sciences (STATS 13, UCLA)
		<b>Discrete Math</b>	Design & Analysis of Algorithms (CSE 101, UCSD) Linear Algebra and Applications (MATH 33A, UCLA) Intro to Discrete Structures (MATH 61, UCLA)