

# Jessica Lumian, PhD

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## 7+ Years Bioinformatics Experience

- Generated five novel Cyanobacteria genomes from metagenomic data
- Elucidated environmental stress responses from genome content
- Implemented cutting-edge k-mer-based search algorithm for metagenome classification

## Microbial & Molecular Biology Expertise

- 5+ years experience analyzing microbial mat communities with bioinformatics techniques, including 3 dissertation chapters and 1 first author publication
- 8+ years benchwork experience isolating and culturing microbes, and shotgun and long read library prep

## Bioinformatics Training & Cross-Functional Teamwork

- Organized 30+ bioinformatics workshops on NGS techniques scripting, and cloud computing
- Created [Asset Inventory](#) application product according to stakeholder needs using SQL for data organization
- Product Owner for NIH Common Fund metadata catalog [portal](#)

## PROFESSIONAL HIGHLIGHTS

University of California, Davis, Davis, CA

Sept 2016 – Mar 2022 (PhD), Mar 2022 – Present (Postdoc)

### Product Owner | Postdoctoral Scholar for NIH Common Fund Data Ecosystem

#### *Product Owner/Software Engineer*

- Coordinated product strategy for National Institute of Health [Common Fund Data Ecosystem Portal](#)
- Ran Help Desk to troubleshoot internal issues and answer external customer questions

#### *Bioinformatics Training*

- Organized and led hackathons and training events about bioinformatics and computational skills (NGS sequence analysis, cloud computing with AWS, UNIX skills) and the CFDE Portal

### Research Postdoctoral Scholar at UC Davis Data Intensive Biology Lab

- Pursued cutting-edge bioinformatics research focusing on detection of genomes using large-scale k-mer searching
- Built curriculum and instructed research-focused bioinformatics class using NGS techniques and R

### PhD Candidate at UC Davis

- Assessed phylogeny, metabolic potential, and ecology of novel microbial genomes from diverse environments

### Bioinformatics Training Coordinator at UC Davis

- Coordinated over 30 bioinformatics workshops, developed instructional materials, and participated in teaching. Topics included NGS workflows, cloud computing, and programming languages

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**Governor's Office of Planning and Research**, Sacramento, CA

Apr 2021 – Mar 2022

## **Policy Fellow at the California Initiative to Advance Precision Medicine**

- Developed [Asset Inventory](#) application product based on feedback from stakeholders: biotechnology companies, community organizations, and governmental policy groups
- Piloted impact assessment initiative team including interviewing stakeholders to determine needs, creating report structure, and delegating tasks

## **EDUCATION**

### **PhD in Microbiology, University of California, Davis, Davis, CA, 2022**

- Awarded a competitive National Science Foundation Graduate Research Fellowship Program fellowship
- 1 first author [publication](#)

### **BS in Biochemistry and Molecular Biology, Michigan State University, East Lansing, MI, 2015**

- 10 [publications](#) about structural chemistry
- Participated in collaborative programming of [khmer bioinformatics software](#) (on author list as Jessica Mizzi)

## **SKILLS**

### **Technical Skills**

- Cloud, Data Analysis and Management: Amazon Web Services, Jupyter Notebooks, NGS data analysis
- Workflow Management and Automation: SnakeMake, Git/GitHub
- Programming Languages: Bash scripting, Python, R
- Communication: Markdown, ReStructured Text, Jira

### **Organizational Skills**

- Project management in cross-functional teams
- Customer-facing communication
- Logistical planning of online and in-person scientific events