

**NOW HIRING. APPLY BY MARCH 14TH.**

# **Research Technician (Bioinformatics Analyst)**



## ***Computational Biology Core, Institute for Systems Genomics***

The Computational Biology Core provides data analysis, consulting, and training to support research across UConn. Analysts collaborate on sequencing, workflows, and HPC support while leading workshops and managing projects.

### **DUTIES AND RESPONSIBILITIES**

- Support of the HPC resources.
- Support users through installing open-source scientific software and troubleshooting user code
- Manage and update shared databases
- Develop curriculum and lead training workshops
- Conduct analysis of high throughput genomic data for the research community
- Triage requests received via Slack, and the e-mail-based help desk

### **MINIMUM QUALIFICATIONS**

- M.A./M.S. in computational or biological sciences
- Experience with analyzing high-throughput sequence data
- Proficiency with Linux
- Comfort interacting with an HPC system
- Strong presentation and communication skills
- Positive attitude and ability to work as part of a team (multi-tasking extraordinaire)

### **PREFERRED QUALIFICATIONS**

- Comfort with R and/or Python programming
- Proficiency with open-source software installation

### **APPOINTMENT TERMS**

This temporary position (6 months) requires 40 hours per week at \$34.00 per hour, with the potential to transition into a long-term staff scientist role. It can be on-site or hybrid, but candidates must reside in Connecticut. Sponsorship is not available.

### **HOW TO APPLY**

To apply, please email a resume and cover letter detailing your qualifications to [karelyn.lambert@gmail.com](mailto:karelyn.lambert@gmail.com), with the subject line "Research Technician Application."

Screening will begin immediately.

