# **Aidan Lakshman**

ahl27@pitt.edu • ahl27.com • 1 (724) 612 9940

### **EDUCATION**

### University of Pittsburgh, School of Medicine

Doctor of Philosophy, Biomedical Informatics; expected graduation Summer 2025

Dissertation: Comparative genomic methods to reveal functional associations among proteins

### **University of Central Florida**

Bachelor of Science, Mathematics; magna cum laude, 2020

- Burnett Honors College
- National Merit Scholar

# WORK EXPERIENCE

# University of Pittsburgh, Pittsburgh, PA

Graduate Researcher, 2020 - 2025

- Designed novel algorithms to predict gene function using evolutionary signal; preliminary results formed the basis of a successfully funded U01 grant, results in review at *Nature Biotechnology*
- Optimized network clustering algorithms to process graphs with millions of nodes in constant memory complexity
- Created domain-specific network clustering algorithms to predict metabolic pathways from genome data
- Managed the Wright Lab's technical infrastructure, CI/CD pipelines, and code repositories
- Researched new approaches to infer causal relationships among variables in the presence of data missingness

# **Amazon Web Services**, Herndon, VA [Virtual]

Software Development Engineer Intern, Summer 2020 & 2021

- Led a team to implement a robust testing framework for Research Service Workbench on AWS (RSW)
- Streamlined RSW user experience by redesigning frontend components and building new backend infrastructure
- Implemented frontend components using React, backend components with Node.js and AWS Lambda

# Carnegie Mellon University, Intelligent Coordination and Logistics Lab

Robotics Institute Summer Scholar, Summer 2018

- Optimized traffic signal control algorithms with Bayesian hierarchical modelling to predict bus behavior
- Developed assistive technology for mobility impaired pedestrians using cellular and DSRC GPS data

# Software Engineering Institute, CERT Division, Carnegie Mellon University

Data Science / Software Engineering Intern, Summer 2017

- Identified trends in malware execution behavior using Apache Spark and Python
- Developed a Python program to simulate web traffic and user activity for cyberdefense training environments

### GRANT FUNDING

### **R Consortium**, Infrastructure Steering Committee

"Critical Updates to Biostrings", 2024 – 2025 [Award: \$8,000]

- Became the primary maintainer of Biostrings, an open source R package with >1M downloads per year
- Optimized internal methods, implemented CI/CD pipelines, handled bug reports, and added unit testing

### **University of Central Florida**

Burnett Research Scholars Grant, 2018 – 2019 [Award: \$3,000]

Optimized embodied evolutionary robotic systems for multi-foraging problems by incentivizing exploration

# SKILLS

# **High Performance Computing**

- Over 3.5 million compute hours on HTCondor systems
- Passed AWS Cloud Practitioner Certification Exam (2020-2023)

# **Programming Languages**

- Expert proficiency: R (packages: Biostrings, SynExtend, froth; additional contributions to base R)
- Work Experience: C, Fortran, Python, JavaScript, Bash, PowerShell
- Other Experience: C#, Java, Haskell, Forth, Assembly (6502)

# **Computer Engineering**

- Designed and built a cloud storage system with multiple layers of data redundancy
- Built a 6502 computer on a breadboard and developed a Forth OS from scratch in 6502-Assembly

### **PUBLICATIONS**

Lakshman, A. and Wright, E.S. "EvoWeaver: Large-scale prediction of gene functional associations from coevolutionary signals" (Under Revision at *Nature Biotechnology*). [Preprint available on request]
Lakshman, A. and Wright, E.S. "ExoLabel: Scalable network clustering for massive datasets" (In preparation).

# SELECTED TALKS

### **ISMB 2024**

Predicting Gene Functional Associations from Coevolutionary Signals with EvoWeaver

# useR! 2024

Community Detection for Extremely Large Networks

### **Great Lakes Bioinformatics Conference**

Scalable Community Detection for Large Networks

Also organized and co-chaired special session "Scalable Analysis for Big Biological Data"

# RECOMB 2024

EvoWeaver: Large-scale prediction of gene functional associations from coevolutionary signals

■ Poster Presentation, won Best Poster award

# R Project Sprint 2023\*

• Refactored and optimized R's **dendrapply** function

# Evolution 2023\*

Protein Function from Coevolutionary Signal

#### Bioconductor 2022\*

Using comparative genomics to predict protein coevolution networks

■ Led a two hour workshop (materials available at ahl27.com/tutorials)

#### **Evolution 2022**

Protein Functional Inference using Coevolutionary Signal

# **NLM Informatics Training Conference 2022**

Ensemble Methods Improve de novo Prediction of Protein Functional Association Networks

\*Awarded merit-based travel funding

# TEACHING & ADVISING

### **Undergraduate Mentor**

Advisor, Fall 2022

- Mentored undergraduate students for a semester-long research internship program
- Designed an individualized curriculum to teach R programming for bioinformatics

### **UPMC DDCF-UI Program**

Advisor, Summer 2022

- Mentored undergraduate students for a summer-long research internship program
- Gave lectures to intern cohort

# R Programming for Scientific Research, Univ. Pittsburgh

Teaching Assistant, Fall 2021

- Graduate level course in R programming (roughly 20 students)
- Gave lectures, graded assignments, and wrote quizzes

# Artificial Intelligence Club, Univ. Central Florida

Director, 2018 – 2020

- Gave regular lectures on machine learning to classes of >30 undergraduates
- Led multiple weekly journal clubs for undergraduate students
- Coordinated sponsorship opportunities and guest speakers