# **TIMOTHY CROMBIE**

# **Postdoctoral Researcher**

Northwestern University

## Contact

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# **Programming**

R

Python bash Nextflow

### **Tools**

RStudio Jupyter git/GitHub HPC (SLURM)

# **Data Science**

Advanced statistical analysis
Computational pipelines
Machine learning
Data visualization
Markdown reports

#### Research

Experimental design
High-throughput optimization
Genetics and genomics
R/Python package development
Interdisciplinary collaboration

### **EDUCATION**

Ph.D., Biology 2009 - 2015

University of Florida

B.S., Biology 1999 - 2004

Arizona State University

### **EXPERIENCE**

#### **Postdoctoral Researcher**

2017 - Present

Northwestern University

- Generate whole-genome sequence data and apply population genomics tools to explore patterns of C. elegans sequence variation
- Optimize high-throughput assays using imaging platforms and liquid handling robots to measure various phenotypes of C. elegans
- Devise Nextflow pipelines to process experimental datasets using machine learning tools on a high-performance computing cluster
- Develop R and Python packages to facilitate reproducible data processing and distribute packages using version control on GitHub
- Mentor and manage undergraduate and graduate researchers in experimental design, data collection, statistical analysis, and reporting

#### **Postdoctoral Researcher**

2016 - 2017

University of Florida

- Built a 600-line panel of recombinant inbred lines (RILs) with C. elegans
- Prepared whole-genome sequencing libraires for the RIL panel and processed sequence data to summarize genetic variation in the lines
- Optimized a high-throughput competitive fitness assay to measure the competitive fitness of RILs using a large particle sorting instrument

#### **Graduate Researcher**

2009 - 2015

University of Florida

 Used liquid handling robots, automated imaging systems, gene expression assays, and enzyme activity assays to explore interactions of molecular stress responses during combined stress exposures

#### **Laboratory Technician**

2005 - 2009

University of California, Santa Barbara

- Ran Sanger sequencing reactions and analyzed sequence data
- Managed laboratory inventory, ordering, and undergraduate training

### SELECTED PUBLICATIONS

**Crombie TA** et al. 2022. <u>Local adaptation and spatiotemporal patterns of genetic diversity revealed by repeated sampling of Caenorhabditis elegans across the Hawaiian Islands</u>. *Molecular ecology* 

Nyaanga J, **Crombie TA**, Widmayer SJ, and Andersen, EC. 2021. <u>easyXpress:</u> An R package to analyze and visualize high-throughput C. <u>elegans</u> microscopy data generated using CellProfiler. PLoS ONE

**Crombie TA**, Tang L, Choe KP, and Julian D. 2016. <u>Inhibition of the oxidative</u> stress response by heat stress in Caenorhabditis elegans. The Journal of Experimental Biology