

# WADE ROBERTS

I have 10 years of experience conducting research independently and collaboratively. My current work heavily involves computational biology, but I also have previous experience as a field and wet lab biologist. My experience has resulted in strong expertise in genomics, transcriptomics, and phylogenetics. I have extensive experience generating and working with NGS and long read sequencing. I am highly skilled in data wrangling, statistical analysis, and data visualization using R (base and Tidyverse), Python, and the command line interface. I regularly design and execute workflows using R, Bash, and Slurm, with some experience using Snakemake. I strive for reproducibility and execute end-to-end reports and visualization using RMarkdown. For my career, I am focused on pursuing my passion for applying genomics and computational biology to scientific discovery.

## EXPERIENCE

- current  
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2018

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Postdoctoral Researcher

University of Arkansas

Fayetteville, AR

- Managed and lead multiple research projects into genome evolution and the transcriptomic response of diatoms to environmental changes. Expertise in genomic, metagenomic, transcriptomic, and phylogenomic approaches.
  - Sequenced new reference genomes for 50+ species using Illumina and Oxford Nanopore technologies. Increased the number of available genomes for diatoms 4-fold.
  - Performed bioinformatic analyses and designed workflows to analyze genomic data from 280+ genomes and transcriptomes. Provided reproducibility, interpretation, and data visualization using R and Markdown. Expansive practical knowledge of current genomics software and methodologies.
  - Used high-performance computing (HPC) servers and virtual compute nodes daily. Regularly used databases to manage and curate biological metadata.
  - Reviewed scientific and technical writing for dozens of journal articles and SOPs. Communicated research to different audiences at conferences and journal publications.
- 2018  
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2012

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Graduate Research Assistant

Washington State University

Pullman, WA

- Managed and lead multiple research projects in phylogenetics and the evolution of gene expression. Expertise in molecular genetics, plant biology, and phylogenetics.
  - Designed experiments to sequence and analyze RNA-seq data collected across multiple developmental time points in 12 species.
  - Performed bioinformatic and phylogenetic analyses to gather, clean, and analyze publicly available genetic and trait data for 500+ species.
  - Communicated research through presentations at national conferences and journal publications.
  - Designed, built, and maintained in-house computing and storage servers.

## EDUCATION

- 2018  
|  
2012

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Ph.D., Molecular Plant Sciences

Washington State University

Pullman, WA

- Dissertation: Evolutionary genomics of flower diversification in Achimenes (Gesneriaceae)

2012  
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2007

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B.S., Biology | B.A., Art

Whitworth University

Spokane, WA
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- View this Resume online with links at [wrroberts.github.io/cv/](https://wrroberts.github.io/cv/)
- ## CONTACT
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[github.com/wrroberts](https://github.com/wrroberts)

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[wrroberts.github.io](https://wrroberts.github.io)

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1 (509) 631-4330
- ## LANGUAGE SKILLS
- | Language/Skill | Proficiency Level (Relative) |
|----------------|------------------------------|
| R              | High                         |
| Linux          | High                         |
| Bash           | High                         |
| CLI            | High                         |
| Python         | Medium-High                  |
| Markdown       | Medium                       |
| AWK            | Medium                       |
| sed            | Medium                       |
| SQL            | Low                          |
- ## MORE INFO
- See full CV at [wrroberts.github.io](https://wrroberts.github.io) for more complete list of positions and publications.
- Made with the R package [pagedown](https://github.com/jgm/pagedown).  
The source code is available on [github.com/wrroberts/cv](https://github.com/wrroberts/cv).  
Last updated on 2023-06-08.



## TEACHING AND OUTREACH

- current  
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2020
- **myDiatoms**  
<https://www.mydiatoms.org/>
    - A citizen science project to introduce students, naturalists, and the public to diatoms.
- 2023  
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2020
- **Genome Assembly Workshop**  
University of Arkansas  Fayetteville, AR
    - Designed from bottom up a course to teach all stages of genome assembly, annotation, and analysis.
    - Covers raw data cleaning, assembly, contaminant removal, genome size estimation, ploidy estimation, and gene prediction. Uses R, Python, Bash, and Anaconda.
- 2023  
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2018
- **Northwest Arkansas Regional Science Fair**  
 Fayetteville, AR
    - Science project judging for middle and high school students.
- 2018  
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2013
- **Graduate Teaching Assistant**  
WSU Biology Department  Pullman, WA
    - TA and lectured for 10 semesters.
    - Courses included Intro Biology, Intro Botany, and Systematic Botany.
    - Revised course materials with new activities to better engage students with active learning and focus on PNW flora.
- 2017  
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2016
- **Plant Science Day**  
Washington State University  Pullman, WA
    - Co-chair, organizer, and exhibit lead. Provided learning activities about plant biology for local students and their families.