Wade R. Roberts

Postdoctoral Fellow

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RESEARCH INTERESTS

Evolutionary genomics; Phylogenomics and comparative biology; Diatoms; Gesneriaceae; Evolutionary-developmental biology; Genetics of flower development

EDUCATION

2018	Ph.D., Molecular Plant Sciences , Washington State University, Pullman, WA Dissertation: Evolutionary genomics of flower diversification in the magic flowers (<i>Achimenes</i> , Gesneriaceae).
	Advisor: Eric H. Roalson.
	Thesis committee: Amit Dhingra, Joanna L. Kelley, Andrew McCubbin
2012	B.S., Biology, cum laude, Whitworth University, Spokane, WA
2012	B.A., Art (Sculpture and Ceramics), cum laude, Whitworth University, Spokane, WA
	EMPLOYMENT

2018	Postdoctoral Fellow, University of Arkansas, PI: Andrew Alverson
2013-2018	Graduate Teaching Assistant, Washington State University
2012-2013	Graduate Research Assistant, Washington State University
2010-2012	Greenhouse Worker, Whitworth University

PUBLICATIONS

Published:

- 8. Onyshchenko A, **Roberts WR**, Ruck EC, Lewis JA, Alverson AJ. 2021. The genome of a nonphotosynthetic diatom provides insights into the metabolic shift to heterotrophy and constraints on the loss of photosynthesis. *New Phytologist* 232(4):1750-1764. DOI: https://doi.org/10.1111/nph.17673.
- 7. Gargas CB*, **Roberts WR***, Alverson AJ. 2020. Genome sequences of bacteria associated with the diatom *Cyclotella cryptica* strain CCMP332. *Microbiology Resource Announcements* 9:e01030-20. DOI: https://doi.org/10.1128/MRA.01030-20.
- 6. **Roberts WR**, Downey KM, Ruck EC, Traller JC, Alverson AJ. 2020. Improved reference genome for *Cyclotella cryptica* CCMP332, a model for cell wall morphogenesis, salinity adaptation, and lipid production in diatoms (Bacillariophyta). *G3: Genes, Genomes, Genetics* 10(9): 2965-2974. DOI: https://doi.org/10.1534/g3.120.401408.

- Roberts WR, Roalson EH. 2020. Co-expression clustering across flower development identifies modules for diverse floral forms in *Achimenes* (Gesneriaceae). *PeerJ* 8: e8778. DOI: https://doi.org/10.7717/peerj.8778.
- 4. Kleinkopf JA, **Roberts WR**, Wagner WL, Roalson EH. 2019. Diversification of Hawaiian *Cyrtandra* (Gesneriaceae) under the influence of incomplete lineage sorting and hybridization. *Journal of Systematics and Evolution* 57(6): 561-578. DOI: https://doi.org/10.1111/jse.12519.
- 3. **Roberts WR**, Roalson EH. 2018. Phylogenomic analyses reveal extensive gene flow within the magic flowers (*Achimenes*). *American Journal of Botany* 105(4): 726-740. DOI: https://doi.org/10.1002/ajb2.1058.
- 2. **Roberts WR**, Roalson EH. 2017. Comparative transcriptome analyses of flower development in four species of *Achimenes* (Gesneriaceae). *BMC Genomics* 18(1): 240. DOI: https://doi.org/10.1186/s12864-017-3623-8.
- 1. Roalson EH, **Roberts WR**. 2016. Distinct processes drive diversification in different clades of Gesneriaceae. *Systematic Biology* 65(4): 662-684. DOI: https://doi.org/10.1093/sysbio/syw012.

Preprints:

Çiftçi O, Alverson AJ, van Bodegom P, Duijsings D, **Roberts WR**, Mertens A, van de Vijver B, Trobajo R, Mann D, Pirovano W, van Eijk I, Gravendeel B. Phylotranscriptomics reveals the reticulate evolutionary history of a widespread diatom species complex. *BioRxiv*. DOI: https://doi.org/10.1101/2022.04.11.487918.

(* indicates equal contribution)

CONTRIBUTED DATASETS

Phylogenomics of Thalassiosirales. 2022. NCBI BioProject: PRJNA825288.

Thalasssiosira oceanica CCMP1005 genome version 2.0. 2021. Zenodo: https://doi.org/10.5281/zenodo.4589594.

Nitzschia sp. Nitz4 Genome sequencing and assembly. 2020. NCBI BioProject: PRJNA412514.

- Cyclotella cryptica strain: CCMP332 Genome sequencing and assembly. 2020. NCBI BioProject: PRJNA628076.
- **Roberts WR**, Roalson EH. 2019. Data from: Phylogenomic analyses reveal extensive gene flow within the magic flowers (*Achimenes*). Dryad, Dataset, https://doi.org/10.5061/dryad.9202s.
- Kleinkopf JA, **Roberts WR**, Wagner WL, Roalson EH. 2019. Data from: Diversification of Hawaiian *Cyrtandra* (Gesneriaceae) under the influence of incomplete lineage sorting and hybridization. Dryad, Dataset, https://doi.org/10.5061/dryad.s7h937n.

Diversification of Hawaiian Cyrtandra (Gesneriaceae). 2019. NCBI BioProject: PRJNA531912.

Transcriptome sequencing of flower development in Achimenes. 2018. NCBI BioProject: PRJNA435759.

- Phylogenomic analysis of gene flow between species of *Achimenes* (Gesneriaceae). 2017. NCBI BioProject: PRJNA401042.
- Comparative transcriptome analyses of flower development in four species of *Achimenes* (Gesneriaceae). 2016. NCBI BioProject: PRJNA340450.
- Roalson EH, **Roberts WR**. 2016. Data from: Distinct processes drive diversification in different clades of Gesneriaceae, Dryad, Dataset, https://doi.org/10.5061/dryad.1br13.

RESEARCH GRANTS AND AWARDS

2020	Co-PI: Community Science Program, Joint Genome Institutes
	[ID 506537100]: 100 Diatom Genomes Project. PI: Thomas Mock.
2016	Co-PI: Doctoral Dissertation Improvement Grant, National Science Foundation
	[DEB 1601003]: Evolution of gene expression in floral diversification of Neotropical
	Gesneriaceae. PI: Eric Roalson. \$19,323.
2016	PI: Elvin McDonald Research Endowment Fund, The Gesneriad Society.
	Characterizing a red-to-blue flower color transition in <i>Achimenes</i> (Gesneriaceae). \$1,750.
2012	PI: Global Plant Sciences Initiative Fellowship, Washington State University. \$10,000.
2016	Best Poster Presentation, Inland Northwest Genomics Symposium.
2015	Travel Award, Botany meeting, Edmonton, AB, American Society of Plant Taxonomists. \$300.
2011	Synthetic Biology Research Experience (REU) Fellow, National Science Foundation, North
	Carolina State University. PI: Jenny Xiang.

PRESENTATIONS

Departmental Seminars:

- **Roberts WR.** 2017. Evolutionary genomics in the magic flowers (*Achimenes*, Gesneriaceae). Washington State University.
- **Roberts WR.** 2016. Floral diversification in Gesneriaceae: macroevolutionary and genomics approaches. Washington State University.

Contributed Talks:

- **Roberts WR**, Alverson AJ. 2021. Phylogenomic analysis of the centric diatom order Thalassiosirales based on genome and transcriptome data. Botany meeting, Virtual.
- **Roberts WR**, Alverson AJ. 2020. Evidence for a recent whole-genome duplication in a diatom genome. Botany meeting, Virtual.
- **Roberts WR**, Alverson AJ. 2019. Whole genome shotgun phylogenomics in the Thalassiosirales. North American Diatom Symposium, Eatonton, GA.
- **Roberts WR**, Roalson EH. 2018. Gene co-expression network connectivity is an important determinant of selective constraint during flower diversification in the magic flowers (*Achimenes*, Gesneriaceae). Botany meeting, Rochester, MN.
- Kleinkopf JA, **Roberts WR**, Wagner WL, Roalson EH. 2018. Detecting hybridization in Hawaiian *Cyrtandra* (Gesneriaceae) using genome-wide data. Botany meeting, Rochester, MN.

- **Roberts WR**, Roalson EH. 2017. Genomic evidence for gene flow between species of magic flowers (*Achimenes*, Gesneriaceae). Evolution meeting, Portland, OR.
- **Roberts WR**, Roalson EH. 2017. Dissecting floral diversification in the magic flowers (*Achimenes*, Gesneriaceae). Biological Sciences Graduate Student Symposium, Washington State University.
- **Roberts WR**, Roalson EH. 2015. Understanding flower diversification in *Achimenes* (Gesneriaceae) using a comparative transcriptomics approach. Botany meeting, Edmonton, AB.

Posters:

- Ruck EC, **Roberts WR**, Nakov TN, Alverson AJ. 2019. Parallel sequencing of diatom plastid genomes using a bait-capture approach.
 - North American Diatom Symposium, Eatonton, GA.
- **Roberts WR**, Roalson EH. 2018. Phylogenomic analyses reveal extensive gene flow within the magic flowers (*Achimenes*, Gesneriaceae).
 - Botany meeting, Rochester, MN.
 - Academic Showcase, Washington State University.
- **Roberts WR**, Roalson EH. 2017. Exploring phylogenetic relationships in *Achimenes* (Gesneriaceae) using transcriptome sequencing.
 - School of Biological Sciences Graduate Student Symposium, Moscow, ID.
 - Plant Sciences Retreat, Washington State University, Pullman, WA.
- **Roberts WR**, Roalson EH. 2016. Using comparative transcriptomics to understand flower diversification: an example from *Achimenes* (Gesneriaceae).
 - Inland Northwest Genomics Research Symposium, Moscow, ID.
 - Academic Showcase, Washington State University.
 - Plant Sciences Retreat, Washington State University.
- **Roberts WR**, Roalson EH. 2015. Understanding flower diversification in *Achimenes* (Gesneriaceae) using a comparative transcriptomics approach.
 - Pan-Am Evo-Devo inaugural meeting, Berkeley, CA.
- **Roberts WR**, Roalson EH. 2015. Geographic processes drive diversification in different clades of Gesneriaceae.
 - School of Biological Sciences Graduate Student Symposium, Moscow, ID.
 - Plant Sciences Retreat, Washington State University, Pullman, WA.
- **Roberts WR**, Yi Y, Xiang L, Xiang J. 2011. Cloning and evolutionary analysis of *SEPALLATA3* genes in dogwoods—deciphering the genetic links to bract petaloidy.
 - North Carolina State University Undergraduate Summer Research Symposium, Raleigh, NC.

TEACHING EXPERIENCE

2018,	Guest Lecturer, Washington State University
	Systematic Botany (undergraduate, Biol 332)
2013-2018	Graduate Teaching Assistant, Washington State University
	Systematic Botany, Biol 332 (3 semesters)
	Introductory Botany, Biol 120 (3 semesters)
	Introductory Biology: Organismal Biology, Biol 106 (5 semesters)
2012	Supplemental Instructor, Whitworth University
	General Biology I, Biol 140 (1 semester)
2010-2012	Undergraduate Teaching Assistant, Whitworth University
	Plant Biology, Biol 153 (2 semesters)

STUDENT MENTORSHIP

Graduate students:

2020	Kathryn Dupree, Ph.D. student, University of Arkansas
	Population genomics of <i>Lonicera maackii</i>
2017-2018	Joseph Kleinkopf, M.S. student, Washington State University
	Phylogenomics of Hawaiian Cyrtandra
2017	Raimundo Luciano Soares Neto, visiting scholar, Universidade Federal de Pernambuco
	DNA extractions and lab work for Cleomaceae

Undergraduate students:

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SERVICE AND OUTREACH

2020-current	myDiatoms, citizen science outreach project, University of Arkansas
2022	Student project judge, Northwest Arkansas Science and Engineering Fair
2021	Session chair, Botany meeting, virtual
2020	Student project judge, Northwest Arkansas Science and Engineering Fair
2019	Student project judge, Northwest Arkansas Science and Engineering Fair
2019	Volunteer, Secchi Day at Beaver Lake, Rogers, AR
2018	Volunteer botanist, Palouse Conservation District, Steptoe Butte State Park, WA
2016-2018	Faculty representative, Molecular Plant Sciences Graduate Student Organization
2017	Co-lead organizer, Plant Science Day, Washington State University
2014-2016	Co-chair, Palouse Discovery Science Center Committee
2016	Exhibit leader, Plant Science Day, Washington State University
2015	Native plant guide, Palouse Outdoor Science Day, Washington State University
2014 - 2015	Greenhouse tour guide, Biology Open House, Washington State University

Scientific Peer review: Biodiversitas Journal of Biological Diversity (1) – Biological Procedures Online (2) – BMC Genomics (1) – Communications Biology (2) – Ecology and Evolution (2) – Evolutionary Bioinformatics (1) – Genes (5) – Molecular Ecology Resources (2) – PeerJ (5) – PhytoKeys (12) – Plant Cell Reports (1) – Plant Growth Regulation (3) – Plants (1) – PLoS One (2) – Systematic Biology (2) – Taxon (4)

Professional Society Membership: Botanical Society of America; Society for the Study of Evolution; The Gesneriad Society

FIELDWORK AND FLORISTICS

Northwest Arkansas (2018-current); Eastern Washington (2011-2018); Hudson Biological Reserve, Smoot Hill, WA (2016-2018); Palouse Prairie, Whitman County, WA (2016-2018).

HONORIFIC NAMES

2007 'Wade Riley', M. Roberts, Tall Bearded Iris, The American Iris Society