Wade Roberts

Ph.D. Candidate

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Education

Washington State University, Pullman, WA I Aug 2012 – May 2017

Ph.D. in Molecular Plant Sciences

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

Advisor: Dr. Eric Roalson

Committee: Drs. Amit Dhingra, Joanna Kelley, and Andrew McCubbin

Whitworth University, Spokane, WA | Sep 2007 - May 2012

B.S. with Honors in Biology

B.A. with Honors in Art, sculpture and ceramics Cumulative GPA: 3.60: Laureate Society

Advisors: Dr. Lee Anne Chaney and Ms. Katie Creyts

Publications

Roalson E.H., Roberts W.R. 2016. Distinct processes drive diversification in different clades of Gesneriaceae. Systematic Biology. In press. DOI: 10.1093/sysbio/syw012

Roberts W.R., Roalson E.H. Characterization and comparative analyses of floral transcriptomes in four species of *Achimenes* (Gesneriaceae). (in prep)

Roberts W.R., Crabb J.L., Dhingra A., Roalson E.H. Micropropagation and *Agrobacterium*-mediated genetic transformation of *Achimenes erecta* (Gesneriaceae). (in prep)

Presentations

Using comparative transcriptomics to understand flower diversification: an example from *Achimenes* (Gesneriaceae)

Botany, Savannah, GA I Jul 2016 (invited talk)

WSU SBS Graduate Student Symposium, Pullman, WA I Mar 2016 (invited symposium + poster)

WSU Academic Showcase, Pullman, WA I Mar 2016 (poster) WSU Plant Sciences Retreat, Pullman, WA I Mar 2016 (poster)

Understanding flower diversification in *Achimenes* (Gesneriaceae) using a comparative transcriptomics approach

Pan-Am Evo Devo, Berkeley, CA I Aug 2015 (poster) Botany, Edmonton, AB I Jul 2015 (invited talk)

Geographic processes drive diversification in different clades of Gesneriaceae

WSU Plant Sciences Retreat, Pullman, WA I Mar 2015 (poster)

WSU SBS Graduate Student Symposium, Pullman, WA I Feb 2015 (poster)

Research

Research Assistant Roalson Lab, Washington State University I Aug 2012 – Present PI: Dr. Eric Roalson, School of Biological Sciences

Transcriptome sequencing, assembly, and characterization in *Achimenes* (Gesneriaceae) Developmental genetics and biochemistry of flower color in *Achimenes* (Gesneriaceae)

Phylogenetics, biogeography, and diversification of Gesneriaceae

Global Carex Project, contributor

Construction and maintenance of high performance computational server

Independent Undergraduate Research Pond Lab, Whitworth University I Sep – Dec 2011 PI: Dr. Finn Pond, Biology Department

Bioinformatics and cloning of putative R-body genes from Rhodospirillum rubrum

NSF REU Fellow Xiang Lab, North Carolina State University I May – Aug 2011

PI: Dr. Jenny Xiang, Department of Plant and Microbial Biology

Fellowship in Synthetic Biology

Cloning and phylogenetic analysis of SEP3 genes from dogwoods

Teaching Graduate Teaching Assistant Washington State University I Aug 2013 - Present

> Biol 332 Systematic Botany (x2) Biol 120 Introductory Botany (x2)

Biol 106 Introductory Biology: Organismal Biology (x5)

Undergraduate Teaching Assistant Whitworth University I Feb 2010 – May 2012

Bio 363 Genetics

Bio 153 Plant Biology (x2)

Bio 141 General Biology I: Organismal Biology

Service + Outreach Co-Lead Palouse Discovery Science Center Committee I Nov 2014 - Present

Organized hands-on workshops in plant science for children and families

Creation and implementation of permanent plant biology exhibit

Exhibit Lead Plant Science Day | Apr 2016

Organized and lead exhibit on plant pigments and pH for local children and families

Native Plant Guide Palouse Outdoor Science Day I Apr 2015

Organized and lead short hikes for children and families to identify and collect wildflowers

Greenhouse Tour Guide Biology Science Day I Oct 2014, 2015

Lead informational tours of the living plant collection at Washington State University

Recruitment Mentor Molecular Plant Sciences Graduate Program I Mar 2013, 2014

Lead invited recruits to and from faculty interviews

Programming

Languages: proficient in R, Python, Perl, Bash Operating systems: Unix, Linux, MacOS, Windows

Bioinformatics and Genomics

Sequencing platforms: Illumina, Ion Torrent, Roche 454, Sanger

Assembly: Trinity, Velvet, Oases, SOAP, CLC

Analysis: BLAST, Bioconductor, GATK, KEGG, OrthoMCL Phylogenetics: RAxML, MrBayes, BEAST, PAML, HyPhy Machine learning: clustering, PCA, HMMER, WGCNA

Molecular Biology

DNA: extraction, purification, PCR, Southern blotting, cloning

RNA: extraction, purification, cDNA preparation, qPCR, RT-PCR, RNAseq library construction

Genetic engineering: RNAi

Culture: bacterial cell culture, plant tissue culture, micropropagation Biochemistry: HPLC, TLC, pigment extraction and purification

Data Visualization

R (ggplot2), Python (matplotlib), Adobe Suite

Selected Awards 2016 NSF Doctoral Dissertation Improvement Grant (\$19,323)

2015 Student Travel Grant, American Society of Plant Taxonomists (\$300)

2012 Global Plant Sciences Initiative Fellowship, Washington State University (\$10,000) 2011 NSF REU Fellowship in Synthetic Biology, North Carolina State University (\$5,000)

2007 Trustee Scholarship, Whitworth University (\$18,000 x4)

Skills