Amanda Charbonneau

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Research and Professional Experience

June 2019 - present	Bioinformatics Consultant , Reimers Lab, MSU Create code and scripting necessary to complete a large psychiatric genomics study. GWAS results from multiple found studies and SNP locations for a number of genomic features are combined using a regression framework (LDSR) to determine whether traits are enriched for SNPS in various classes of genomic feature.
May 2018 - present	Program Manager , Common Fund Data Ecosystem, (formerly the <u>DCPPC</u>) Organize and manage a cross-disciplinary, multi-institution consortium to build the community and infrastructure required to meaningfully connect Common Fund data assets. Facilitate inter-group interactions as well as collaborations and outreach with other scientific organizations. Maintain and create policies and procedures and act as reporting intermediary between awardees and the NIH.
2011 – 2018	Ph.D in Genetics, MSU Completed: 4/27/2018 Research Topic: Determining the loci that underlie the differences between weedy R. raphanistrum and its non-weedy relatives. Used population genomics techniques to try to determine the source population(s) for weedy radish and crop radishes, as well as how they have evolved phenotypically and genetically.
Summer 2014 & 2015	Course Coordinator, <u>Next-Gen Sequence Analysis Workshop</u> Logistics and troubleshooting for two-week intensive learning course, in addition to
	teaching (see below).
2007 – 2011	teaching (see below). Research Technologist , Biochemistry and Molecular Biology, MSU Genomic screening of secondary metabolites in tomato trichomes from wild and cultivated species. Genetic and biochemical characterization of genes causing qualitative differences between wild and cultivated varieties of tomato. Screening of morphological and biochemical phenotypes for Arabidopsis 2010 initiative
2007 – 2011 2005 – 2007	Research Technologist , Biochemistry and Molecular Biology, MSU Genomic screening of secondary metabolites in tomato trichomes from wild and cultivated species. Genetic and biochemical characterization of genes causing qualitative differences between wild and cultivated varieties of tomato. Screening of morphological and biochemical phenotypes for Arabidopsis 2010

2001 – 2005 BA Biochemistry & Criminal Justice, Olivet College, Olivet, Michigan, 49076

Teaching & Outreach

Summer 2012, 2013, 2014, 2015, 2016 & 2018, 2019	Instructor , <u>Next-Gen Sequence Analysis Workshop</u> <i>Teach a wide variety of computing topics including basic unix,</i> <i>bioinformatics and cloud computing</i>
2017 - present	Maintainer, Data Carpentry Genomics Lessons Remote and Cloud Computing
2015 – present	Instructor , <u>Software Carpentry</u> and <u>Data Carpentry</u> Workshops taught for: <u>Notre Dame</u> , <u>U of M</u> , <u>James Madison</u> , <u>MSU</u> , <u>MSU</u>
2008 – present	Docent , <u>Potter Park Zoo</u> , Lansing , Michigan Teach the scientific method & introduce research methods for the "BIG Zoo Lesson". Use observation of animal behavior to teach children core science principles. Train animals and animal handlers for outreach
2017 - 2018	Genetics Officer, Forum and career workshop coordinator <i>Organize professional events for the Genetics Program</i>
2014 - 2018	Education Staff , <u>Potter Park Zoo</u> , Lansing , Michigan Organize and manage individual educational special events such as birthdays and overnights
Summer 2016, 2017 & 2018	Instructor , Evolution, MSU Develop and teach entirely online course through D2L
Spring 2016, 2017 & 2018	Lab Instructor, Cells and Development, MSU Teach cell culture and experimental design
Fall 2016 & 2017	Teaching Assistant , Histology, MSU Teach microscopy and micro-anatomy. Half-time recitation appointment
2010, 2015, & 2016	Presenter , <u>Grandparents University</u> , <u>MSU</u> Science of Smells and Tastes
Summer 2016	Instructor, Fundamental Genetics, MSU Traditional on-campus course in basic genetics
Feb 2016	Instructor , Introduction to LaTeX, <u>DIB Intermediate Bioinformatics</u> <u>Training</u> <i>Develop and teach LaTeX on Overleaf, for beginners</i>
Jan 2016	Instructor , <u>RNAseq Workshop</u> , ICER, MSU Teach single workflow to novices over two days
Fall 2015 - Spring 2016	Organizer/Instructor, <u>NGS Continuation course</u> Monthly evening session to teach practical computing to biologists

Fall 2015	Teaching Assistant , Genetics, MSU Half-time recitation appointment
2013 – May 2015	GK-12 Fellow , Kellogg Biological Station Act as a 'visiting scientist' in local schools helping to both teach and improve upon the science curriculum for elementary, middle and high school classrooms
2013 - 2015	Judge, University Undergraduate Research and Arts Forum, MSU
Sept. 2013	Teaching Assistant , <u>Software Carpentry</u> short course Teach a wide variety of computing topics including basic unix, programing and data management

Publications

Clarke, Daniel, et.al. (2019) FAIRshake: toolkit to evaluate the findability, accessibility, interoperability, and reusability of research digital resources. BioRxiv. <u>https://doi.org/10.1101/657676</u>

Charbonneau A, Tack D, Lale A, et al. (2018) Weed evolution: Genetic differentiation among wild, weedy, and crop radish. *Evol Appl.* ;11:1964–1974. https://doi.org/10.1111/eva.12699

Charbonneau, Amanda (2018) Rapid Adaptation of Floral Phenotypes in Weedy radish, *R.r. Raphanistrum.* Michigan State University, ProQuest Dissertations Publishing.

LaTeX tutorial presented for DIB Intermediate Bioinformatics Training

Publicly available lessons for GK-12 Bioenergy Sustainability Project [2013, 2013, 2014, 2015]

Publicly available lessons for Next-Gen Sequencing Workshop [2012, 2013, 2014, 2015, 2016]

Invited blog post, **BEACON** Researchers at Work, June 2014

Schilmiller, A., **Charbonneau**, **A**., Last, R. (2012) Identification of a BAHD acetyltransferase that produces protective acyl sugars in tomato trichomes. <u>PNAS</u>, 40, 16377-16382.

Schilmiller, A., Shi, F., Kim, J., Charbonneau, A., Holmes, D., Jones, A., and Last, R. (2010) Mass spectrometry screening reveals widespread diversity in trichome specialized metabolites of tomato chromosomal substitution lines. <u>Plant Journal</u>, 62, 391-403.

Schilmiller, A., Schauvinhold, I., Larson, M., Xu, R., Charbonneau, A., Schmidt, A., Wilkerson, C., Last, R., and Pichersky, E. (2009) Monoterpenes in the glandular trichomes of tomato are synthesized from a neryl diphosphate precursor rather than geranyl diphosphate. <u>PNAS</u>, 26, 10865–10870.

Presentations

Midwest PopGen Conference, 2017	Talk: Weed evolution: Genetic differentiation among wild, weedy, and crop radish
Evolution Conference, 2014	Talk: Evolution of Weed phenotypes in Radish

BEACON Congress, 2013	Talk: The Evolution of Weediness
Ecological Genomics Symposium, 2012	Poster: Evolution of weedy and crop phenotypes in radish
BEACON Congress, 2012	Poster: Evolution of crops and weeds: Loss of over- wintering phenotypes in wild radish

Attended Workshops and Symposiums

Feb 2016	DIB Intermediate Bioinformatics Training, Bodega Bay, CA
March 2015	Data Carpentry Genomics and Assessment Hackathon Helped to create the lesson content for the Genomics workshop
Jan. 2015	Software Carpentry Instructors Workshop
July 2013	18th <u>Summer Institute in Statistical Genetics</u> Modules taken: Quantitative Genetics, Mixed Models in Quantitative Genetics, Coalescent Theory
May 2012	Software Carpentry Scientific computing and software design
Awards	
2013 – 2015	Fellow – Kellogg Biological Station GK-12 Partner with rural school teachers to help improve science lessons and learn teaching skills, two full years of financial support
2013	Honorable Mention – National Science Foundation Graduate Research Fellowship Program
2011 – 2012	Excellence in BioMolecular Science Fellowship – College of Natural Sciences, MSU <i>One year of full financial support</i>
2011	'Good Egg Award' – <u>Potter Park Zoo</u> , Lansing Michigan <i>Docent of the year award for more than 400 hours of service in 2011</i>

References

Available upon request