

Maximilian Oliver Press, PhD

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EDUCATION

University of Washington

Seattle, WA

PhD, Genome Sciences

June 2016

Dissertation: "Observations about the effects of epistasis on evolution and complex traits."

Reed College.

Portland, OR

BA, Biology. Phi Beta Kappa

May 2008

Thesis: "Microevolution of *Gasterosteus aculeatus* in the Johnson Creek watershed."

RESEARCH EXPERIENCE

University of Washington

Seattle, WA

Postdoctoral Fellow; advisor: Christine Queitsch.

2016-2017

Highly accurate short tandem repeat genotyping reveals heritable variation, functional effects, and signatures of selection.

- Optimized sequencing assays for genotyping short tandem repeats (STRs).
- Optimized software pipelines for high-throughput sequencing
- Performed functional and evolutionary analysis of natural STR variation in *Arabidopsis thaliana*.

Graduate Research Assistant; advisors: Christine Queitsch and Elhanan Borenstein. 2010-2016

Evolutionary constraint of gene acquisition in prokaryotic evolution.

Developed computational methods for studying epistasis in bacterial genome evolution

Contribution of microsatellites to phenotypic and epistatic variance.

- Developed genetic analysis methods for studying microsatellite variation and epistasis in *Arabidopsis thaliana*

Independent genetic pathways for thermoresponsive flowering and thermomorphogenesis in *A. thaliana*.

- Phenotyping and genetic analysis of reaction norms in *A. thaliana* strains.

Fred Hutchinson Cancer Research Center

Seattle, WA

Research Technician; supervisor: Sue Biggins.

2008-2010

- Genetic and molecular characterization of histone function in chromosome segregation.

Reed College

Portland, OR

Undergraduate Researcher; advisor: Robert Kaplan

2007-2008

- Molecular and phenotypic analysis of differentiation in stickleback populations isolated by urban barriers.

RELATED PROFESSIONAL EXPERIENCE

Phase Genomics Inc.

Seattle, WA

Senior Bioinformatics Engineer

2017-present

- Performed computational biology research and technical writing in an industrial setting.
- Designed algorithms, developed software in a cutting-edge genomics service company.
- Worked actively with customers to provide solutions to problems using Hi-C data.

FELLOWSHIPS AND AWARDS

- 2015: UW Graduate School Fund for Excellence and Innovation Travel Grant

- 2011-2013: NIH NHGRI Genomics Training Grant 2T32HG35-16
- 2008: Phi Beta Kappa (Reed College)
- 2007-8: Miller Undergraduate Research Foundation grant, Reed College
- 2004-8: 3 Commendations for Excellence in Scholarship, Reed College

PUBLICATIONS (see also [Pubmed](#), [bioRxiv](#), [Google scholar](#))

- **Press MO**, Queitsch C (2017). "Variability in a Short Tandem Repeat Mediates Complex Epistatic Interactions in *Arabidopsis thaliana*." *Genetics*, 205(1): 455-464.
- **Press MO**, Lanctot A, Queitsch C (2016). "PIF4 and ELF3 act independently in *Arabidopsis thaliana* thermoresponsive flowering." *PLOS ONE* 11(8): e0161791.
- **Press MO**, Queitsch C, Borenstein E (2016). "Evolutionary assembly patterns of prokaryotic genomes." *Genome Research* 26: 826-833.
- Carlson KD, Sudmant PH, **Press MO**, et al. (2015). "MIPSTR: a method for multiplex genotyping of germ-line and somatic STR variation across many individuals." *Genome Research* 25(5):750-761.
- Rival P*, **Press MO***, Bale J*, et al. (2014). "The conserved *PFT1* tandem repeat is crucial for proper flowering in *Arabidopsis thaliana*" *Genetics* 198(2): 747-754.
- **Press MO**, Carlson KD, Queitsch C (2014). "The overdue promise of short tandem repeat variation for heritability." *Trends in Genetics* 30(11) 504-512.
- **Press MO***, Li* et al. (2013). "Genome-scale co-evolutionary analysis identifies functions and clients of bacterial Hsp90." *PLOS Genetics* 9(8): e1003631
- Undurraga S, **Press MO**, et al. (2012). "Background-dependent effects of polyglutamine variation in the *Arabidopsis thaliana* gene *ELF3*." *PNAS* 109(47):19363-7.
- Ranjitkar P, **Press MO**, et al. (2010). "An E3 ubiquitin ligase prevents ectopic localization of the centromeric histone H3 variant via the centromere targeting domain." *Molecular Cell* 40(3): 455-64.

*: equal contribution to publication.

Manuscripts in review

- **Press MO** et al. (2017). "Massive variation of short tandem repeats with functional consequences across strains of *Arabidopsis thaliana*." bioRxiv: <https://doi.org/10.1101/145128>

Working manuscripts

- **Press MO** et al. (2017). "Hi-C deconvolution of a human gut microbiome yields high-quality draft genomes and reveals plasmid-genome interactions." bioRxiv: <https://doi.org/10.1101/198713>
- English translation of Woltereck R (1909). "Weitere experimentelle Untersuchungen über Art-veränderung, speziell über das Wesen quantitativer Artunterschiede bei Daphniden". *Verhandlungen der deutschen zoologischen Gesellschaft* 19 (1909): 110–73. In progress, available at <https://github.com/maximilianpress/miscellaneous/blob/master/experimental-investigations-variation.pdf>.
 - This classic paper originates the idea of the reaction norm in heredity and organismal development, and has not been previously translated to English.

SELECTED PRESENTATIONS

- Talk: Thermomorphogenesis Meeting 2016: "PIF4 and ELF3 Act Independently in *Arabidopsis thaliana* Thermoresponsive Flowering". Halle, Germany
- Talk: The Allied Genetics Conference, July 2016: "The variable ELF3 polyglutamine is an epistatic hub." Orlando, FL
- Talk: Congress of the Society for Molecular Biology and Evolution, July 2015: "Evolutionary Assembly Patterns of Prokaryotic Genomes." Vienna, Austria

- Talk: 6th International Conference on the Hsp90 chaperone machine, September 2012: “Evolutionary Inference of bacterial Hsp90 functions.” Les Diablerets, Switzerland

TEACHING AND MENTORING

University of Washington

- **Teaching assistant** (GENOME 351: Human Genetics for non-majors). Spring 2014
- **Teaching assistant** (GENOME 371: Introductory Genetics). Fall 2012

Reed College

- **Tutor** (Biology 101/102: Introductory Biology), 2005-2008

PEER REVIEW:

- **Reviewer**: *Nature Communications*, *Nature Methods*, *PLoS Evolutionary Biology*, *Matters*.
- **Sub-reviewer for C. Queitsch**: *PNAS*, *Nature Methods*, *PLOS Genetics*, *Human Genetics*, *American Journal of Primatology*.
- **Sub-reviewer for E. Borenstein**: *RECOMB 2012*, *Scientific Reports*.

SCHOLARLY SOCIETY MEMBERSHIPS

Genetics Society of America.

Society for Molecular Biology and Evolution.

Peer Community in Evolutionary Biology.

SERVICE AND OUTREACH ACTIVITIES

- **Co-instructor**, GRE preparation course. July 2014, July 2015, (short class for underrepresented minority undergraduates).
- **Organizer**, 2012 NHGRI UW Genome Training Grant Symposium.
- **Instructor**, Basic ideas of bioinformatics, June 2012, (short class for HS teachers).
- **Departmental coordinator**, 2012 Science Education Partnership, Genome Sciences.
- **Restoration volunteer**, Adopt-a-Stream Foundation, (Summer-Winter 2005). Wetlands restoration field work.
- **Volunteer**, Students for Empowering, Educating, Diversity and Service (SEEDS), Reed College chapter (2004-2008). Taught middle school students about wetland restoration.

REFERENCES

- **Dr. Christine Queitsch**, University of Washington Dept. Genome Sciences.
queitsch@uw.edu
- **Dr. Elhanan Borenstein**, University of Washington Dept. Genome Sciences.
elbo@uw.edu
- **Dr. Joe Felsenstein**, University of Washington Dept. Genome Sciences.
joe@gs.washington.edu
- **Dr. Sue Biggins**, Fred Hutchinson Cancer Research Center, Howard Hughes Medical Institute.
sbiggins@fhcrc.org