Schuyler D. Smith

Ph.D. Student in Bioinformatics and Computational Biology Department of Agriculture and Biosystems Engineering College of Engineering Iowa State University of Science and Technology *D* +1 (413) 212-9110

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3 Google Scholar

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Education

Ph.D. Iowa State University of Science and Technology, Ames, IA

2017 - pres.

Bioinformatics and Computational Biology

Dissertation:

Research focus: metagenomic community analysis, microbial ecology.

Advisor: Dr. Adina Howe Committee Members: Dr. Tadashi Fukami Co-Advisor: Dr. Ido Friedberg Dr. Dan Nettleton Dr. Gwyn Beattie

M.S. Texas A&M University, College Station, TX

2013 - 2015

Quantitative Genetics and Plant Breeding

Thesis: Molecular Characterization of the Texas Maize Breeding Program. Research focus: quantitative genetics, marker analysis, NIR phenotyping.

Advisor: Dr. Seth Murray

B.S. Iowa State University of Science and Technology, Ames, IA

2008 - 2012

Genetics

Experience

Iowa State University of Science and Technology

January, 2017 - pres.

Genomics and Environmental Research in Microbial Systems Lab, Ames, Iowa

Graduate Research Assistant - Ph.D.

Identifying microbial interactions in complex systems.

University of Wisconsin-Madison

June, 2015 - August, 2016

Potato Breeding and Genetics Laboratory, Madison, Wisconsin

Graduate Research Assistant - Ph.D.

Development of workflows and pipelines to automate and scale genotyping-by-sequencing and genomic prediction of quantitative traits.

United States Department of Agriculture - ARS

June, 2014 - December, 2014

Arid-Land Agricultural Research Center. Maricopa, Arizona

Biological Science Technician (Internship)

Developed high-throughput phenotyping platforms for crop traits such as canopy cover, plant height, and seed counting.

Texas A&M University

January, 2013 - May, 2015

Maize Breeding and Genetics Program, College Station, Texas

Graduate Research Assistant - M.S.

Developed a molecular characterization of the Texas maize germplasm and oleic-acid screening using near-infrared spectroscopy.

Monsanto Company

Huxley Research Station. Huxley, Iowa

Maize Plant Breeding Intern

Conducted research project on optimizing a high-throughput phenotyping platform looking at seed characteristics. Led crews in pollinating nurseries, and harvest.

DuPont Pioneer

May, 2011 - December, 2011

May, 2012 - November, 2012

Willmar Research Station. Willmar, Minnesota

Maize Product Trait Development Intern - 6 month

Conducted a QTL study for important problematic proprietary trait. Led crews for data collection in yield trials, pollination in nurseries, and harvesting.

United States Department of Agriculture - ARS

Soybean Genomics Laboratory - Graham Lab. Ames, Iowa

Student Undergraduate Research Assistant

Assisted post-doctorate researcher via PCRs, DNA extractions and preparations, gel-imaging, among other laboratory procedures.

January, 2010 - May, 2011

Publications

J Choi, EL Rieke, TB Moorman, ML Soupir, HK Allen, **SD Smith**, A Howe. Practical implications of erythromycin resistance gene diversity on surveillance and monitoring of resistance. January, 2018. FEMS microbiology ecology. academic.oup.com/femsec/94

Smith, Schuyler D., Heffner, Elliot, Murray, Seth C. Molecular analysis of genetic diversity in a Texas maize breeding program. 2015. Maydica. 60. cabdirect.org/abstract/201

Posters

- Smith, S.D., Villanueva, P.E., Fukami, T., Howe, A. Co-Occurrence Networks Reveal Key OTUs in Flower Nectar Microbiomes Across Dispersal Treatments. Presented at the 17th ISME. 2018, August 12-17. Leipzig, Germany & the NSF Research Traineeship (NRT) Annual Meeting. 2018, September 27-28. Arlington, Virginia.
- Smith, S.D., Howe, A. Examining Antibiotic Resistance Gene (ARG) horizontal transfer and introduction through farmland soil microbiomes as a result of modern farming practices. Presented at the 3rd Annual Front Rang Computational & Systems Biology Symposium: Microbiome. 2017, June 12-13. Fort Collins, Colorado.
- Smith, S.D., Endelman, J.B. Genotyping by Sequencing for Autotetraploid Species. Presented at the 5th International Conference on Quantitative Genetics. 2016, June 12-17. Madison, Wisconsin. & the 11th Annual National Association of Plant Breeders Meeting. 2016, August 15-18. Raleigh, North Carolina.

Conferences Attended

- 2018 NSF Research Traineeship (NRT) Annual Meeting. 2018, September 27-288. Arlington, Virginia.
- 17th Annual International Symposium on Microbial Ecology. 2018, August 12-17. Leipzig, Germany.
- 4th Annual Bioinformatics and Computational Biology Symposium: he Past and Future of Bioinformatics and Computational biology. 2018, March 30. Ames, Iowa.
- 3rd Annual Front Range Computational & Systems Biology Symposium: Microbiome. 2017, June 12-13. Fort Collin, Colorado.
- 3rd Annual Bioinformatics and Computational Biology Symposium: The Breadth and Depth of Bioinformatics Analysis. 2017, March 31. Ames, Iowa.
- 11th Annual National Association of Plant Breeders Meeting. 2016, August 15-18. Raleigh, North Carolina
- 5th International Conference on Quantitative Genetics. 2016, June 12-17. Madison, Wisconsin.
- American Seed Trade Association Annual Meeting. 2015, December 9-11. Chicago, Illinois.

• NCCC: Potato Breeding and Genetics Technical Committee. 2015, December 7-9. Chicago, Illinois.

Awards, Fellowships, Honors, Recognition
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Iowa State University of Science and Technology

• Selected as representative for the 2018 NRT Annual Meeting	2018
• NSF-NRT Predictive Plant Phenomics Fellow	2017 - 2018
• College of Engineering Interdepartmental Research Fellow	2017 - pres.
Academic Recognition Award	2008 - 2012

Texas A&M University

• Willie May Harris Fellow 2013 - 2014

Workshops Taught

Developer / Instructor

• Introduction to Data Analysis for Biology Graduate Students.

Developer and Instructor for Tutorial: R Basic

Teaching Assistant

• EDAMAME 2017 Explorations in Data Analyses for 2017 & 2018 Metagenomic Advances in Microbial Ecology.

• Introduction to Data Analysis for Biology Graduate Students. 2017

• BCB Data Analysis Language Workshops

Introduction to Unix 2017 & 2018
Introduction to Python 2017

Graduate Coursework Completed

M.S. Plant Breeding and Quantitative Genetics:

- Quantitative Genetics Plant Breeding I
- Statistics in Research I Plant Breeding II

• Statistics in Research II • Experimental Design • Host-Plant Resistance

Ph.D. Bioinformatics and Computational Biology:

- Bioinformatic Algorithms Fundame
- Statistical Bioinformatics
- Bioinformatic Systems
- Genomic Sciences
- Linear Mixed Models
- Fundamentals of Predictive Plant Phenomics
- Plant Genetics
- Biometric Procedures in Plant Breeding
- Advanced Plant Breeding

• Molecular and Quantitative

Genetics in Plant Breeding

- Selection Theory
- Tools for Reproducible Research