Schuyler D. Smith

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Google Scholar

github.com/schuyler-smith

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Education

Ph.D.	Bioinformatics and Computational Biology Iowa State University of Science and Technology, Ames, IA	ABD
M.S.	Quantitative Genetics Texas A&M University, College Station, TX	2015
B.S.	Genetics Iowa State University of Science and Technology, Ames, IA	2012
Experi	ience	
Iowa State University of Science and Technology Genomics and Environmental Research in Microbial Systems Lab, Ames, Iowa Graduate Research Assistant - Ph.D.		
University of Wisconsin-Madison Potato Breeding and Genetics Laboratory, Madison, Wisconsin Graduate Research Assistant - Ph.D.		2015 - 2016
United States Department of Agriculture - ARS Arid-Land Agricultural Research Center. Maricopa, Arizona Biological Science Technician (Internship)		2014
Texas A&M University Maize Breeding and Genetics Program, College Station, Texas Graduate Research Assistant - M.S.		2013 - 2015
Monsanto Company Huxley Research Station. Huxley, Iowa Maize Breeding Intern		2012
DuPont Pioneer Willmar Research Station. Willmar, Minnesota Maize Product Trait Development Intern - 6 month		2011
Soybea	States Department of Agriculture - ARS an Genomics Laboratory - Graham Lab. Ames, Iowa graduate Research Assistant	2010 - 2011

Publications

- Schuyler D. Smith, P Colgan, F Yang, EL Rieke, ML Soupir, TB Moorman, HK Allen, A Howe. Investigating the dispersal of antibiotic resistance associated genes from manure application to soil and drainage waters in simulated agricultural farmland systems. PLOS One. journal.pone.0222470 September, 2019.
- **Schuyler D. Smith** phylosmith: an R-package for reproducible and efficient microbiome analysis with phyloseq-objects. Journal of Open Source Software. 10.21105/joss.01442 June, 2019.
- 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. FEMS microbiology ecology. 10.1093/femsec/fiy006 January, 2018.
- Schuyler D. Smith, E Heffner, SC Murray. Molecular analysis of genetic diversity in a Texas maize breeding program. Maydica. cabdirect.org/abstract/201 2015.

Posters

- Smith, S.D. phylosmith: an R-package for reproducible and efficient microbiome analysis with phyloseq-objects. Presented at the 27th ISMB/18th ECCB. 2019, July 21-25. Basel, Switzerland.
- 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.

Software

phylosmith: an R-package for reproducible and efficient microbiome analysis with phyloseq-objects.

Workshops Taught

• Data Carpentry	y: Genomics hington University, Washing	D.C.	2020
o o	2020		
 Data Carpentry Upper Gwyr 	2019		
• Girls in Science	2019		
Science Cent	2019		
• Data Analyses	2010		
Kellog Resea	2017 & 2018		
o o	Data Analysis for Biology	,	
Iowa State U	2017 & 2018		
• Iowa State BCI	B Data Analysis Language	Workshops	
Introduction	2017 & 2018		
Introduction to Python			2017
· ·	y of Science and Technology		
· ·	presentative for 2018 NSF-N		2018
• College of Engineering Interdepartmental Research Fellow			2017 - pres.
NSF-NRT Predictive Plant Phenomics Fellow			2017 - 2018
Academic Recognition Award			2008 - 2012
Texas A&M Universi			
• Willie May Har			2013 - 2014
V			
Certifications			
Data Carpentry	y - Course Instructor		2017
Applicable Clri	Π_{G}		
Applicable Ski	IIS 		
• R	• Shell (BASH)	• LATEX	• Linux
• C++	• SQL	• Markdown	 Windows
• Python	• html	• Git	• MacOS