Schuyler D. Smith

Ph.D. Candidate - Bioinformatics and Computational Biology Department of Agricultural and Biosystems Engineering College of Engineering Iowa State University **A**mes, IA 50010 +1-(413)-212-9110

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8 Google Scholar

github.com/schuyler-smith

schuyler-smith.github.io

Education

Ph.D.	Bioinformatics and Computational Biology Iowa State University, Ames, IA	2021			
M.S.	Plant Breeding and Quantitative Genetics Texas A&M University, College Station, TX	2015			
B.S.	Genetics Iowa State University, Ames, IA	2012			
Experience					
Graduate Research Assistant - Ph.D. Genomics and Environmental Research in Microbial Systems Lab, Ames, Iowa Iowa State University					
Gradua Pota Univ	2015 – 2016				
Biologi Arid Unit	2014				
Maiz	nte Research Assistant - M.S. ze Breeding and Genetics Program, College Station, Texas as A&M University	2013 - 2015			
Hux	Breeding Intern ley Research Station. Huxley, Iowa Isanto Company	2012			
Maize l Will DuP	2011				

Publications

- **Schuyler D. Smith**, Choi, J., Howe, A. Diversity of Antibiotic Resistance genes and Transfer Elements-Quantitative Monitoring (DARTE-QM): a method for sequencing microbiome functional profiles. 2021.
- Yu, W., Lawrence, N. C., Sooksa-nguan, T., **Smith, S.D.**, Tenesaca, C., Howe, A. C., Hall, S. J. *Microbial linkages to soil biogeochemical processes in a poorly drained agricultural ecosystem.* 2021.
- **Schuyler D. Smith**, Colgan, P., Yang, F., Rieke, E.L., Soupir, M.L., Moorman, T.B., Allen, H.K., Howe, A. *Investigating the dispersal of antibiotic resistance associated genes from manure application to soil and drainage waters in simulated agricultural farmland systems*. 2019.
- **Schuyler D. Smith** *phylosmith:* an *R-package* for reproducible and efficient microbiome analysis with *phyloseq-objects*. 2019.
- Choi, J., Rieke, E.L., Moorman, T.B., Allen, H.K., **Smith, S.D.**, Howe, A. *Practical implications of erythromycin resistance gene diversity on surveillance and monitoring of resistance*. 2018.
- **Schuyler D. Smith**, Heffner, E., Murray, S.C. *Molecular analysis of genetic diversity in a Texas maize breeding program.* 2015.

Software

phylosmith: an R-package for reproducible and efficient microbiome analysis with phyloseq-objects. ssBLAST: an R-package of efficient C++ functions to parse BLAST outputs and wrangle corresponding FASTO/FASTA files.

simple.dada: an R-package for streamlined implementation of the dada2 processing pipeline.

schuylR: an R-package of various functions that I often find useful.

genome tools: an R-package with various functons for genomic data analysis.

linux box sync: a shell program to sync a local Box directory to the cloud.

hpcc R setup: a shell program to setup R on the MSU-HPCC.

Posters

- Smith, S.D. phylosmith: an R-package for reproducible and efficient microbiome analysis with phyloseg-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.

Workshops Taught

Awards, Fellowships, Honors, & Recognitions

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 Selected P3 representative for 2018 NSF-NRT Annual Meeting 	2018
College of Engineering Interdepartmental Research Fellow	2017 - 2020
NSF-NRT Predictive Plant Phenomics Fellow	2017 - 2018
Academic Recognition Award	2008 - 2012
xas A&M University	

2013 - 2014

Tex

• Willie May Harris Fellow

Applicable Skills

• R	 Shell (BASH) 	• \text{MEX}	 Linux
• C++	• HTML	 Markdown 	 Windows
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