

# Sarah Stevens

## PhD Candidate, Microbiology Doctoral Training Program

sstevens2@wisc.edu sarah.lr.stevens@gmail.com sarahlrstevens.info

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### Education

University of Wisconsin - Madison  
*PhD Candidate - Microbiology Doctoral Training Program- 2012-Present*  
SciMed GRS Fellow  
University of Illinois Urbana - Champaign  
*Bachelor of Science - Molecular and Cellular Biology- 2007-2011*  
Minor in Informatics  
James Scholar Honors

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### Research

University of Wisconsin - Madison, McMahon Lab  
*Graduate Research Assistant - 2012-Present*  
Classifying reference genomes (composite genomes from metagenomes and single-cell genomes) for unculturable but common freshwater bacteria and understand how the related populations change through time using our metagenomic time series.  
University of Illinois Urbana - Champaign, Metcalf Lab  
*Visiting Researcher - 2011-2012*  
Assembled genomes from isolates belonging to the archaeal genus *Methanosarcina* and analyzed shared genomic features of members of this genus.  
University of Illinois Urbana - Champaign, Whitaker Lab  
*Undergraduate Research Assistant - 2010-2011*  
Studied insertion sequences in *Sulfolobus islandicus* in an independent project, Culturing archaea, organized sample/culture database, cleaned-up and prepped lab equipment.

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### Organizations

Molecular Microbial Ecology and Evolution (MoMiEE) focus group  
*Co-Chair - 2014-Present*  
MoMiEE provides an opportunity for researchers to discuss new computational tools and learn bioinformatics skills together. In addition, I lead python study group for those interesting in learning python.  
Software Carpentry  
*Instructor - 2015-Present*  
DOE Joint Genome Institute  
*Intern/Affiliate - Jan. 2015-Mar. 2015*

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### Presentations

**Stevens, S. L. R.**, Bendall, M. L., Chan, L.-K., Malfatti, S., Schwientek, P., Tremblay, J., ... McMahon, K. D. Malmstrom, R. R. Genome-wide Selective Sweeps in Natural Bacterial Populations Revealed by Time-series Metagenomics (Oral Presentation). 15th International Symposium on Microbial Ecology. August 24, 2014. Seoul, South Korea. <https://goo.gl/6iunz0>

**Stevens, S. L. R.**, Bendall, M. L., ... McMahon, K. D. Malmstrom, R. R. Dynamics of Sequence-Discrete Bacterial Populations Inferred Using Metagenomes (Poster Presentation). 15th International Symposium on Microbial Ecology. August 24, 2014. Seoul, South Korea. <https://goo.gl/qsYL32>

**Stevens, S. L. R.**, Bendall, M. L., ... McMahon, K. D. Malmstrom, R. R. Dynamics of Sequence-Discrete Bacterial Populations Inferred Using Metagenomes (Poster Presentation). SciMed GRS Poster Session 2014. Sept. 03, 2014. Madison, WI.

**Stevens, S. L. R.**, Bendall, M. L., ... McMahon, K. D. Malmstrom, R. R. Dynamics of Sequence-Discrete Bacterial Populations Inferred Using Metagenomes (Poster Presentation). DOE Joint Genome Institute User Meeting 2014. Walnut Creek, CA. <https://goo.gl/6iunz0>

**Stevens, S. L. R.** and K. D. Malmstrom. Complementing Metagenomes with Single Amplified Genomes (Poster Presentation). Raper Symposium 2013. University of Wisconsin - Madison.

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### Publications

Bendall, M. L.\*, **Stevens, S. L. R.\***, Chan, L.-K., Malfatti, S., Schwientek, P., Tremblay, J., ... Malmstrom, R. R. (*submitted to ISMEJ*). Genome-wide selective sweeps and gene-specific sweeps in natural bacterial populations. \*Equal contributors

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### Teaching

McMahon Git Workshop  
*Instructor - July 2015*  
Bioinformatics Basics: Microbial Ecology and Evolution - Illinois Mathematics and Science Academy Intersession  
*Instructor - Jan. 2015*  
Microbiology 304: Biology of Microorganisms Laboratory - University of Wisconsin - Madison  
*Teaching Assistant/Instructor - Fall 2013*

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