

# Dean Sanders

Telephone: 920-207-6953

Email: [dmsanders@wisc.edu](mailto:dmsanders@wisc.edu)

Github: [sandman2127.github.io](https://github.com/sandman2127)

LinkedIn: [Dean's LinkedIn Profile](#)

ResearchGate: [Dean's Research Gate Profile](#)

## Education:

### University of Wisconsin-Madison

Ph.D. Genetics

Madison, WI

Fall 2013 - Fall 2018

### University of Wisconsin-Madison

M.S. Bacteriology

Madison, WI

Fall 2011 - Spring 2013

### University of Wisconsin-Oshkosh

B.S. Microbiology

Oshkosh, WI

Fall 2008 - Spring 2011

## Skills:

### Computational experience:

- **Designed** bioinformatic pipelines for:
  - Structural Variant Analysis
  - Genotype By Sequencing
  - *de novo* genome Assembly
  - ChIP-Seq and Cut & Run analysis
- **Python, Julia and R** for program design and plotting
  - Pandas, Numpy, Multiprocessing, and Cython
  - dplyr, ggplot2
  - Jupyter notebook (Python 3 and R)
- **4 years experience** with distributed & cloud computing
  - AWS
  - Slurm
  - SGE
  - Condor
- **7 years experience** with software dev & reproducibility practices:
  - Versioning control (git)
  - Containerized environments (singularity, docker, conda)
- **10 years experience** in unix command line & file systems
  - Awk and GNU Parallel for scripting, data preprocessing and pipeline automation
  - Regex

**Analytical techniques:** Personal experience with ESI-MS/MS orbi-trap mass spectrometry and HPLC, sample prep and MS/MS data analysis

**General molecular biology:** Site directed mutagenesis, plasmid isolation, molecular cloning, PCR, Sanger sequencing

**Microscopy:** Performed Epifluorescence and Confocal Microscopy

## Experience:

### **Data Scientist II at the University of Wisconsin Bioinformatics Resource Center**

with Dr. Derek Pavelec and Dr. Mark Berres, University of Wisconsin, Madison.  
December 2018 - Present.

- ❖ Independent project leader on all assigned work
- ❖ Project Experience:
  - *de novo* genome assembly
  - Structural and SNP variant analysis
  - ChIP-Seq & Cut & Run analysis
  - Genotype by sequencing

- GWAS and QTL analysis

**Graduate research assistant investigating histone lysine to methionine mutations and loss of DNA methylation during plant development and stress in *Arabidopsis thaliana*** with Dr. Xuehua Zhong, University of Wisconsin, Madison. December 2013 - November 2018.

- ❖ Bioinformatic analysis of mRNA, ChIP, SNP and Bisulfite sequencing data
- ❖ Mass spectrometry sample preparation, injection and customized MS/MS data analysis
- ❖ General molecular cloning, transgenic plant production

**Graduate research assistant examining bacterial biofilm disruption and production of antimicrobial secondary metabolites by the social amoeba *Dictyostelium*** with Dr. Marcin Filutowicz and Dr. Kalin Vetsigian. University of Wisconsin, September 2011- May 2013.

- ❖ Epifluorescence and confocal microscopy
- ❖ C-18 solid phase extraction, HPLC and LC/MS-TOF

#### Communication: Mentoring experience:

- Personally mentored two graduate and four undergraduate students (4/6 now pursuing higher education)

#### Teaching assistantship:

- 3 semesters TA experience (>20 hours lecture time)

#### Python study group leader:

- Member of combee python study group since Fall 2016, leader in Fall 2017

#### Scientific meeting presentations:

- 6 conference presentations, 9 public outreach in the Wisconsin Institute for Discovery

#### Selected Publications:

Islam A, **Sanders D**, Mishra A, Joshi V (2021) Genetic Diversity and Population Structure Analysis of the USDA Olive Germplasm Using Genotyping-By-Sequencing (GBS) *Genes* 12, 1-15

Jiang J, Lui J, **Sanders D**, Wang B, Zhong X (2021) UVR8 interacts with de novo DNA methyltransferase and suppresses DNA methylation in *Arabidopsis*. *Nature Plants*. 7, 184–197

**Sanders D**, Fieweger R, Lu L, Qian S, Dowell J, Denu JM, Zhong X (2017) Histone lysine-to-methionine mutations reduce histone methylation and cause developmental pleiotropy. *Plant Physiology*. 173, 2243-2252.

#### Patent:

Marcin Filutowicz, Katarzyna Dorota Borys, **Dean Sanders** (2014). Dictyostelid amoeba and biocontrol uses thereof. US20140056850 A1. Amoebagone, LLC