

**Elizabeth McDaniel**  
Microbiology PhD Student  
(214)546-9748 | [emcdaniel@wisc.edu](mailto:emcdaniel@wisc.edu) | <https://elizabethmcd.github.io>

## Education

### **University of Wisconsin – Madison**

*Aug. 2016-present*

Ph.D. Candidate in the Microbiology Doctoral Training Program

- Laboratory of Dr. Katherine McMahon, Departments of Bacteriology and Civil and Environmental Engineering

### **University of Arkansas – Fayetteville**

*Aug. 2012-May 2016*

Bachelor of Science in Biology with Statistics Minor, Cum Laude Honors

- GPA: 3.71/4.0
- Honors Thesis: Natural Variation of the Ena1p Sodium Pump in *S. cerevisiae*

## Research Experience

### **Graduate Research Assistant**

*Jan. 2017 - present*

*McMahon Lab – Departments of Bacteriology and Civil & Environmental Engineering, University of Wisconsin – Madison, Madison WI*

- Dissertation project on bacterial communities of engineered wastewater systems using a combination of genome-resolved metagenomics approaches and enrichment culture techniques
- Developed computational workflows and approaches for high-throughput assembly and analysis of freshwater lake metagenomes
- Lab data and systems administrator for storing large-scale metagenomic data and associated metadata

### **Undergraduate Research Assistant**

*Aug. 2013-Aug. 2016*

*Lewis Lab – Department of Biological Sciences, University of Arkansas, Fayetteville, AR*

- Studied the natural variation of stress defense mechanisms in the budding yeast *S. cerevisiae*
- Developed a protocol for creating transposon deletion libraries in wild strains of *S. cerevisiae*

### **Summer Undergraduate Research Intern**

*May-Aug. 2014*

*Broach Lab – Department of Biochemistry and Molecular Biology, Penn State Hershey, Hershey, PA*

- Investigated interactions of chromosomes during quiescence in the budding yeast *S. cerevisiae* using fluorescence microscopy

## Peer-Reviewed Publications

**3. Peterson B.D., McDaniel E.A., Schmidt A.G., Lepak R.F., Tran P.Q., Marick R.A., Ogorek J.M., DeWild J.F., Krabbenhoft D.P., McMahon K.D. Mercury methylation genes identified across diverse anaerobic microbial guilds in a eutrophic sulfate-enriched lake. *Environmental Science and Technology*. Nov. 2020, 54 (24), 15840-15851; DOI: 10.1021/acs.est.0c05435**

**2. McDaniel E.A., Peterson B., Stevens S.L.R., Tran P.Q., Anantharaman K., McMahon K.D.. Expanded Phylogenetic Diversity and Metabolic Flexibility of Mercury-Methylating Microorganisms. *mSystems*. Aug 2020, 5 (4) e00299-20; DOI: 10.1128/mSystems.00299-20**

**1. McDaniel E.A., Stuecker T.N., Veluvolu M., Gasch A.P., Lewis J.A. Independent Mechanisms for Acquired Salt Tolerance versus Growth Resumption Induced by Mild Ethanol Pretreatment in *Saccharomyces cerevisiae*. *mSphere. Editor's Pick*. Nov 2018, 3 (6) e0057418; DOI: 10.1128/mSphere.00574-18**

## Preprints and Submissions

**\* denotes equal contribution**

**1. McDaniel E.A.\*, Moya-Flores F.\*, Keene Beach N., Camejo P.Y., Oyserman B.O., Kizaric M., Khor E.H.,**

Noguera D.N., McMahon K.D. **Metabolic differentiation of co-occurring *Accumulibacter* clades revealed through genome-resolved metatranscriptomics.** *bioRxiv*. Nov. 2020. DOI: 10.1101/2020.11.23.394700. Submitted.

### Software, Teaching Resources, and Editorials (Non Peer-Reviewed)

3. Eren A.M., Kiehl E., Shaiber A., Veseli I., Miller S.E., Schechter M.S., Fink I., Pan J.N., Yousef M., Fogarty E.C., Trigodet F., Watson A.R., Esen O.C., Moore R.M., Clayssen Q., Lee M.D., Kivenson V., Graham E.D., Merrill B.D., Karkman A., Blankenberg D., Eppley J.M., Sjoden A., Scott J.J., Vazquez-Campos X., McKay L.J., **McDaniel E.A.**, Stevens S.L.R., Anderson R., Fuessel J., Fernandez-Guerra A., Maignien L., Delmont T.O., Willis A.D. **Community-led, integrated, reproducible multi-omics with anvi'o.** Jan. 2021, 6, 3-6. *Nature Microbiology (Commentary)*. DOI: 10.1038/s41564-020-00834-3.

2. **McDaniel E.A. Analyzing 16S Amplicon Data in R Course** (version 1.0). July 2020. DOI: 10.5281/zenodo.3942708. Course website at <http://elizabethmcd.github.io/R-amplicons/>.

1. **McDaniel E.A.**, Anantharaman, K., McMahon K.D. **metabolisHMM: Phylogenomic analysis for exploration of microbial phylogenies and metabolic pathways.** Dec. 2019. *bioRxiv*. DOI: 10.1101/2019.12.20.884627.

### Oral Presentations

#### Invited Talks and Plenary Sessions

5. Examining Long-Term Microbial Population Dynamics At Multiple Scales Using Enrichment Bioreactors As Model Ecosystems. **Microbial Ecology and Water Engineering (MEWE) conference.** Hiroshima, Japan. Nov. 2019.

4. Expanded Phylogenetic Diversity and Metabolic Flexibility of Microbial Mercury Methylation. Evolution Series Seminar. **University of Wisconsin – Madison JF Crow Institute for the Study of Evolution.** Sept. 2019.

3. Integrating Anvi'o Tools into your Workflow: Insights from a Biological Nutrient Removal (BNR) System. **Resolving Microbial Communities at Strain-Level Symposium.** Penryn, UK. Aug. 2018.

2. Natural Variation in Yeast Uncovers Novel Regulation of the Ena1p Sodium Pump. **Southeastern Regional Yeast Meeting.** Tuscaloosa, AL. March 2016.

1. Natural Variation in Yeast Uncovers Novel Regulation of the Ena1p Sodium Pump. **Southeastern Regional Yeast Meeting.** Little Rock, AR. March 2015.

#### Internal Seminars

4. Characterizing Ecological Roles and Interactions of '*Candidatus Accumulibacter phosphatis*' in Engineered Bioreactors. **Microbiology Doctoral Training Program.** University of Wisconsin – Madison. Sept. 2020

3. "Best" Practices in Metagenomic Binning and Annotation. **Computational Biology, Ecology, and Evolution (ComBEE) 'Omics Study Group session.** University of Wisconsin – Madison. ComBEE 'Omics Study Group Session. Dec. 2019.

2. Eco-systems biology of a Microbial Community Performing Enhanced Biological Phosphorus Removal. **Microbiology Doctoral Training Program Seminar.** University of Wisconsin – Madison. Nov. 2018.

1. Bacterial Communities of Lab-Scale Wastewater Enrichments. **UW-Madison Bioscience Opportunities Preview Weekend Lightning Talk.** Sept. 2018.

#### Virtual Seminars and Plenary Sessions

2. Characterizing Ecological Roles and Interactions of '*Candidatus Accumulibacter phosphatis*' in Wastewater Systems. **MicroSeminar.** August 2020. Recording available at <https://bit.ly/2Q5NCzb>

1. Applying Genome-Resolved Metagenomics to Elucidate the Microbial “Black Box” in Model Wisconsin Lakes. **Incorporating Data Science and Open Science Techniques in Limnology and Oceanography Virtual Summit**. July 2020. Recording available at <https://bit.ly/2BAOSGC>

### Poster Presentations

7. ComBEE: Computational Biology, Ecology, and Evolution: Enhancing computational literacy in the life sciences through peer-led study groups. Presented at **UW-Madison Data Science Hub Data Science Research Bazaar** and **Wisconsin Institute for Discovery “Illuminating Discovery”** events. Jan. & Feb. 2020. (Poster constructed and presented on behalf of ComBEE Team)

6. Microbiomes of Freshwater Lakes and Engineered Wastewater Systems. **Water @ UW Symposium**. Madison, WI. Oct. 2018. (Poster constructed and presented on behalf of McMahon Lab)

5. Expanded Phylogenetic and Metabolic Diversity of Microbial Mercury Methylation. Presented at **International Society of Microbial Ecology Meeting**. Leipzig, Germany. Aug. 2018., and **Department of Bacteriology Raper Symposium**. Madison, WI. Sept. 2018.

4. Long-Term Population Dynamics of ‘*Candidatus Accumulibacter phosphatis*’ in Enhanced Biological Phosphorus Removal Sequencing-Batch Reactors. **Population, Evolutionary, Quantitative Genetics Conference**. Madison, WI. May 2018.

3. Comparative Genomics of Microbial Methylmercury Production. Presented at **Joint Genome Institute User Meeting: Genomics of Energy and Environment**. San Francisco, CA. Mar. 2018, and **Madison Microbiome Meeting**. Madison, WI. April 2018.

2. Natural Variation in Yeast Uncovers Novel Regulation of the Ena1p Sodium Pump. Presented at **South Central Branch of the American Society for Microbiology Joint Meeting**. Fayetteville, AR. Sept. 2014, **27<sup>th</sup> International Conference on Yeast Genetics and Molecular Biology**. Levico, Terme, Trentino, Italy. Sept. 2015, **Arkansas IDEa Network of Biomedical Research Excellence Meeting**. Fayetteville, AR. Nov. 2015.

1. Chromosome Interactions in Quiescent Yeast. **Penn State Hershey Summer Undergraduate Research Internship Programs Symposium**. Hershey, PA. Aug. 2014.

### Honors and Awards

Civil and Environmental Engineering Becker Travel Supplement Award - \$500	Nov. 2019
O.N. Allen Soil and Environmental Microbiology Small Grants Award - \$3,290	Aug. 2019
Department of Bacteriology Betley-Allen Fellowship Award	May 2019
University of Wisconsin-Madison Student Travel Grant - \$600	April 2019
Microbiology Doctoral Training Program Travel Award - \$1000	May 2018
Department of Bacteriology Travel Award - \$1000	May 2018
Southeastern Regional Yeast Meeting Travel Award - \$250	Mar. 2016
University of Arkansas Honors College Research Grant - \$1200	Jan. 2016
University of Arkansas Honors College Travel Grant - \$1200	Aug. 2015
Southeastern Regional Yeast Meeting Travel Award - \$250	Mar. 2015
University of Arkansas Honors College Research Grant - \$2500	Jan. 2015
University of Arkansas Academic Scholarship - \$2500	2014-2015
ASM South Central Branch Meeting 2 <sup>nd</sup> Place Poster Award	Sept. 2014
University of Arkansas Academic Scholarship - \$1000	2013-2014
University of Arkansas Symphony Orchestra Scholarship - \$1000	2012-2016
University of Arkansas New Arkansan Non-Resident Tuition Award - \$39,040	2012-2016

### Teaching, Mentoring, and Service

**Bioinformatics Workshops (Helper/Teaching Assistant/Instructor/Course Development)**  
**Certified Carpentries Instructor as of June 2019**

Virtual Software Carpentry Workshop; Helper

Aug. 2020

Amplicon Analysis in R Workshop; <i>Instructor and Developer</i>	April 2020
Virtual Software Carpentry Mini-Workshop; <i>Helper</i>	April 2020
Research Bazaar Software Carpentry Workshop; <i>Helper</i>	Jan. 2020
Carpentries Genomics Introduction to R; <i>Instructor</i>	Aug. 2019
Data Carpentry Workshop Introduction to R; <i>Instructor</i>	June 2019
Software Carpentry Workshop; <i>Helper</i>	June 2019
Microbiome & Data Science Hubs Git/Github Pages Workshop; <i>Instructor and Developer</i>	Mar. 2019
Resolving Microbial Communities at Strain-Level Resolution Symposium; <i>Teaching Assistant</i>	Aug. 2018
ComBEE Git/Github Pages Workshop; <i>Instructor and Developer</i>	Sept. 2017
ComBEE Anvi'o Workshop; <i>Teaching Assistant</i>	May 2017
ComBEE Git Workshop; <i>Teaching Assistant</i>	Mar. 2017
ComBEE R Study Group Introduction to R; <i>Instructor and Developer</i>	Sp. 2017

### **Computational Biology, Ecology, and Evolution (ComBEE) Study Group Co-Chair** Jan. 2017-present

ComBEE is a peer-led group for computational biology researchers on the UW-Madison campus. We hold study groups on the R and python programming languages, current topics in genomics, and host seminars from graduate students, postdocs, and professors on topics broadly ranging in ecology and evolution. Website at [combe-uw-madison.github.io](http://combe-uw-madison.github.io).

### **McMahon Lab Mentoring** 2017-present

*Graduate Student Mentor for Undergraduate Students, Research Experience Undergraduate Interns (REUs), and Masters Students carrying out independent projects and/or routine maintenance and sampling*

- *Kaela Amundson*: Characterization and Enrichment of Microorganisms Capable of Extracellular Electron Transfer. *Undergraduate Mentee* Fa. 2017-Sp. 2018.
- *Matthew Wolff*: Metagenomics of Freshwater Lake and Engineered Wastewater Microbial Communities. *Undergraduate Mentee* Sp.-Fa. 2018
- *Kali Denis*: Time-Series Analysis of Under-Ice Freshwater Bacterial Communities. *Undergraduate Mentee* Sp. 2018.
- *William Dunn*: Ecophysiology of *Thermoleophilia*. *Undergraduate Mentee*. Fa 2020.
- *EBPR Team*: Mentored to date **6 undergraduate** students performing routine maintenance and sampling of lab-scale bioreactors simulating Enhanced Biological Phosphorus Removal (EBPR). Additionally mentored **2 masters** students through their thesis projects on EBPR.

<b>O.N. Allen Soil and Environmental Microbiology Small Grants Review Panel</b>	Sp. 2021
<b>Topics in Biotechnology Guest Lecturer on Wastewater Microbial Ecology</b>	Sp. 2020
<b>Microbiology 551: Senior Capstone Guest Lecturer on 16S Workflows and Bioinformatics</b>	Sp. 2020
<b>MEWE Multi-Omics Methods for Water Engineering Workshop Developer and Assistant</b>	Nov. 2019
<b>Microbiology Doctoral Training Program Admissions Committee</b>	Fa. 2019
<b>Data Science Hub Research Bazaar Planning Committee</b>	Fa. 2019
<b>Microbiology 304: Biology of Microorganisms Laboratory Teaching Assistant</b>	Sp. 2018
<b>Ad-hoc Journal Reviewer: ISMEJ (1), mSystems (1), mSphere (1), Env Micro (1)</b>	2017-present
<b>BIOL 2323: General Genetics Drill Instructor</b>	Sp. 2015
<b>University of Arkansas Office of Admissions Ambassador</b>	Aug. 2013-Dec. 2015

### **Professional Development**

<b>Google Career Readiness Data Analyst Challenge</b>	Summer 2020
- Google Cloud Essentials Quest Badge	
- From Data to Insights with Google Cloud Platform Specialization	
<b>DELTA Research Mentor Training</b>	Summer 2019
<b>Carpentries Instructor Training and Certification</b>	April 2019
<b>IMG Microbial Genomics and Metagenomics Workshop, FISABIO</b>	June 2017
<b>Anvi'o Workshop, University of Chicago</b>	April 2017
<b>Data Carpentry Workshop, University of Wisconsin-Madison</b>	Jan. 2017