DREW T. WAGNER

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EDUCATION

→ Doctor of Philosophy, Biochemistry • Fall 2017

University of Texas at Austin Advisor: Adrian Keatinge-Clay

→ Elements of Computing Program: Data Science Track • Fall 2017

University of Texas at Austin

→ Bachelor of Science, Biochemistry and Molecular Biology with Honors • Spring 2010

University of California Santa Cruz

EXPERIENCE

→ Graduate Student • Keatinge-Clay Lab • August 2012 - Present

Department of Molecular Biosciences, University of Texas at Austin

- Active researcher on several completed and ongoing projects focusing on the structural and mechanistic characterization of polyketide synthase domains.
- Developed enzyme assays and mutagenesis screens to characterize substrate tolerance and mechanism of action of biosynthetic enzymes.
- Employed recombinant DNA technology and X-ray crystallography to design and construct custom fusion proteins for synthetic biology applications.

→ Research Technician • June 2011 - August 2012

Molecular Sciences Institute - Berkeley, CA

- Constructed and maintained recombinant *S. cerevisiae* strain collection.
- Performed metabolic engineering experiments on *S. cerevisiae* strains using flow cytometry and fluorescent microscopy leading to development of yeast strains for production of commercially relevant chemicals.
- Statistical analysis of *S. cerevisiae* gene expression data.

→ Research Assistant • Ottemann Lab • June 2010 – June 2011

Department of Microbiology and Environmental Toxicology – University of California Santa Cruz

• Assisted with design and execution of *in vitro* experiments on chemotaxis proteins of the bacterial pathogen *H. pylori*.

TEACHING

- Spring 2017 BCH369 Fundamentals of Biochemistry
- Fall 2016, Spring 2015, Fall 2014, Fall 2013 BCH339F Foundations of Biochemistry
- Spring 2016 BIO361 Human Infectious Diseases
- Fall 2015 BIO326M Intro to Medical Microbiology and Immunology
- Spring 2014 BCH369L Biochemistry Laboratory
- Spring 2011 (UCSC) BIOC100C Integrative Biochemistry of Metabolic Pathways

COMPUTATIONAL SKILLS

- Programming
 Python, R, Bash, git/github
- Database
 MySQL, MongoDB
- Data Visualization PyMol, R studio, ggplot, Tableau
- Web HTML, CSS, Javascript, Node.js, jQuery

PUBLICATIONS

- 1. **Wagner DT**, Zhang Z, Meodod R, Piel P, Keatinge-Clay AT. Structure and Function of a Pyran Synthase Domain from a trans-Acyltransferase Polyketide Synthase. [In Preparation]
- 2. **Wagner DT***, Zeng J*, Bailey CB*, Gay DC, Yuan F, Manion HR, Keatinge-Clay AT. Structural and Functional Trends in Dehydrating Bimodules from *trans*-Acyltransferase Polyketide Synthases. Structure. 2017 Jun.

- Zeng J, Wagner DT, Zhang Z, Moretto L, Addison JD, Keatinge-Clay AT. Portability and Structure of the Four-Helix Bundle Docking Domains of trans-Acyltransferase Modular Polyketide Synthases. ACS Chem Biol. 2016 Sep.
- Wagner DT*, Stevens DC*, Mehaffey MR, Manion HR, Taylor RE, Brodbelt JS, Keatinge-Clay AT. α-Methylation follows condensation in the gephyronic acid modular polyketide synthase. Chem Commun (Camb). 2016 Jul.
- Stevens DC*, Wagner DT*, Manion HR, Alexander BK, Keatinge-Clay AT. Methyltransferases excised from trans-AT polyketide synthases operate on N-acetylcysteamine-bound substrates. J Antibiot (Tokyo). 2016 Jul.
- Gay DC*, Wagner DT*, Meinke JL, Zogzas CE, Gay GR, Keatinge-Clay AT. The LINKS motif zippers transacyltransferase polyketide synthase assembly lines into a biosynthetic megacomplex. J Struct Biol. 2016 Mar
- 7. Fage CD, Isiorho EA, Liu Y, **Wagner DT**, Liu HW, Keatinge-Clay AT. The structure of SpnF, a standalone enzyme that catalyzes [4 + 2] cycloaddition. Nat Chem Biol. 2015 Apr.
- Gay G, Wagner DT, Keatinge-Clay AT, Gay DC. Rapid modification of the pET-28 expression vector for ligation independent cloning using homologous recombination in Saccharomyces cerevisiae. Plasmid. 2014 Nov.
- Hughes AJ, Tibby MR, Wagner DT, Brantley JN, Keatinge-Clay AT. Investigating the reactivities of a
 polyketide synthase module through fluorescent click chemistry. Chem Commun (Camb). 2014 May.
- Zdraljevic S, Wagner D, Cheng K, Ruohonen L, Jäntti J, Penttilä M, Resnekov O, Pesce CG. Single-cell measurements of enzyme levels as a predictive tool for cellular fates during organic acid production. Appl Environ Microbiol. 2013 Dec.

^{* =} denotes shared first authorship