

# Christopher Hamm

*Data Scientist*

37437 State Hwy 16  
Woodland, CA 95695  
USA

**M** +1 (517) 802 8503

**T** +1 (530) 669 6017

**E** [cahamm@ucdavis.edu](mailto:cahamm@ucdavis.edu)

**W** [butterflyology.github.io](https://butterflyology.github.io)

## Education

**PhD**, *Michigan State University, Entomology.* **2012**

**PhD**, *Michigan State University, Ecology, Evolutionary Biology & Behavior.* **2012**

**MS**, *California State University, Fresno, Biology.* **2008**  
*With Distinction*

**BS**, *California State University, Fresno, Biology.* **2004**  
*Magna Cum Laude*

## Experience

### Vocational

**Data Science Training Lead**, *Bayer Crop Sciences.* **2018–Present**

Naveen Singla, supervisor

Achievements:

- Develop, promote, and distribute best practices curriculum for enterprise data scientists
- Organize, coordinate, and teach data fluency workshops
- Develop proprietary curriculum for company specific workflows
- Create and implement strategic vision to improve company-wide data fluency
- Provide *ad hoc* statistical support

**Data Scientist**, *Monsanto Company.* **2017–2018**

Shawn L. Stricklin, supervisor

Achievements:

- Developed and implemented a program to simulate the vegetable breeding pipeline
- Implemented predictive modeling to predict genotype sample submission frequency
- Organized and coordinated Software Carpentry and Instructor Training workshops
- Created gradient boosted regression model to identify disease susceptible genotypes
- Provided *ad hoc* statistical support

**Postdoctoral researcher**, *University of California, Davis.* **2016–2017**

C. Titus Brown, supervisor.

Achievements:

- Oversaw development and deployment of Reproducible Research with R lessons for the Data Carpentry Foundation

**Postdoctoral scholar**, *University of Kansas*.

**2014–2016**

James R. Walters, supervisor.

Achievements:

- Developed tools to analyze large empirical and simulated molecular datasets
- Implemented and interpreted complex statistical modeling on ecological and genomic data, including differential expression analysis
- Formalized novel statistical models to describe bacterial infection frequency while correcting for relatedness

**Postdoctoral researcher**, *University of California, Davis*, **2012–2014**  
Davis, California.

David J. Begun and Michael Turelli, supervisors.

Achievements:

- Coordinated multi-laboratory effort to assess the use of a bacterial infection to control an invasive insect
- Executed comparative genomic analysis of invasive species
- Created bioinformatics pipeline to import, trim, and map genomic data

## Statistical methods

Deep learning in R & Python using Keras and TensorFlow

Linear and mixed-effects models in R

Forecasting with ARIMA and prophet

Hierarchical modeling and inference in R and stan

## Computer skills

**Data analysis:** R, Python, SQL

**Reproducibility:** RMarkdown,  $\text{\LaTeX}$

**Version control:** git, GitHub

**Other:** BASH