

Christopher Hamm

Data Science Training Lead

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Summary

Data Science Training Lead and Lead Data Scientist with a strong research and experimental background with 10+ years experience using reproducible methods and rigorous analysis to generate unique insights and create value. Strong teacher using research-based pedagogy and a Software Carpentry Foundation Trainer.

Experience

2018–Present **Data Science Training Lead, Lead Data Scientist, Bayer - Decision Science.**
Nalini Polavarapu, supervisor.

- **Technical Project Management:**
 - Supervised 30+ machine learning projects
 - Develop, promote, and distribute best practices curriculum for enterprise data scientists
 - Develop proprietary curriculum for company specific workflows
 - Created neural network forecasting models for supply chain
- **Mentorship & Training:**
 - Platinum Pinnacle Award for creating and implementing company-wide digital fluency curriculum
 - Organize, coordinate, mentor, and teach data fluency workshops - have instructed over 1200 learners in 3 years.
 - Coordinate 15 digital fluency trainers
- **Member:**
 - Data Science Center of Excellence
 - Model Governance Team
 - Data Management Team

2017–2018 **Data Scientist, Monsanto Company.**
Shawn L. Stricklin, supervisor.

- **Individual Contributor:**
 - Developed and implemented a program to simulate the vegetable breeding pipeline
 - Implemented predictive modeling to predict genotype sample submission frequency
 - Created gradient boosted regression model to identify disease susceptible genotypes
 - Provided *ad hoc* statistical support to 10 plant breeders
 - Organized and coordinated Software Carpentry and Instructor Training workshops

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

C. Titus Brown, supervisor.

Achievements:

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

2014–2016 **Postdoctoral scholar**, *University of Kansas*.

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

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

Education

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

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

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

With Distinction

2004 **BS**, *California State University, Fresno*, Biology.

Magna Cum Laude

Statistical methods

Deep learning in R & Python using Keras and TensorFlow

Deep learning in python with pytorch and fastai

Machine learning in R & python

Linear and mixed-effects models in R

Forecasting with ARIMA, prophet, and LSTM

Hierarchical Bayesian modeling and inference in R and stan

Computer skills

Data analysis R, Python

Reproducibility docker, knitr, RMarkdown, \LaTeX , jupyter notebooks,

Version git, GitHub, GitLab

control

Cloud AWS ML Pipeline & SageMaker, Applied AI with IBM Watson
computing

Misc. BASH, SQL