Keith Hultman, Ph.D.

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Skills

- Analytic Methods: Experience with linear and logistic regression, k-means and agglomerative clustering, decision trees and random forest, A/B testing, sentiment analysis of text, and time series forecasting
- Data Software: Fluent in R for data analysis and machine learning including implementing R Shiny interactive applications; proficient in creating data visualizations using Tableau and ggplot2; additional experience with Perl, SQL, and Python and AWS cloud computing
- Writing: Experience in technical writing, including 7 peer-reviewed publications, grant writing, and editing for scientific journals
- Presenting: Charismatic and confident public speaker with award-winning experience presenting to technical and lay audiences

Experience

Visiting Assistant Professor, Elmhurst College

Aug 2016 - Jun 2017

- Designed and taught upper level Bioinformatics course focusing on reproducible research methods in genomics using R markdown
- Implemented RStudio Server on AWS Cloud for student lab use
- Developed an interactive R Shiny application to analyze alumni survey

Postdoctoral Research Fellow, Northwestern University Sep 2009 - Aug 2013

- Developed research program involving next-generation sequencing of neural crest transcripts during epithelial-to-mesenchymal transition
- Led a team of five on independent research projects
- Established zebrafish as a model system in the laboratory and managed
 126-tank facility housing up to 7000 fish
- Analyzed 36 GB of RNA sequence data and identified >10 genes uniquely expressed in neural crest cells that were previously unknown
- Automated methods for image acquisition and analysis for phenotypic effects (ImageJ) that reduced microscope usage time by 90%

Adjunct Professor

Sep 2013 - Aug 2016

- Instructor for Introductory Biology, Molecular Genetics, Development, and Human Genetics and Society at Elmhurst College and Loyola University
- Developed and published a hypothesis-driven dog genotyping laboratory

Selected Independent Projects

- (2016) Developed an ensemble predictive model for product sales of bakery goods in Grupo Bimbo Kaggle competition, decreased error rate by 50%
- (2006) Conducted pharmacological screen with >1200 drugs and developed quantification methods for cell migration assay
- (2003) Identified gene duplications using Perl and the BLAST algorithm in genomic sequence

Education

Elmhurst College, Elmhurst, Illinois

May 2017

Masters in Data Science

Washington University School of Medicine, St. Louis, Missouri

May 2009

Ph.D. Molecular Genetics and Genomics

University of Saint Thomas, St. Paul, Minnesota

May 2001

B.A. Biology