## Eric T. Mittman

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Objective

Statistician, data scientist

Education

PhD candidate in Statistics, Iowa State University, expected Winter 2017

GPA: 3.83

M.S. in Statistics, Iowa State University, Ames, Iowa B.A. in Philosophy, Luther College, Decorah, Iowa

Experience

AES Consultant, Iowa State University, Ames, IA March 2016-present

- Provide statistical support for graduate students and post-docs in various areas of research
- Help clients select and interpret appropriate displays and analyses for their data
- Write and edit R and SAS code

Research Assistant, Iowa State University, Ames, IA May 2014-present

- Research and implement methods for statistical analysis of high-dimensional RNA-seq data
- Write code in R, C++ and CUDA
- Present research to faculty and students and at conferences

Teaching Assistant, Iowa State University, Ames IA

August 2013-May 2014

- Instructed large introductory Statistics course
- Wrote and administered midterm and final exams
- Recorded and tabulated grades on Blackboard

**Secondary Math Instructor**, Lake Mills Community School August 2009 - May 2013

- Taught high school courses in algebra, trigonometry, precalculus and computer science
- Monitored student progress and communicated important information to students, parents and administrators
- Wrote lesson plans and assessments with aim of improving student acheivement

Computer Skills Languages: R, C++, Python, SAS, Latex

Software/Tools: RStudio, Emacs, Git, SQLite, Microsoft Office

Publications

Niemi, J., Mittman, E., Landau, W., & Nettleton, D. (2015). Empirical Bayes analysis of rna-seq data for detection of gene expression heterosis. Journal of agricultural, biological, and environmental statistics, 20(4), 614-628.

**Presentations** 

Mittman, E., Niemi, J. (2016, August). Dirichlet process mixture model for high-dimensional gene expression data, *Contributed talk*, Section on Statistics in Genomics and Genetics, Joint Statistical Meetings, Chicago, IL.

Mittman, E., Niemi, J. (2016, May). Semiparametric Bayesian inference for high-dimensional gene expression experiments, *Poster*, Annual Conference on Applied Statistics in Agriculture, Manhattan, KS.