William Michael Landau

Education

- PhD Statistics, Iowa State U, 2016
- MS Statistics, Iowa State U 2013
- BS Math, U of Chicago 2011

Contact

- will.landau@gmail.com
- wlandau.github.io
- linkedin.com/in/wlandau
- github.com/wlandau

Publications

- Landau, W., Niemi, J., and Nettleton, D., "Fully Bayesian analysis of RNA-seq counts for the detection of gene expression heterosis", Accepted to the *Journal of the American Statistical Association* on April 28, 2018.
- Landau, W. (2018), "The drake R package: a pipeline toolkit for reproducibility and high-performance computing". *Journal of Open Source Software*, 3(21), 550, https://doi.org/10.21105/joss.00550.
- Niemi, J., Mittman, E., Landau, W., and Nettleton, D. (2015), "Empirical Bayes Analysis of RNA-seq Data for Detection of Gene Expression Heterosis," *Journal of Agricultural*, *Biological, and Environmental Statistics*, 20, 1-15. Available at link.springer.com.
- Landau, W. and Liu, P. (2013), "Dispersion Estimation and Its Effect on Test Performance in RNA-seq Data Analysis: A Simulation-Based Comparison of Methods," *PLOS One*, 8.
 Available at journals.plos.org.
- Ratliff, B., Womack. C., Tang, X., Landau, W., Butler, L., and Szpunar, D. (2010), "Modeling the Rovibrationally Excited C2H4OH Radicals from the Photodissociation of 2-Bromoethanol at 193 nm," *Journal of Physical Chemistry*, 114, 4934-4945. Available at ncbi.nlm.nih.gov.

Articles under review

 Landau, W. and Niemi, J., "A fully Bayesian strategy for high-dimensional hierarchical modeling using massively parallel computing," submitted to the *Journal of Computational* and *Graphical Statistics* on March 8, 2016. Preprint available at arxiv.org.

Open Source Software

- drake, an R-focused workflow manager for reproducibility and high-performance computing. Part of rOpenSci.
- downsize, and R package to toggle between the test and production versions of large workflows.
- R packages fbseq, fbseqCUDA, and fbseqOpenMP from dissertation research.
- R utilities eply, grapes, and wildcard.

Awards

- 2017 Lilly Innovator Award. Awarded for leading a successful team effort to modernize Lilly's internal process for contributing open source software.
- Student Paper Award, American Statistical Association Section on Statistical Computing, Jan 2016. Awarded for an early draft of the preprint at arxiv.org/abs/1606.06659.
- Vince Sposito Statistical Computing Award, Iowa State University, Aug 2013.
- GlaxoSmithKline Industrial Scholarship, Iowa State University, Sep 2011.
- Alumni Scholarship, Iowa State University, Aug 2011.

Skills

- Reproducible research, hierarchical models, Bayesian methods, Markov chain Monte Carlo, statistical computing, high-dimensional data analysis, genomics data analysis, exploratory analysis, visualization, linear and nonlinear models, data mining, machine learning, predictive modeling, multivariate analysis.
- High-performance computing, R, R package development, general-purpose graphics processing unit (GPU) computing, CUDA, shell scripting, LaTeX, HTML, CSS.
- Past experience with C/C++, MPI, OpenMP, Python, JavaScript, AWK, Fortran.

Research statistician

- October 2016 Present
- Eli Lilly and Company
- Helped design and plan early- and mid-phase clinical trials as the lead statistician for three new autoimmune therapies.
- Supported late-phase clinical trial teams with advanced analytics, including clinical program simulation and tailored therapeutics.
- Developed interactive R Shiny web applications to facilitate decision-making and simulation.
- Created the drake package to enhance reproducibility and high-performance computing in R.

Research assistant

- May 2013 Aug 2016
- RNA-sequencing Working Group, Department of Statistics, Iowa State University.
- Funded by NIH grant R01GM109458 with Drs. Dan Nettleton and Jarad Niemi.
- Developed a new fully Bayesian analysis method for high-dimensional genomic datasets using hierarchical models.
- Implemented massively parallelized Markov chain Monte Carlo.
- Created the

fbseq

R package to distribute the analysis method.

- Implemented and distributed parallel computing backends for CUDA GPUs (fbseqCUDA) and OpenMP (fbseqOpenMP).
- · Created the

remakeGenerator,

parallelRemake, and

downsize packages to manage, ameliorate, expedite, and accelerate computationally heavy reproducible workflows that are under heavy development.

Seminar instructor

- Aug Dec, 2012 and 2013.
- Department of Statistics, Iowa State University.
- GPU computing seminar series at wlandau.github.io/gpu.
- Educated faculty and graduate students on massively parallel computing with generalpurpose graphics processing units.
- Constructed, curated, and distributed slides, video, and example code at wlandau.github.io/gpu and on YouTube.

Course instructor

- Jan May, 2012 and 2013.
- Department of Statistics, Iowa State University.
- STAT 305: Engineering Statistics (wlandau.github.io/stat305).

Grader

- Aug Dec, 2011.
- Department of Statistics, Iowa State University.
- STAT 231: Engineering Probability.
- STAT 105: Introduction to Engineering Statistics.

Leadership at Eli Lilly and Company

- Led a successful team effort to modernize Lilly's internal procedure for contributing open source software.
- Served as a volunteer moderator in the 2017 National Science Bowl (high school Indiana Regionals).

Leadership at Iowa State University

- Founder and leader, Cloud Computing Working Group, Sep Dec 2015.
- Member, Computation Advisory Committee, Sep 2015 May 2016.
- Volunteer instructor, Office of Precollegiate Programs for Talented and Gifted (OPPTAG), Mar 13, 2014.
- Fellow, Preparing Future Faculty, Aug 2013 May 2014.
- Assistant Coach, Boxing Club, Aug 2013 Dec 2013.

References

- Jarad Niemi, PhD advisor and major professor, niemi@iastate.edu.
- Dan Nettleton, lead principal investigator of the RNA-sequencing Working Group (Iowa State Department of Statistics), dnett@iastate.edu.
- Peng Liu, MS advisor and major professor, pliu@iastate.edu.
- Additional references available on request.

Hobbies

· Climbing, Brazillian Jiu Jitsu, sailing, windsurfing