### 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". *Journal of the American Statistical Association*, https://doi.org/10.1080/01621459.2018.1497496.
- 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.

## Open Source Software

- drake, an R-focused pipeline toolkit for reproducible computation and high-performance computing. Part of rOpenSci.
- txtq, a minimalist, serverless, socketless message queue for interprocess communication.
- downsize, and R package to toggle between the test and production versions of large workflows.
- R packages fbseq, fbseqCUDA, and fbseqOpenMP. A toolkit for the fully Bayesian analysis
  of genomic count data.

### **Awards**

- 2019 NumFOCUS New Contributor Recognition. Awarded for inclusive and collaborative work with rOpenSci. https://numfocus.org/blog/2019-numfocus-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, statistical computing, hierarchical models, Bayesian methods, Markov chain Monte Carlo, 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
- Developed internal statistical tools and capabilities for the design, simulation, and analysis of clinical trials.
- Served as the lead statistician in early-phase autoimmune asset teams.
- Supported late-phase clinical trial teams with advanced analytics, including clinical program simulation and tailored therapeutics.
- Published open-source software packages drake and txtq to improve 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