DANIEL R. KICK, PHD

- · Experience with statistics, programming, machine learning, and deep learning.
- · Presented to scientific and general audiences over 25 times since 2016.
- · Led 4 teaching assistants and mentored 7 research students.
- · Developed statistical tool used by >700 students as of 2021.

┛ PROFESSIONAL AND RESEARCH EXPERIENCE

Present 2021

Research Geneticist

Jacob Washburn Lab, USDA-ARS

- · Employed deep neural networks, machine learning models, and best linear unbiased predictors to improve corn yield prediction accuracy in diverse environments (see Kick et al., 2023).
- · Designed project plan and submitted grant application titled "Environmentally Aware Deep Learning Based Genomic Selection And Management Optimization For Maize Yield" to the NIFA Agriculture and Food Research Initiative's Education and Workforce Development Program (Decision Pending).
- · Communicated with stakeholders through 8 presentations (4 national, 2 regional, and 2 outreach presentations).
- · Mentored 2 students conducting a high throughput root phenotyping experiment and wrote scripts for data organization and analysis.
- · Created and taught a Python data visualization workshop titled "Tools and Techniques for a Jupyter Based Scientific Workflow".
- · Collaborated with plant biologists, contributing statistical expertise (manuscript in preparation).
- · Completed Software Carpentries instructor certification, taught R for Reproducible Scientific Analysis, and assisted in teaching Data Management with SQL.
- · Designed and completed a professional development curriculum with the guidance of an industry scientist via the Bayer-University Mentoring Program.
- · Served as a panel member on "Next-Generation Omics" at the 2022 University of Missouri Division of Biological Sciences Retreat.

2021 2015

Graduate Researcher

David Schulz Lab, University of Missouri

- · Author on 6 publications: 4 original research and 2 eLife Insight publications.
- · Assessed the efficacy of machine learning models to identify cell identity from mRNA and contig abundances. Applied cluster estimation, hyperparameter tuning, unsupervised machine learning, and supervised machine learning. Identified and learned needed skills primarily through self study. Collaborated with molecular biology project lead. (see Northcutt¹, Kick¹, et al. 2019).
- · Demonstrated activity desynchronization induces changes in neuronal connections. Defined research question and experiments. Developed novel approach to quantify changes in cell activity (see Kick and Schulz 2022).
- · Investigated activity dependent changes in neuronal excitability, conductances, and ion channel mRNA abundances. Designed experiments, collected data, performed analysis, and developed novel method for quantifying changes in cell activity using in silico simulations.
- · Collaborated with electrophysiologists, assisting with data analysis.
- · Collaborated with computational neuroscientists, contributing biological and statistical expertise (in preparation).
- · Mentored 5 students and oversaw their projects.
- · Communicated results through 17 presentations (6 national, 6 regional, and 5 outreach and recruitment).
- · Served as a peer mentor of 3 PhD students in use of R for reproducible data analysis, created internal documents on same.

CONTACT INFO

- in daniel-kick-5a449b9a
- **8** Google Scholar
- github.com/danielkick
- **6** danielkick.com

EDUCATION PhD: Biological Sciences

University of Missouri, Columbia, MO (2021)

Machine Learning Methods for Biomedical Informatics. Ouantitative Methods in the Life Sciences, and Grant Writing

Bachelor of Science: Biology

Truman State University, Kirksville, MO (2015)

Next Generation Sequence Data and Analysis, Bioinformatics

Leadership role in the biological honors society Tri-Beta

Technical Skills

- **Q**: R Programming (6 years) experience with tidyverse, lme4, caret, ggplot2, shiny, & package creation.
- **?**: Python Programming (2) years) experience with pandas, numpy, plotly, scikit-learn, keras, pytorch.
- ☑: Miscellaneous Experience with high performance computing (bash, slurm), virtual environments (conda, singularity, docker), version control (git, GitHub), literate programming (Rmarkdown, Jupyter).

For a pdf with links scan here



2021	•	Lead Teaching Assistant, Animal Physiology Lab
 2020		Biological Sciences, University of Missouri
		 Developed statistics web application used by more than 700 students as of 2021 with shiny (source, deployed) for data visualization, testing assumptions, and fitting frequentist, non-parametric, and Bayesian models. Led 4 Teaching Assistants and coordinated adaptation of lab curriculum to be fully online due to COVID-19 pandemic. Mentored next Lead Teaching Assistant, created documentation on best practices.
2020	•	Teaching Assistant, Animal Physiology Lab
 2018		Biological Sciences, University of Missouri
		 Updated and refined curriculum, delivered lectures and ensured experiments were conducted safely, and modeled student grade distributions to identify and adjust for differences in grading.
2018	•	Curriculum Consultant, Animal Physiology Lab
		Biological Sciences, University of Missouri
		Redesigned course material to incorporate primary literature and data analysis.
2016	•	Teaching Assistant, Animal Physiology Lab
 2015		Biological Sciences, University of Missouri
		· Delivered weekly lectures, ensured experiments were conducted safely, provided timely feedback on assignments.
2013	•	Undergradute Researcher
 2015		University of Missouri, University of Connecticut, and Truman State University
		 Designed a hydroponic system for maize root phenotyping – Diane Janick-Buckner and Brent Buckner, Truman State University (2014-2015), Quantified retinal minor splicisome expression using immunohistochemistry – (NSF REU) Rahul Kanadia, University of Connecticut (2014), Measured effectiveness of oligonucleotide treatment for spinal muscular atrophy in mice – (NSF REU) Christian Lorson, University of Missouri (2013).
	Q	HONORS AND AWARDS (3/4)
2019	•	J. Perry Gustafson Award for Outstanding Graduate Research in the Life Sciences
		This award is granted for the quality of a student's independent research and academic achievements. Recipients receive a \$2,000 award.
2018	•	National Institutes of Health T ₃₂ Training Grant Recipient
 2016		This fellowship provides a \$27,000 yearly stipend and travel awards of \$750.
2015	•	Cum Laude & President's Recognition, Truman State University
	<u>Д</u>	SELECTED PUBLICATIONS (3/7 PUBLISHED, 2 IN REVIEW, 2 IN PREP.)
2023	•	Ensemble of BLUP, Machine Learning, and Deep Learning Models Predict Maize Yield Better than Each Model Alone
		<i>Daniel R. Kick</i> , Jacob D. Washburn (In Review) $bioR\chi iv$
2023	•	Yield Prediction Through Integration of Genetic, Environment, and Management Data Through Deep Learning
		Daniel R. Kick, Jason G. Wallace, James C. Schnable, Judith M. Kolkman, Baris Alaca, Timothy M. Beissinger, David Ertl, Sherry Flint-Garcia, Joseph L. Gage, Candice N. Hirsch, Joseph E. Knoll, Natalia de Leon, Dayane C. Lima, Danilo Moreta, Maninder P. Singh, Teclemariam Weldekidan, Jacob D. Washburn G3: Genes, Genomes, Genetics
2022		Timing dependent potentiation and depression of electrical synapses contributes to network stability in the crustacean cardiac ganglion
		Daniel R. Kick and David J. Schulz The Journal of Neuroscience
2019	•	Molecular profiling of single neurons of known identity in two ganglia from the crab Cancer borealis
	1	Adam J. Northcutt ¹ , Daniel R. Kick ¹ , Adriane G. Otopalik, Benjamin M. Goetz, Rayna M. Harris, Joseph M. Santin, Hans A. Hofmann, Eve Marder, and David J. Schulz (¹ denotes co-first authorship) Proceedings of the National Academy of Sciences