DANIEL R. KICK, PHD

- · Experience with statistics, programming, machine learning, and deep learning.
- · Designed NIFA funded research plan (Grant 2023-67012-39485).
- Presented to scientific and general audiences 34 times since 2016.
- · Invited presentations at University of Michigan, Truman State University, and Iowa State University.
- · Developed statistical tool used by >700 students as of 2021.
- · Data and models downloaded 410 times and viewed over 1105 times.
- · Led 4 teaching assistants and mentored 7 research students.

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PROFESSIONAL AND RESEARCH EXPERIENCE

Present | 2021

Research Geneticist

Jacob Washburn Lab, USDA-ARS

- Secured \$225,000 to develop "Environmentally Aware Deep Learning Based Genomic Selection And Management Optimization For Maize Yield" from the National Institute of Food and Agriculture (NIFA) (Grant 2023-67012-39485).
- Employed deep neural networks, machine learning models, and best linear unbiased predictors to improve corn yield prediction accuracy in diverse environments.
- Communicated with stakeholders via 19 presentations (6 national, 9 regional, 4 outreach).
- · Mentored 2 students conducting a high throughput root phenotyping experiment.
- Created and taught a Python data visualization workshop titled "Tools and Techniques for a Jupyter Based Scientific Workflow".
- · 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).
- Defined research question and experiments. Developed novel approaches to quantify changes in cell activity.
- · Collaborated with electrophysiologists, assisting with data analysis.
- · Collaborated with computational neuroscientists, contributing domain expertise.
- · Mentored 5 students and oversaw their projects.
- · Communicated results through 18 presentations (6 national, 6 regional, 6 outreach).
- Served as a **peer mentor of 3 PhD students** in use of R for reproducible data analysis, created internal documents on same.

Lead Teaching Assistant, Animal Physiology Lab

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.

CONTACT INFO

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

EDUCATION PhD: Biological Sciences

University of Missouri, Columbia, MO (2021)

Machine Learning Methods for Biomedical Informatics, Quantitative 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

- **R**: R Programming (7 years) and experience with tidyverse, lme4, caret, ggplot2, shiny, & package creation.
- e: Python Programming (3 years) and experience with pandas, numpy, plotly, scikit-learn, keras, pytorch.
- ☑: Miscellaneous Experience with high performance computing (bash, slurm), virtual environments (conda, singularity), version control (git, GitHub), literate programming (Rmarkdown, Jupyter), crop growth modeling (APSIM Next Generation).

For a pdf with links scan here.





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
2015		· 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 –(2014-2015), Quantified retinal minor splicisome expression using immunohistochemistry – (2014), Measured effectiveness of oligonucleotide treatment for spinal muscular atrophy in mice – (2013).
	Q	HONORS AND AWARDS (4/5)
2025		NIFA Fellowship (AFRI EWD)
 2023		\$225,000 awarded (Grant # 2023-67012-39485) over two years to create and environmentally aware deep learning genomic selection models and prepare the recipient to transition into industry.
2019		J. Perry Gustafson Award for Outstanding Graduate Research in the Life Sciences
2019	Ĭ	This \$2,000 award is granted for the quality of their research and academic achievements.
2018		NIH T ₃₂ Training Grant Recipient
2016 2016		This provides a \$27,000 yearly stipend and \$750 yearly to facilitate presenting research at scientific conferences.
2015	•	Cum Laude & President's Recognition, Truman State University
	Д	SELECTED PUBLICATIONS (SHOWING 4/9 PUBLISHED, 0/2 IN PREP.)
2023		Ensemble of Best Linear Unbiased Predictor, Machine Learning, and Deep Learning Models Predict Maize Yield Better Than Each Model Alone
		Daniel R. Kick, Jacob D. Washburn in Silico Plants
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. Kick¹ , 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