# DANIEL R. KICK, PHD

- · Quantitative biologist with 7 years of experience in academic and government laboratories.
- Experience with statistical modeling, machine learning, deep learning, and data visualization.
- · Experience with leadership, management, mentorship, and teaching.
- · Developed and deployed an educational statistical tool used by >700 students as of 2021.

## **-**

## ฝ PROFESSIONAL AND RESEARCH EXPERIENCE

Present | 2021

#### Research Geneticist

Jacob Washburn Lab, USDA-ARS

- Evaluated deep neural networks, machine learning models, and best linear unbiased predictors to improve yield prediction accuracy in diverse environments.
- Communicated with stakeholders through 7 presentations (3 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. Grace Sidberry (2021-pres.), Madi Michell (2022-pres.)
- Created and taught a Python data visualization workshop at University of Missouri Bioinformatics in Plant Science group 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

#### Graduate Researcher

David Schulz Lab, University of Missouri

- · Author on 6 publications: 4 original research and 2 eLife Insight publications.
- Communicated results with stakeholders through 17 presentations (6 national, 6 regional, and 5 outreach and recruitment).
- Collaborated with electrophysiologists studying spinal cord injury. Assisted with data organization, cleaning, and analysis (manuscript in preparation).
- Collaborated with computational neurosciencists, contributing biological and statistical expertise (in preparation).
- Mentored 5 students and oversaw their projects. Abby Beckerdite (2016-2019), Ayla Ross (2019), Katlyn Sullivan (2018), Kelly Hiersche (2017), & Rody Kingston (2016)
- $\cdot$  Served as a peer mentor for 3 PhD students in use of R for reproducible data analysis, created internal guides to share knowledge on same.
- · Assessed the efficacy of machine learning models to recapitulate neural cell identity from mRNA and contig abundances. Applied cluster estimation, hyperparameter tuning, unsuprevised machine learning, and supervised machine learning techniques. Identified theoretical and practical resources, then learned needed knowledge and skills primarily through self study. Collaborated with molecular biology project lead (Adam Northcutt).
- Demonstrated that activity desynchronization induces degree dependent changes in conductance between neurons. Defined research question and approach, designed and implemented experiments using current clamp, voltage clamp, dynamic clamp, and pharmaceutical application. Developed novel approach to quantify changes in cell activity. Applied resampling techniques, mixed effects models, and akaike information criteria, and asympotic regression.
- Investigated activity dependent changes from elevated depolarization in neuronal excitability, conductances, and ion channel mRNA abundances in small neural networks. Designed and executed experiments, collected data, performed analysis, developed novel method for quantifying changes in cell activity using *in silico* simulations.

#### CONTACT INFO.

■ hello@danielkick.com

Ø danielkick.com

github.com/danielkick

in daniel-kick-5a449b9a

**9** Google Scholar

# 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, Analysis of Variance and Experimental Design, Non-Parametric Statistics, and Economic & Medicinal Botany

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.

#### 

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  Biological Sciences, University of Missouri
2020	<ul> <li>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.</li> <li>Lead 3-4 Teaching Assistants and coordinated adaptation of lab curriculum to be fully online due to Covid-19.</li> <li>Mentored successor, created documentation on best practices.</li> </ul>
2020	Teaching Assistant, Animal Physiology Lab
 2018	Biological Sciences, University of Missouri
	<ul> <li>Analyzed student grade distributions to adjust for grader effects at request of the instructor of record.</li> <li>Continued duties below.</li> </ul>
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
	· Provided 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
	<ul> <li>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)</li> </ul>
	HONORS AND AWARDS (2/5)
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 T32 Training Grant Recipient
2016 <b>Д</b>	This fellowship provides a \$27,000 yearly stipend and travel awards of \$750.
	SELECTED PUBLICATIONS (3/7, 5 REVIEWED)
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
	Adam J. Northcutt <sup>1</sup> , <i>Daniel R. Kick</i> <sup>1</sup> , Adriane G. Otopalik, Benjamin M. Goetz, Rayna M. Harris, Joseph M. Santin, Hans A. Hofmann, Eve Marder, and David J. Schulz ( <sup>1</sup> <i>denotes co-first authorship</i> ) Proceedings of the National Academy of Sciences