## DANIEL R. KICK, PHD

- Experience predicting maize yield across environments using deep learning, machine learning, statistical modeling to facilitate crop improvement.
- Designed funded research project (USDA NIFA) to create environmentally aware deep learning models for trait prediction in row crops (\$225,000, Grant 2023-67012-39485).
- · Initiated and led a multi-institutional research collaboration; collaborated with domain experts.
- Presented to scientific and general audiences 36 times since 2016.
- · Invited presentations at University of Michigan, Truman State University, Iowa State University, and University of Georgia's Al in Plant Breeding Symposium.
- · Developed data analysis app used by >700 students as of 2021.
- · Data and models downloaded over 775 times and viewed over 1565 times.
- · Led 4 teaching assistants and mentored 9 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).
- Initiated and led a multi-institutional research collaboration and wrote custom libraries for same
- Employed deep neural networks, machine learning models, best linear unbiased predictors, and scientific models (i.e. process based) to improve trait prediction accuracy in diverse environments.
- Communicated with stakeholders via 21 presentations (7 national, 10 regional, 4 outreach).
- Organized and led deep learning community attended by students, university professors, and government scientists.
- 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.
- Mentored and advised 4 students (including through the 2024 pilot of the USDA Postdoc Mentorship Program) with projects involving high throughput phenotyping, a genome wide association study (GWAS), and representing spatial effects.
- · Served as a panel member on "Next-Generation Omics" at the 2022 University of Missouri Division of Biological Sciences Retreat.
- Designed and completed a professional development curriculum with the guidance of an industry scientists via the Bayer-University Mentoring Program and the Maize Genetics Mentoring Program.

### 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 predict cell identity from mRNA and contig abundances. Applied cluster estimation, hyperparameter tuning, unsupervised machine learning, and supervised machine learning.
- Defined research question and experiments. Developed novel methodolgies and approaches to address research questions.
- · Collaborated with molecular biology project lead, providing data analysis and statistical knowledge. (see *Northcutt<sup>1</sup>, Kick<sup>1</sup>, et al. 2019*).
- · Collaborated with biologists and physiologists, assisting with data analysis.
- Served as a **peer mentor of 3 PhD students** in use of R for reproducible data analysis, created internal documents on same.
- Mentored 5 students and provided oversight and advice on their projects and analysis.
- · 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.

### CONTACT INFO

- → hello@danielkick.com
- in daniel-kick-5a449b9a
- **9** Google Scholar
- github.com/DanielKick-USDA
- github.com/danielkick
- **6** 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**

- **\(\rightarrow\):** Python (3 years) experience with pandas, numpy, plotly, scikitlearn, keras, pytorch.
- **R**: R (8 years) experience with tidyverse, lme4, caret, ggplot2, shiny, & package creation.
- E: Miscellaneous Experience with high performance computing (bash, slurm), virtual environments (conda, singularity), version control (git, GitHub), literate programming (Quarto, Rmarkdown, Jupyter, nbdev), Scientific modeling (crop growth modeling APSIM Next Generation), basic SQL (SQLite, PostgreSQL).

For a pdf with links scan here.



2021	Lead Teaching Assistant, Animal Physiology Lab
 2020	Biological Sciences, University of Missouri
	<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>Led 4 Teaching Assistants and coordinated adaptation of lab curriculum to be fully online due to COVID-19 pandemic.</li> <li>Mentored next Lead Teaching Assistant, created documentation on best practices.</li> </ul>
2020	Teaching Assistant, Animal Physiology Lab
 2018	Biological Sciences, University of Missouri
	· Iteratively refined data analysis components of the curriculum, modeled student grade distributions to identify and adjust for differences in grading.
2018	Curriculum Consultant, Animal Physiology Lab
	Biological Sciences, University of Missouri
	· Redesigned and updated course material to incorporate data analysis and primary literature.
2016	Teaching Assistant, Animal Physiology Lab
 2015	Biological Sciences, University of Missouri
2013	· Delivered weekly lectures, ensured experiments were conducted safely, provided timely feedback on assignments.
2015	Undergradute Researcher
2013	University of Missouri, University of Connecticut, and Truman State University
2013	<ul> <li>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).</li> </ul>
•	HONORS AND AWARDS (SHOWING 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
	This \$2,000 award is granted for the quality of their research and academic achievements.
2018	NIH T32 Training Grant Recipient
 2016	This provides a \$27,000 yearly stipend and \$750 yearly to facilitate presenting research at scientific conferences.

Cum Laude & President's Recognition, Truman State University



2015

2023

2023

2019

## SELECTED PUBLICATIONS (SHOWING 3/9 PUBLISHED, 5 IN PREP. OR REVIEW)

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

### 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

### Molecular profiling of single neurons of known identity in two ganglia from the crab Cancer borealis

Adam J. Northcutt<sup>1</sup>, **Daniel R. Kick**<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> denotes co-first authorship) Proceedings of the National Academy of Sciences