Kelly L. Sovacool

sovacool.dev

in kelly-sovacool

kelly-sovacool

Education

2018-present **PhD Bioinformatics**, *University of Michigan*,

Dept. of Computational Medicine and Bioinformatics

2014-2018 **BS Biology**, *University of Kentucky*,

Dept. of Biology, Minor: Computer Science

Research Experience

2019-present Graduate Student Researcher, Schloss Lab,

Dept. of Microbiology and Immunology, University of Michigan

- o Developing bioinformatics pipelines & software for microbial ecology.
- Analyzing metagenomic and 16S rRNA amplicon sequence data.
- Applying machine learning methods to microbiome-based classification problems.
- 2018-2019 **Rotation Student Researcher**, *Program in Biomedical Sciences*, University of Michigan
- 2015-2018 Undergraduate Lab Assistant, Moseley Bioinformatics Lab,

Dept. of Molecular and Cellular Biochemistry, University of Kentucky

 Developed a computational tool for identifying sets of orthologous and paralogous gene products in whole genomes to facilitate collinearity analysis and detection of gene duplication events.

2016-2018 BIO395 Independent Research Student, Weisrock Lab,

Dept. of Biology, University of Kentucky

- Developed scripts and a SNP calling pipeline for amplicon sequence data.
- Population structure analysis of the *Ambystoma tigrinum* species complex.
- Bayesian species delimitation of the *Desmognathus fuscus* species complex.
- 2015-2016 Undergraduate Lab Assistant, Jaromczyk Lab,

Dept. of Computer Science, University of Kentucky

- o Maintained the Epichloë festucae genome project database.
- Analyzed RNA-seq data of *Chenopodium quinoa* and coffee ringspot virus.

Awards

Grants & Fellowships

2020 Conference Travel Grant, Rackham Graduate School, (\$800)

University of Michigan

2019-2021 NIH T32 Bioinformatics Training Program Fellow

University of Michigan

Dec 2017 **Oswald Research & Creativity Award**, *UK Office of Undergraduate Research*, 2nd place in the Biological Sciences category (\$200) University of Kentucky

May-Aug Summer Research Grant, UK Office of Undergraduate Research, (\$2,000)

2017 University of Kentucky

2014-2018 **Presidential Scholarship**, (out-of-state full tuition)

University of Kentucky

Honors

May 2018 Graduated Cum Laude with Departmental Honors in Biology University of Kentucky

May 2018 Biology Undergraduate Research Award Nominee

University of Kentucky

2014-2018 Lewis Honors College

University of Kentucky

Service

2021-present CoderSpaces co-host, U-M ISR Data Science Hub

Host office hours at a weekly virtual help session for data science practitioners.

2018-present **Executive Committee Member: Open Source Technical Lead**, Girls Who Code at UM-DCMB

- Plan, apply for funding, develop curriculum, and maintain resources to teach introductory Python programming & data science to young women.
- Facilitate collaborative development and maintenance of our teaching resources.
- Organize our year-round Club for high school women.
- Organize our annual Data Science Summer Experience for high school women in the Detroit area.

2019-present **Organizer and Maintainer**, *U-M Software Carpentry*

- Co-lead development & maintenance of a custom curriculum for workshops teaching programming skills for reproducible research.
- o Maintain the website and organize workshops.

2019-present Graduate Student Coordinator, U-M Data Analysis Networking Group

- Organize monthly meetings & a one-day symposium for researchers to sharpen their data analysis skills.
- Funded by a Rackham Interdisciplinary Workshop grant.

2009-present Live Sound Engineer, for various churches and non-profit organizations.

- Set up, maintain, repair, and operate front of house, monitor, and recording systems during sound checks, rehearsals, services, and concerts.
- o Train new sound techs in the art and science of live sound.

2021-present **Peer review for scientific journals**, *PLOS ONE (2)*

Teaching Experience

2019-present Facilitator & Capstone Project Mentor, Girls Who Code at UM-DCMB

- o 12-22 July 2021. Virtual Data Science Summer Experience.
- o Aug 2020 May 2021. Weekly virtual Club for high schoolers.
- o 06-16 July 2020. Virtual Data Science Summer Experience.
- o Aug 2019 May 2020. Weekly Club for high schoolers.
- o 15-19 July 2019. Data Science Summer Experience in Detroit, MI.
- o Jan-May 2019. Weekly Club for high schoolers.

2018-present **Software Carpentry instructor**, *U-M Carpentries*, umcarpentries.org

- o 11-12 Jan 2021. Lead instructor for U-M WISE-sponsored virtual workshop.
- o 06-07 Jan 2020. Lead instructor for U-M WISE-sponsored workshop.
- o 01-02 July 2019. Workshop instructor.
- o 22-23 May 2019. Workshop helper.
- o 01 Mar 2019. Workshop helper.
- o 17-18 Dec 2018. Workshop helper.
- 25 Apr 2019 **DNA Day Ambassador**, *MI DNA Day*, Epigenetics & scientific journeys, Ann Arbor, MI
- 20 Mar 2019 **Workshop helper**, *Graduate Society of Black Engineers and Scientists*, Data visualization with Python workshop, University of Michigan
- 16 Mar 2019 Capstone Activity Leader, Females Excelling More in Math, Engineering, & Science, Binary numbers through Ozobots with GWC at UM-DCMB, University of Michigan
 - 2012-2018 **Tutor**, *freelance*, for high school and college students in Biology, Calculus, Chemistry, Computer Science, and Bioinformatics.

Presentations

- Jun 2020 **ASM Microbe**, OptiFit: a fast method for fitting amplicon sequences to existing OTUs, Poster, Cancelled due to COVID-19.
- Apr 2018 **Showcase for Undergraduate Scholars**, Developing a Global Homology Analysis for Comparative Genomics, Poster, University of Kentucky
- Apr 2018 **Systems Biology & Omics Integration Seminar**, Developing a Global Homology Analysis for Comparative Genomics, Talk, University of Kentucky
- Apr 2018 National Conference on Undergraduate Research, Developing a Global Homology Analysis for Comparative Genomics, Poster, University of Central Oklahoma
- Apr 2016 **Showcase for Undergraduate Scholars**, *Processing RNA-seq Reads of Plants Infected with the Coffee Ringspot Virus*, Poster, University of Kentucky
- Apr 2016 **UT-KBRIN Bioinformatics Summit**, Processing RNA-seq Reads of Plants Infected with the Coffee Ringspot Virus, Poster, Cadiz, KY

Apr 2015 **Showcase for Undergraduate Scholars**, *The Effect of Meditation on Performance*, Poster, University of Kentucky

Continuing Education

- 27-30 Jan Building Tidy Tools workshop at rstudio::conf, RStudio, PBC
 - 2020 San Francisco, CA
- 17-19 Dec Winter School in Research Software Engineering, US Research Software
 - 2019 Sustainability Institute
 Seattle, WA
- 03-04 Jun Software Carpentry Instructor Training, The Carpentries
 - 2019 University of Michigan
- 01-05 May PyCon Education Summit & Conference, Python Sofware Foundation
 - 2019 Cleveland, OH
- 24-26 Apr MICROBIOL 612.2, Riffomonas minimalR workshop
 - 2019 University of Michigan

Software Contributions

- mikropml User-Friendly R Package for Supervised Machine Learning Pipelines.

 Co-author and maintainer. https://github.com/SchlossLab/mikropml
 - mothuR R package to work with mothur output files.

 Co-author and maintainer. https://github.com/SchlossLab/mothuR
 - mothur Command-Line Tool for Processing 16S rRNA Gene Sequence Data.

 Contributor. https://github.com/mothur/mothur

Publications

- B. D. Topçuoğlu, Z. Lapp, K. L. Sovacool, E. Snitkin, J. Wiens, and P. D. Schloss. Mikropml: User-Friendly R Package for Supervised Machine Learning Pipelines. *JOSS*, 6(61):3073, May 2021.
- K. M. Everson, L. N. Gray, A. G. Jones, N. M. Lawrence, M. E. Foley, K. L. Sovacool, J. D. Kratovil, S. Hotaling, P. M. Hime, A. Storfer, G. Parra-Olea, R. Percino-Daniel, X. Aguilar-Miguel, E. M. O'Neill, L. Zambrano, H. B. Shaffer, and D. W. Weisrock. Geography is more important than life history in the recent diversification of the tiger salamander complex. *PNAS*, 118(17), Apr. 2021.
- A. K. Hagan, N. A. Lesniak, M. J. Balunas, L. Bishop, W. L. Close, M. D. Doherty, A. G. Elmore, K. J. Flynn, G. D. Hannigan, C. C. Koumpouras, M. L. Jenior, A. J. Kozik, K. McBride, S. B. Rifkin, J. M. A. Stough, K. L. Sovacool, M. A. Sze, S. Tomkovich, B. D. Topcuoglu, and P. D. Schloss.

Ten simple rules to increase computational skills among biologists with Code Clubs. *PLOS Computational Biology*, 16(8):e1008119, Aug. 2020.

M. M. Goodin, M. Farman, H. Inocencio, C. Jang, J. W. Jaromczyk, N. Moore, and K. Sovacool. Processing RNA-Seq data of plants infected with coffee ringspot virus. In *Proceedings of the 15th Annual UT-KBRIN Bioinformatics Summit 2016: Cadiz, KY, USA. 8-10 April 2016.* BMC Bioinformatics, Aug. 2016.