Kelly Sovacool

PhD Candidate

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Education

2018-present PhD Bioinformatics, Dept. of Computational Medicine and Bioinformatics, University of Michigan.

o Advisor: Patrick D. Schloss

2014-2018 BS Biology, Dept. of Biology, University of Kentucky.

Minor: Computer Science

Skills

Languages & R, Python, Bash, C++, Snakemake, R Markdown, Quarto, LaTeX, conda,

Tools git, GitHub, SLURM.

Research gut microbiome composition, amplicon sequence analysis, metagenomics, su-

pervised machine learning, data visualization, reproducible manuscripts.

R package maintenance, continuous integration, high performance computing.

Research Experience

2019-present Graduate Student Researcher, Schloss Lab, Dept. of Microbiology and *Immunology*, University of Michigan.

- O Develop and benchmark bioinformatics pipelines and software for microbial ecol-
- O Analyze 16S rRNA gene amplicon sequence data.
- O Apply machine learning methods to gut microbiome classification problems in colorectal cancer and C. difficile infection.
- O Collaborate with other scientists on microbiome projects and mentor junior lab

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.

> O Developed a computational tool in Python 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.
 - O Developed bash scripts and a SNP calling pipeline in Snakemake.
 - O Population structure analysis of the Ambystoma tigrinum species complex.
 - O 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.
 - O Analyzed RNA-seq data of Chenopodium quinoa and coffee ringspot virus.

Teaching Experience

- Jan-Apr 2023 **Graduate Student Instructor**, *Dept. of Computational Medicine & Bioinformatics*, University of Michigan.
 - O BIOINF 576: Tool Development for Bioinformatics
 - 2019-2022 **Facilitator & Capstone Project Mentor**, *Girls Who Code at U-M DCMB*, University of Michigan.
 - Weekly Club during the school year and week-long Summer Experience for high schoolers to learn Python for data science
- 2018-present Workshop Instructor & Helper, U-M Carpentries, University of Michigan.
 - 2-day Software Carpentry workshops teaching computational skills for reproducible research
 - Jun 2022 Instructor, Virtual.
 - Intro to R & RNA-Seq Workshop for ASM Microbe conference attendees
 - Apr 2019 **DNA Day Ambassador**, *Michigan DNA Day*, Pioneer High School, Ann Arbor, MI.
 - Epigenetics & Scientific Journeys
 - Mar 2019 Workshop helper, Graduate Society of Black Engineers and Scientists, University of Michigan.
 - Data Visualization with Python Workshop
 - Mar 2019 Capstone Activity Leader, Females Excelling More in Math, Engineering, & the Sciences, University of Michigan.
 - Binary Numbers through Ozobots with GWC at U-M DCMB
 - 2012-2018 Tutor, freelance.
 - for high school and college students in Biology, Calculus, Chemistry, Computer Science, and Bioinformatics.

Service

2019-present Executive Committee Member: Open Source Technical Lead, Girls Who Code at U-M DCMB.

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

2019-present Organizer & Maintainer, U-M Carpentries.

- O Co-lead development & maintenance of a curriculum for workshops teaching programming skills for reproducible research.
- O Maintain the website, develop curriculum, and organize workshops.
- Collaborate with U-M Women in Science and Engineering to organize workshops for women+.

2021-present Mentor, Schloss Lab.

O Mentor an undergraduate student in building reproducible machine learning models to predict C. difficile infection severity from gut microbiome composition.

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

O Hold office hours at a weekly virtual help session for data science practitioners

2021-present Peer reviewer.

o PLOS ONE (1)

2019-2021 Graduate Student Coordinator, U-M Data Analysis Networking Group.

- Organize monthly meetings & a one-day symposium for researchers to sharpen their data analysis skills.
- O Apply for funding through a Rackham Interdisciplinary Workshop grant.

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

Open Source Contributions

Software

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

Co-author and maintainer

schtools, Schloss Lab tools for reproducible microbiome research (R package).

Co-author and maintainer

mikropml snakemake workflow, Template for running mikropml with Snakemake.

Co-author and maintainer

mothur, Command-Line Tool for Processing 16S rRNA Gene Sequence Data.

Contributor

mothur snakemake workflow, Snakemake template for amplicon sequence analysis with mothur.

Co-author

Curricula

U-M Software Carpentry, Intro to R- the Unix shell- and git for workshops on reproducible research.

O Co-author and maintainer

Girls Who Code at U-M DCMB, Intro to Python for Data Science for Girls Who Code clubs..

Co-author and maintainer

Code Clubs, Short coding tutorials for lab meetings.

Contributor

U-M DANG!, repro-packs: Organizing projects for reproducibility and headache prevention.

Author

Intro to R & RNA-seq, Workshop for 2022 ASM Microbe attendees.

Contributor

Presentations

Talks

- Nov 2022 **Bioinformatics Student Research Hour**, *Predicting _ C. difficile_ infection severity from the taxonomic composition of the gut microbiome*, University of Michigan.
- Feb 2022 **Seminar for the KG Jebsen Center for Genetic Epidemiology**, *Intro to git & GitHub*, (Virtual) Norwegian University of Science and Technology.
- Mar 2021 Bioinformatics Student Research Hour, OptiFit: a fast method for fitting amplicon sequences to existing OTUs, (Virtual) University of Michigan.
- Apr 2018 Systems Biology and Omics Integration Seminar, Developing a Global Homology Analysis for Comparative Genomics, University of Kentucky.

Posters

- Jun 2022 **ASM Microbe**, *Predicting the severity of _ C. difficile_ infections from the taxonomic composition of the gut microbiome*, Washington, DC.
- Jun 2020 **ASM Microbe**, OptiFit: a fast method for fitting amplicon sequences to existing OTUs, Virtual.
- Apr 2018 Showcase for Undergraduate Scholars, Developing a Global Homology Analysis for Comparative Genomics, University of Kentucky.
- Apr 2018 National Conference on Undergraduate Research, Developing a Global Homology Analysis for Comparative Genomics, University of Central Oklahoma.
- Apr 2016 **Showcase for Undergraduate Scholars**, *Processing RNA-seq Reads of Plants Infected with the Coffee Ringspot Virus*, University of Kentucky.
- Apr 2016 UT-KBRIN Bioinformatics Summit, Processing RNA-seq Reads of Plants Infected with the Coffee Ringspot Virus, Cadiz, KY.
- Apr 2015 **Showcase for Undergraduate Scholars**, *The Effect of Meditation on Performance*, University of Kentucky.

Awards

- Grants and Fellowships
- 2022 **Conference Travel Grant**, \$900, Rackham Graduate School
 sity of Michigan.
- 2020-2021 Rackham Interdisciplinary Workshop Grant, \$500, Rackham Graduate School
 School str> University of Michigan.
 - 2020 **Conference Travel Grant**, \$800, Rackham Graduate School
 sity of Michigan.
- 2019-2020 Rackham Interdisciplinary Workshop Grant, \$500, Rackham Graduate School
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- 2019-2021 NIH T32 Bioinformatics Training Program Fellow, Bioinformatics Graduate Program
 Vniversity of Michigan.

- 2017 Oswald Research & Creativity Award
 2nd place in Biological Sciences, \$200, Office of Undergraduate Research
 br> University of Kentucky.
- 2017 **Summer Research Grant**, \$2,000, Office of Undergraduate Research
 University of Kentucky.
- 2014-2018 **Presidential Scholarship**, *out-of-state full tuition*, University of Kentucky.

Honors

- 2018 Graduated Cum Laude with Departmental Honors in Biology, University of Kentucky.
- 2018 Biology Undergraduate Research Award Nominee, University of Kentucky.
- 2014-2018 Lewis Honors College, University of Kentucky.

Continuing Education

- Jan 2020 **Building Tidy Tools workshop at rstudio::conf**, *Rstudio*, *PBC*, San Francisco, CA.
- Dec 2019 Winter School in Research Software Engineering, US Research Software Sustainability Institute, Seattle, WA.
- Jun 2019 **Software Carpentry Instructor Training**, *The Carpentries*, University of Michigan.
- May 2019 **PyCon Education Summit & Conference**, *The Python Software Foundation*, Cleveland, OH.
- Apr 2019 MICROBIOL 612.2, Riffomonas Minimal R Workshop, University of Michigan.

Publications

^{*} Indicates co-first author

- Goodin, M. M., Farman, M., Inocencio, H., Jang, C., Jaromczyk, J. W., Moore, N., & Sovacool, K. L. (2016, August). Processing RNA-Seq data of plants infected with coffee ringspot virus. *Proceedings of the 15th Annual UT-KBRIN Bioinformatics Summit 2016: Cadiz, KY, USA. 8-10 April 2016.* https://doi.org/10.1186/s12859-016-1154-y
- 2. Hagan, A. K., Lesniak, N. A., Balunas, M. J., Bishop, L., Close, W. L., Doherty, M. D., Elmore, A. G., Flynn, K. J., Hannigan, G. D., Koumpouras, C. C., Jenior, M. L., Kozik, A. J., McBride, K., Rifkin, S. B., Stough, J. M. A., Sovacool, K. L., Sze, M. A., Tomkovich, S., Topcuoglu, B. D., & Schloss, P. D. (2020). Ten simple rules to increase computational skills among biologists with Code Clubs. *PLoS Comput Biol*, 16(8), e1008119. https://doi.org/10.1371/journal.pcbi.1008119
- 3. Everson, K. M., Gray, L. N., Jones, A. G., Lawrence, N. M., Foley, M. E., Sovacool, K. L., Kratovil, J. D., Hotaling, S., Hime, P. M., Storfer, A., Parra-Olea, G., Percino-Daniel, R., Aguilar-Miguel, X., O'Neill, E. M., Zambrano, L., Shaffer, H. B., & Weisrock, D. W. (2021). Geography is more important than life history in the recent diversification of the tiger salamander complex. *PNAS*, 118(17). https://doi.org/10.1073/pnas.2014719118
- 4. *Topçuolu, B. D., *Lapp, Z., *Sovacool, K. L., Snitkin, E., Wiens, J., & Schloss, P. D. (2021). Mikropml: User-Friendly R Package for Supervised Machine Learning Pipelines. *JOSS*, 6(61), 3073. https://doi.org/10.21105/joss.03073
- *Duda, M., *Sovacool, K. L., Farzaneh, N., Nguyen, V. K., Haynes, S. E., Falk, H., Furman, K. L., Walker, L. A., Diao, R., Oneka, M., Drotos, A. C., Woloshin, A., Dotson, G. A., Kriebel, A., Meng, L., Thiede, S. N., Lapp, Z., & Wolford, B. N. (2021). Teaching Python for Data Science: Collaborative development of a modular & interactive curriculum. JOSE, 4(46), 138. https://doi.org/10.21105/jose.00138
 *Lapp, Z., *Sovacool, K. L., Lesniak, N., King, D., Barnier, C., Flickinger, M., Krüger,
- *Lapp, Z., *Sovacool, K. L., Lesniak, N., King, D., Barnier, C., Flickinger, M., Krüger, J., Armour, C. R., Lapp, M. M., Tallant, J., Diao, R., Oneka, M., Tomkovich, S., Anderson, J. M., Lucas, S. K., & Schloss, P. D. (2022). Developing and deploying an integrated workshop curriculum teaching computational skills for reproducible research. JOSE. https://doi.org/10.21105/jose.00144
- 7. Sovacool, K. L., Westcott, S. L., Mumphrey, M. B., Dotson, G. A., & Schloss, P. D. (2022). OptiFit: An Improved Method for Fitting Amplicon Sequences to Existing OTUs. mSphere. https://doi.org/10.1128/msphere.00916-21 1752 N St., N.W., Washington, DC
- 8. Barron, M. R., Sovacool, K. L., Abernathy-Close, L., Vendrov, K. C., Standke, A. K., Bergin, I. L., Schloss, P. D., & Young, V. B. (2022). Intestinal Inflammation Reversibly Alters the Microbiota to Drive Susceptibility to Clostridioides difficile Colonization in a Mouse Model of Colitis. *mBio*, 0(0), e01904–22. https://doi.org/10.1128/mbio.01904–22
- 9. Armour, C. R., Sovacool, K. L., Close, W. L., Topçuolu, B. D., Wiens, J., & Schloss, P. D. (2022). Streamlined implementation of a machine learning model to classify screen relevant neoplasia using reference-based OTU clustering. *bioRxiv*. https://doi.org/10.1101/2022.09.01.506299