

Erin H. Wilson

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Recent CS PhD graduate with 10+ years of computational analysis experience (industry, academia) ▪ Creative scientist who can connect ideas across computing, life sciences, and sustainable technology ▪ Programming skills in machine learning, software development, and data visualization ▪ Effective communicator who can adapt explanations of complex topics to diverse audiences ▪ **Seeking a data scientist position where I can be part of solutions for sustainability and climate health**

Keywords: *data science, machine learning, microbiology, science communication, data visualization, predictive modeling*
Research Tools: *Python (PyTorch, pandas, sci-kit learn, numpy, Altair, seaborn, matplotlib), Tableau, AWS, Git, SQL*

Education

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| Ph.D. Computer Science, University of Washington, Seattle, WA | 2023 |
| ▪ NSF GRFP Fellow | |
| M.S. Computer Science, University of Washington, Seattle, WA | 2019 |
| B.A. Computational Biology, Carleton College, Northfield, MN | 2014 |
| ▪ <i>Magna Cum Laude</i> ; awarded Distinction on senior thesis | |

Work & Research Experience [[Academia](#) || [Industry](#)]

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|--|----------------------|
| Graduate Researcher, University of Washington /Supervisors: <i>Dr. David Beck, Dr. Mary Lidstrom</i> | 2017 - 2023 |
| ▪ Built computational frameworks to accelerate genetic engineering efforts in methane-consuming bacteria, a promising carbon removal platform; established a suite of new methanotroph promoter tools | |
| ▪ Characterized the effectiveness of machine learning models for discovering influential genetic patterns from RNA-seq experiments in microorganisms with limited data | |
| Visiting Researcher, DTU Biosustainability Institute, Denmark /Supervisor: <i>Dr. Lars Nielsen</i> | Apr-Aug 2022 |
| ▪ Used machine learning methods (ICA) to identify independent gene modules in methane-consuming bacteria | |
| Data Science Intern, Zymergen /Mentor: <i>Trent Hauck</i> | Summer 2018 |
| ▪ Prototyped deep learning models for predicting the presence of DNA features in microbe genomes | |
| Associate Scientist, Scientific Computing, Amyris /Mentors: <i>Dr. Amoolya Singh, Dr. Darren Platt</i> | 2014 - 2017 |
| ▪ Genotype Specification Language (GSL): Implemented software features and trained biologists/external collaborators to use open-source DNA design language invented at Amyris. (co-authored: article , textbook chapter , blog post , poster) | |
| ▪ Database Development: Developed software pipeline and database schema to translate metabolic pathway designs into DNA build instructions for DARPA-funded project to produce sustainable materials (" Automated Scientist ") | |
| ▪ Bioinformatics: Helped maintain automated whole genome sequencing pipeline and supported scientists in interpreting mutation and coverage data for engineered genomes. | |
| ▪ Communication: Facilitated technical communication between biologists and software engineers; held weekly office hours for 1x1 sequencing and GSL support; gave technical presentations to Amyris R&D (~60 people), Strain Engineering (~30 people), and Automation & Computing groups (~30 people); engaged with attendees at Bay Area science nights about Amyris' sustainable technology | |
| Intern, Scientific Computing, Amyris /Mentor: <i>Dr. Amoolya Singh</i> | December 2013 |
| ▪ Implemented data visualization tool to overlay experimental data on yeast metabolic pathway | |
| Research Assistant, Computational Biology, University of Minnesota /PI: <i>Dr. Chad Myers</i> | Summer 2013 |
| ▪ Analyzed genetic and chemical-genetic interaction data to predict gene targets for chemical perturbants and coded target prediction pipeline | |
| Research Abroad, Coastal Marine Ecology, University of Queensland /PI: <i>Dr. Annie Bosacker</i> | "Winter" 2013 |
| ▪ Conducted observational field research in various Australian ecosystems (reef island, rainforest, desert gorge, intertidal zones) | |
| Research Assistant, Evolutionary Computing, Carleton College /PI: <i>Dr. Sherri Goings</i> | Summer 2012 |
| ▪ Executed experiments with populations of mutating digital organisms to examine the effects of limited CPU resources on the populations' ability to evolve complex Boolean logic functions | |
| Research Assistant, Genetics, University of California, San Francisco /PI: <i>Dr. Nadav Ahituv</i> | Summer 2011 |
| ▪ Performed chromatin immunoprecipitation sequencing experiments on mouse limb tissue to find DNA elements involved in limb patterning and development | |

Awards & Fellowships

Scan Design Foundation Fellowship	2022
▪ Support for research and cultural exchange between Danish and American students	
NSF Graduate Research Fellow	2019
▪ Research funding from the National Science Foundation	
Marilyn Fries Fellowship	2017
▪ Awarded first year graduate research funding in Computer Science	
Clare Boothe Luce Scholarship for Women in Physics and Computer Science	2012
▪ Received funding for summer research in Evolutionary Computing	

Science Communication

Scientific Outreach & Tutorials

- “Modeling DNA Sequences with PyTorch.” (2022) [Tutorial](#) in Towards Data Science, a Medium publication.
- “The Light Side of Genetic Engineering.” (2019) [Article](#) in OneZero, a Medium publication.
- “Genetic Constructor and GSL - The Best of Both worlds.” (2016) [Blog post](#) with Autodesk Bionano Research.

Selected Publications & Presentations

- L. He, J. D. Groom, **E. H. Wilson**, J. Fernandez, M. C. Konopka, D. A. C. Beck, M. E. Lidstrom. (2023) “A methanotrophic bacterium to enable methane removal for climate mitigation.” *PNAS* ([link](#))
- **E. H. Wilson**., M. E. Lidstrom, D. A. C. Beck. (2023) “Probing the limits of deep learning methods for predicting gene expression in non-model microbes.” Rapid talk and poster at SBFC. Portland, OR
- A. H. Singh, B. B. Kaufmann-Malaga, J. A. Lerman, D. P. Dougherty, Y. Zhang, A. L. Kilbo, **E. H. Wilson**, C. Y. Ng, O. Erbilgin, K. A. Curran, C. D. Reeves, J. E. Hung, S. Mantovani, Z. A. King, M. J. Ayson, J. R. Denery, C. Lu, P. Norton, C. Tran, D. M. Platt, J. R. Cherry, S. S. Chandran, A. L. Meadows. (2023) “An Automated Scientist to Design and Optimize Microbial Strains for the Industrial Production of Small Molecules.” *bioRxiv*. ([link](#))
- **E. H. Wilson**, M. E. Lidstrom, and D. A. C. Beck. (2021) “A multi-task learning approach to enhance sustainable biomolecule production in engineered microorganisms.” Tackling Climate Change with Machine Learning, *ICML workshop*. ([link](#), [recording](#))
- **E. H. Wilson** et al. (2021) “A Computational Framework for Identifying Promoter Sequences in Nonmodel Organisms Using RNA-seq Data Sets.” *ACS Synthetic Biology*. ([link](#), [project page](#))
- **E. H. Wilson** (2020) “Using microorganisms to solve macro problems: untangling the genetic circuitry of methane-eating bacteria.” Invited talks at MIDAS Data Science Symposium, University of Michigan and Virtual Women’s Research Day, University of Washington ([recording](#)).
- **E. H. Wilson**, C. Macklin, and D. Platt. (2018) “Engineering genomes with Genotype Specification Language.” In *Methods in Molecular Biology, Synthetic Biology*. Springer Publishing Company, New York, NY. In Press. ([link](#))
- **E. H. Wilson** et al. (2016) “Genotype Specification Language.” *ACS Synthetic Biology*. ([link](#))

Leadership, Volunteering, & Activities

- Research mentor for an undergraduate student (2020-2023)
- Pre-application Review Service mentor (2021-2022)
 - Provided early application feedback to support prospective PhD students from diverse backgrounds
- Peer mentor (2018-2022) and Peer Mentorship Program Organizer (2019-2020) for incoming PhD students
- New Grad Orientation organizer (2018)
- Youth Outreach at “UW Engineering Discovery Days” and “Introduce a Girl to Coding, Robotics, and Data Science”
 - Developed interactive activity “Programming Organisms with DNA Puzzles!” to teach elementary/middle schoolers about metabolic engineering
- Recreational data visualization: “Mistborn: The Final Eyebrow.” (2021) [Article](#) in Towards Data Science.
- Wildlife rescue hospital volunteer, [PAWS](#) (Seattle, WA) and [Wildcare](#) (San Rafael, CA) (2015 - 2019)
 - Treated and cared for injured songbirds in hospital; Co-led youth nature hikes with Education Department
- [MeadoWatch](#) field data collector, UW Biology citizen science project (2019 - present)
 - Collect wildflower blooming data in Mount Rainier National Park
- Carleton Varsity Athletics (Div III)
 - Women’s Soccer (4 seasons), Women’s Tennis (1 season)
 - Received MIAC (Minnesota) Academic All-Conference honors (2011, 2012, 2013)