**Jarrett D. Phillips, Ph.D.**

**School of Computer Science**

**Department of Integrative Biology**

**University of Guelph**

 [jphill01@uoguelph.ca](mailto:jphill01@uoguelph.ca)

phillipsjarrett1@gmail.com



 [jphill01](https://github.com/jphill01) [Jarrett D. Phillips](https://scholar.google.ca/citations?user=2YE-Y4cAAAAJ&hl=en)

 [Jarrett D. Phillips](https://ca.linkedin.com/in/jarrettdphillips) [Jarrett Phillips](https://orcid.org/0000-0001-8390-386X)

 [Jarrett Phillips](https://www.researchgate.net/profile/Jarrett-Phillips)

 [Dr. Jarrett D. Phillips](https://twitter.com/JarrettDPhilli1)

\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

**EDUCATION**

**PhD.** in Computational Sciences**,** University of Guelph 2016-2022

**Co-Advisors**: Dr. Daniel Gillis and Dr. Robert Hanner

**Advisory Committee Members**: Dr. Deborah Stacey and Dr. Graham Taylor

**Thesis**: A Novel Statistical Framework for Assessment of Intraspecific Haplotype Sampling Completeness: Implications for DNA Barcode Gap Estimation

**Master of Bioinformatics (MBinf.)**,University of Guelph 2013-2014

**Co-Advisors**: Dr. Robert Hanner and Dr. Daniel Ashlock

**Thesis**: Assessing DNA Barcode Haplotype Sampling Diversity in the Ray-finned Fishes (Chordata: Actinopterygii)

**BSc. (Hons.)** in Biological Science,University of Guelph 2009-2013

**RESEARCH EXPERIENCE**

**Postdoctoral Fellow** 2023

University of Guelph

Gillis Lab, School of Computer Science

Hanner Lab, Department of Integrative Biology

* Developed a Bayesian hierarchical binary logistic time-series regression model of seafood fraud in the Canadian supply chain
* Mentored and supervised an undergraduate STAT\*4600 student and a University Research Assistant (URA) working to develop models in R, Stan, and Shiny

**Postdoctoral Fellow** 2022

University of Guelph

Hanner Lab, Department of Integrative Biology

Supervisor: Dr. Robert Hanner

* Mentored and supervised a Master of Bioinformatics (MBINF.) BINF\*6999 student on research project
* Participated in conceptualization and drafting of various manuscripts and invited book chapters

**Summer Research Assistant** 2016

Algoma University

Plant and Soil Ecology Lab, Department of Biology

Invasive Species Research Institute (ISRI)

Supervisor: Dr. Pedro Antunes

* Offered bioinformatics and statistical analysis support in R
* Assisted Principal Investigator and undergraduate thesis student with initial drafting of a manuscript on invasive plant root lesion quantification

**Lab Assistant Volunteer**               2014-2016

Algoma University

Plant and Soil Ecology Lab, Department of Biology

Invasive Species Research Institute (ISRI)

Supervisor: Dr. Pedro Antunes

* Offered bioinformatics and statistical analysis support in R
* Assisted with collaborative and outreach initiatives for the Terrestrial Invasive Plant Species (TIPS) Network Project through drafting communication letters to public and private conservation agencies across Canada and the USA seeking volunteers to collect invasive plant species for root lesion quantification

**Undergraduate Research Assistant** 2013 University of Guelph

Vaccine Discovery Research Group

Supervisor: Dr. Mario Monteiro

* Performed various experimental techniques (gas chromatography-mass spectrometry (GC-MS) and Nuclear Magnetic Resonance (NMR)) on bacterial polysaccharide samples for vaccine synthesis and development under the supervision of qualified graduate students

**TEACHING EXPERIENCE**

In addition to co-teaching three undergraduate courses, I was directly involved in grading assignments, creating course and lab content, as well as materials for midterms and final exams.

**Graduate Teaching Assistant (GTA)** 2016-2020

University of Guelph

**CIS\*3130 – System Modelling and Simulation** 2020

~ 30 students ⋅ Python

**CIS\*1910 – Discrete Structures in Computing I** 2017

~ 300 students (~ 40 students per section)

**CIS\*2460 – Modelling of Computer Systems** 2016-2019

~ 60 students ⋅ R, Excel, Java

**STUDENT SUPERVISION AND MENTORSHIP**

I have either directly or indirectly supervised and mentored 14 undergraduate students in the School of Computer Science and the Department of Integrative Biology, all of whom are, or will be, coauthors on my publications. My primary role has been to guide students through the entire scientific process (*i.e.*, formulating research questions, designing, and setting up experiments to test hypotheses, collecting data and analyzing results and writing the manuscript drafts), among other responsibilities, such as providing feedback on students’ draft reports. Students completed research both as part of a course and as lab volunteers.

**Zaid Al-Gayyali** (with Dan Gillis) 2023

University of Guelph

Summer Undergraduate Research Assistant (URA) ⋅ Bayesian modelling of seafood fraud in the Canadian supply chain

**Fynn De Vuono-Fraser** (with Dan Gillis) 2023

University of Guelph

STAT\*4600 ⋅ Bayesian modelling of seafood fraud in the Canadian supply chain

**Amina Asif** (with Bob Hanner) 2022

University of Guelph

BINF\*6999 ⋅ DNA barcode gap analysis of Canadian disease vectors and agricultural pests

**Navdeep Singh** (with Dan Gillis) 2021

University of Guelph

CIS\*4900 ⋅ HACSim RShiny web application

**Maya Persram** (with Bob Hanner) 2020-present

University of Guelph

Hanner Lab volunteer ⋅ R reporting ecological meta-analysis

**Ashley Chen** (with Bob Hanner) 2020-present

University of Guelph

Hanner Lab volunteer ⋅ R reporting ecological meta-analysis

**Olivia Friesen Kroeker** (with Bob Hanner)2020-present

University of Guelph

Hanner Lab volunteer ⋅ R reporting ecological meta-analysis

**Scarlett Bootsma** (with Dan Gillis) 2020-2021

University of Guelph

CIS\*4900/4910 ⋅ HACSim simulation study

**Christina Fragel** (with Bob Hanner)2018**-**2019

University of Guelph

BINF\*6999 ⋅ DNA barcode sequence classification with machine learning

**Jiaojia (Paula) Yu** (with Bob Hanner) 2018**-**2019

University of Guelph

BINF\*6999 ⋅ MDMAPR R Shiny app

**Danielle St. Jean** (with Dan Gillis) 2018**-**2019

University of Guelph

MSc. thesis⋅ DNA barcode sequence classification with machine learning

**Steven French** (with Dan Gillis) 2018

University of Guelph

CIS\*4900/4910 ⋅ HACSim R package

**Julia Harvie** (with Bob Hanner) 2018-2019

University of Guelph

MCB\*4500/4510 ⋅ Data mining GenBank and BOLD

**Ankita Bhanderi** (with Bob Hanner) 2018

University of Guelph

BINF\*6999 ⋅ Data mining GenBank and BOLD

**ASSISTANTSHIPS, AWARDS, SCHOLARSHIPS AND GRANTS**

**Graduate Teaching Assistantships** 2017-2020

University of Guelph$34,506.00-35,148.00 CAD

**Graduate Research Assistantships** 2017-2019

University of Guelph $11,000.00 CAD

**CPES Graduate Dean’s Scholarship** 2017

University of Guelph $3500.00 CAD

* Awarded in recognition of achieving over 85% in Master’s coursework

**CPES Graduate Excellence Entrance (GEE) Scholarship**  2016

University of Guelph $30000.00 CAD

* Awarded in recognition of achieving over 85% in Master’s coursework

**Arthur D. Latornell Graduate Travel Grant**  2019

University of Guelph $500.00 CAD

* Awarded for first-class academic standing in Ph.D. coursework
* Supported travel to the 8th International Barcode of Life Conference in Trondheim, Norway to present work related to resource management and conservation

**SoCS Travel Grant** 2019

University of Guelph $1000.00 CAD

* Supported travel to the 8th International Barcode of Life Conference in Trondheim, Norway to present doctoral research

**Guelph Institute for Environmental Research Small Grants Program (GIER SGP)** 2020

University of Guelph $15000 CAD (not funded)

* 1-yeat postdoctoral funding to develop a Bayesian hierarchical binary logistic time-series regression model of seafood fraud in the Canadian supply chain

**NSERC Postdoctoral Fellowship** 2021

University of Waterloo $90000 CAD (not funded)

* 2-year postdoctoral funding to develop an ensemble machine learning model for taxonomic classification of regulated species in Canada

**Food from Thought Advancing Research Impact (ARIF) Fund** 2022University of Guelph $30000.00 CAD

* 1-year postdoctoral funding to develop a Bayesian hierarchical binary logistic time-series regression model of seafood fraud in the Canadian supply chain

**ACADEMIC SERVICE**

**School of Computer Science (SoCS) Search Committee**  2018

University of Guelph

* Associate Professor position in cybersecurity

**School of Computer Science (SoCS) Search Committee** 2017-2018

University of Guelph

* Two-year contractually-limited Assistant Professor position in cybersecurity

**ACADEMIC PEER REVIEW SERVICE**

I have served as a reviewer for 17 manuscripts in 8 different journals, both alone or under the guidance of my PhD. supervisors.

*Ecology and Evolution* (2)2021-2022

*F1000 Research* (1) 2022

*Frontiers in Ecology and Evolution* (2) 2022

*Lifestyle Genomics* (2) 2022

*Mitochondrial DNA Part* *A* (2) 2023

*Molecular Ecology Resources* (6)2019-2022, 2023

*Molecular Biology Reports* (1)2020

*Methods in Ecology and Evolution* (1) 2020

**PROCEEDINGS**

I have presented or attended graduate research at four national or international conferences, including both oral presentations and posters. Sessions listed here also include presentations by supervised undergraduate students.

**Pathway to Increase Standards and Competency of eDNA Surveys (PICSeS)** 2023

**International Conference** (poster presentation)

University of Guelph

**8th** **International Barcode of Life Conference** (oral presentation) 2019

NTNU University Museum and Norwegian Biodiversity Information Centre

**Guelph BioMathematics and Statistics (BioM&S) Symposium**

**Artificial Intelligence and Machine Learning in Biology** (attended)2019

University of Guelph

**CEPS Undergraduate Poster Session** (student poster presentation)2018

University of Guelph

**7th** **International Barcode of Life Conference** (oral presentation) 2017

University of Johannesburg

**6th International Barcode of Life Conference** (poster presentation) 2015

University of Guelph

**SOFTWARE DEVELOPMENT**

I have been directly or indirectly involved with the development and deployment of various software tools for molecular biodiversity analysis.

HACSim (**H**aplotype **A**ccumulation **C**urve **Sim**ulator) ⋅ R package ⋅ R Shiny web app

* A novel nonparametric stochastic (Monte Carlo) local search optimization method of iteratively generating species' haplotype accumulation curves through extrapolation to assess within-species sampling completeness
* R package and Shiny app respectively available for download through the Comprehensive R Archive Network ([CRAN](https://cran.r-project.org/web/packages/HACSim/index.html)) package repository or at [shinyappps.io](https://jphill01.shinyapps.io/HACSim/?_ga=2.135010417.1637014505.1646699846-1055418485.1646260523)
* Publication *in PeerJ Computer Science* was one of the top five most viewed papers in the category *Optimization Theory and Computation*
* Has been downloaded over 22000 times (*c*. 643 times per month) since being published in May 2019

VLF (**V**ery **L**ow **F**requency) ⋅ R package

* A tool to assess PCR errors, sequencing errors, *etc.* in the form of very low frequency variants, within DNA sequences using a sliding window approach
* R package available for download through [CRAN](https://cran.r-project.org/web/packages/VLF/index.html)
* Has been downloaded over 33000 times (*c*. 313 times per month) since publication
* Manuscript published in the *Biodiversity Data Journal*

**REFEREED WORK**

I have been directly or indirectly involved with the conceptualization, research, supervision and eventual publication of several academic and non-academic projects.

**Journal Articles**

Citations: 127 ⋅ H-index: 4 (According to Google Scholar, as of June 6, 2023)

**\*** Indicates students under my direct mentorship or supervision

**\*\*** Indicates students under my indirect mentorship or supervision

*Published or Accepted*

6. **Phillips, J.D.**, Athey, T.B.T., Hanner, R.H. and McNicholas, P.D. VLF: An R package for the analysis of very low frequency variants in DNA sequences. *Biodiversity Data Journal,* e96480*.* DOI: [10.3897/BDJ.11.e98480](https://bdj.pensoft.net/article/96480/).

5. **Phillips, J.D.**, Gillis, D.J. and Hanner, R.H. (2022). Lack of statistical rigor in DNA barcoding likely invalidates the presence of a true species’ barcode gap. *Frontiers in Ecology and Evolution*, 10: 859099. DOI: [10.3389/fevo.2022.859099](https://doi.org/10.3389/fevo.2022.859099).

4. D’Ercole, J., Dincă, V., Opler, P.A., Kondla, N.G., Schmidt, C.B., **Phillips, J.D.**, Robbins, R., Burns, J.M., Miller, S.E., Grishin, N., Zakharov, E.V., deWaard, J.R., Ratnasingham, S. and Hebert, P.D.N*.* (2020). A DNA barcode library for the butterflies of North America. *PeerJ*, 9: e11157. DOI: [10.7717/peerj.11157](https://peerj.com/articles/11157/).

3. **Phillips, J.D.**, \*French, S.H., Hanner, R.H. and Gillis, D.J. (2020). HACSim: An R package to estimate intraspecific sample sizes for genetic diversity assessment using haplotype accumulation curves. *PeerJ Computer Science*, 6(192): 1-37. DOI: [10.7717/peerj-cs.243](https://peerj.com/articles/cs-243/).

2. **Phillips, J.D.**, Gillis, D.J. and Hanner, R.H. (2019). Incomplete estimates of genetic diversity within species: Implications for DNA barcoding. *Ecology and Evolution,* 9(5):2996-3010. DOI: [10.1002/ece3.4757](https://onlinelibrary.wiley.com/doi/10.1002/ece3.4757).

1. **Phillips, J.D.**, Gwiazdowski, R.A., Ashlock, D. and Hanner, R. (2015). An exploration of sufficient sampling effort to describe intraspecific DNA barcode haplotype diversity: examples from the ray-finned fishes (Chordata: Actinopterygii). *DNA Barcodes*, 3: 66-73. DOI: [10.1515/dna-2015-0008](https://biodiversitygenomics.net/site/wp-content/uploads/2016/01/2015%20-%20Phillips%20-%20An%20exploration%20of%20sufficient.pdf).

*Submitted or Under Revision*

1. Young, R.G., \*\*Persram, M., \*\*Friesen, O., \*\*Chen, A., \*\*Yu, J., **Phillips, J.D.** and Hanner, R.H. (Under review). Incomplete and irregular reporting of the R statistical and computing environment highlights the need for citation guidelines to support scientific reproducibility. *PeerJ Computer Science.*

*In Preparation or To Be Submitted*

4. Morey, K.C., **Phillips, J.D.**, Loeza-Quintana, T. and Hanner, R.H. Haplotype diversity reveals challenges and opportunities for developing targeted detection assays for COI in Canadian freshwater fish. Targeted for *Environmental DNA.*

3. **Phillips, J.D.**, \*Singh, N., Hanner, R.H. and Gillis, D.J. The HACSim R Shiny app: A web interface to estimate specimen sampling sufficiency for species genetic diversity assessment with DNA sequence data.

2. D’Ercole, J., Dapporto, L., **Phillips, J.D.**, Dincă, V.E., Vila, R., Talavera, G. and Hebert, P.D.N. Macrogenetics of North American butterflies⎯The impact of Quaternary climatic fluctuations. Targeted for *PNAS.*

1. **Phillips, J.D.**, \*Bootsma, S.E., Hanner, R.H. and Gillis, D.J. Solving the genetic specimen sample size problem with a local search optimization algorithm. Targeted for *Methods in Ecology and Evolution.*

**Book Chapters**

*Submitted or Under Revision*

2. **Phillips, J.D**., Griswold, C.K., Young, R.G., Hubert, N. and Hanner, R.H. A measure of the DNA barcode gap for applied and basic research. Methods in Molecular Biology. Springer.

1. Hubert, N., **Phillips, J.D**. and Hanner, R.H. Delimiting species with single-locus DNA sequences. Methods in Molecular Biology. Springer.

*In Preparation or To Be Submitted*

N/A

**Conference Proceedings**

4. Morey, K., Loeza-Quintana, T., **Phillips, J.** and Hanner R. (2023). Haplotype diversity reveals challenges and opportunities for developing targeted detection assays for *COI* in Canadian freshwater fish. Pathways to Increase Standards and Competency in eDNA Surveys (PISCeS) Conference. Poster.

3. **Phillips, J.D.**, Gillis, D. and Hanner, R. (2019). HACSim: Iterative extrapolation of haplotype accumulation curves for assessment of intraspecific COI DNA barcode sampling completeness Scientific abstracts from the 8th International Barcode of Life Conference, Trondheim, Norway (ed. Torbjørn Ekrem), *Genome*, 62(6): 349-453. Oral presentation.

2. **Phillips, J.D.**, Gillis, D. and Hanner, R. (2017). Intraspecific sample size estimation for DNA barcoding: Are current sampling levels enough? Scientific abstracts from the 7th International Barcode of Life Conference, Johannesburg, South Africa (ed. M. van der Bank), *Genome*, 60(11): 881-1019. Oral presentation.

1. **Phillips, J.D.**, Gwiazdowski, R.A., Ashlock, D. and Hanner, R. (2015). An exploration of sufficient sampling effort to describe intraspecific haplotype diversity in the ray-finned fishes (Chordata: Actinopterygii). Scientific abstracts from the 6th International Barcode of Life Conference, Guelph, ON., Canada (ed. S.J. Adamowicz), *Genome*, 58(5): 163-303. Poster

**NON-REFEREED WORK**

I have communicated aspects of my research to non-technical audiences through various online media.

**Blog posts**

6. **Phillips, J.D.** (2022) Mind the Gap ⎯ The DNA Barcode Gap, That Is. Contributed CEPS Research Highlights article (https://www.uoguelph.ca/ceps/news/2022/08/mind-gap-–-dna-barcode-gap).

5. **Phillips, J.D.** (2020). Barcode Cracking. Contributed CEPS Research Highlights article (https://www.uoguelph.ca/ceps/news/2020/02/barcode-cracking).

4. **Phillips, J.D.** (2020). Protecting Biodiversity Through the Lens of Genetic Diversity. Contributed guest post to the blog of Dr. Daniel Gillis (https://danielgillis.wordpress.com/2020/01/30/protecting-biodiversity-through-the-lens-of-genetic-diversity/).

3. **Phillips, J.D.** (2019). IBOL8 and the Midnight Sun. Contributed guest post to the blog of Dr. Daniel Gillis (https://danielgillis.wordpress.com/2019/07/02/reflections-ibol8-and-the-midnight-sun/).

2. **Phillips, J.D.** (2017). The Big Five and IBOL7. Contributed guest post to the blog of Dr. Daniel Gillis (https://danielgillis.wordpress.com/2017/12/06/reflections-the-big-five-and-ibol7/).

1. **Phillips, J.D.** (2016). Sample size estimation for DNA barcoding: Are current sampling levels enough? Contributed guest post to the DNA Barcoding Blog of Dr. Dirk Steinke (http://dna-barcoding.blogspot.com/2016/01/guest-post-sample-size-estimation-for.html).

**Newsletters**

1. **Phillips, J.D.** (2016). Sample size estimation for DNA barcoding of ray-finned fishes: Are current sampling levels enough? Contributed newsletter article to the Barcode Bulletin, 7(1).

**VOLUNTEER EXPERIENCE**

2. **Pathways to Increase Standards and Competency in eDNA Surveys (PISCeS) Conference**

University of Guelph 2023

* Participated in international eDNA conference hosted by the Hanner Lab
* Duties included registration, upload, and audiovisual support

1. **Wireframing session** 2021

University of Guelph

* Participated in student-led use case mobile app prototype demonstrations for CIS\*3750 – System Analysis and Design in Applications
* Graded students based on several factors via Qualtrics surveys

**REFERENCES**

Dr. Daniel Gillis Dr. Robert Hanner

Associate Professor Full Professor

School of Computer Science Department of Integrative Biology

University of Guelph Biodiversity Institute of Ontario 50 Stone Road East University of Guelph

Guelph, ON. Guelph, ON.

N1G 2W1 50 Stone Road East

[dgillis@uoguelph.ca](mailto:dgillis@uoguelph.ca) N1G 2W1

(519) 824-4120 ext. 53479

[rhanner@uoguelph.ca](mailto:rhanner@uoguelph.ca)

Dr. Deborah Stacey Dr. Graham Taylor

Associate Professor Full Professor

School of Computer Science School of Engineering

University of Guelph University of Guelph

50 Stone Road East 50 Stone Road East

Guelph, ON. Guelph, ON.

N1G 2W1 N1G 2W1

[dastacey@uoguelph.ca](mailto:dastacey@uoguelph.ca) (519) 824-4120 ext. 53644

[gwtaylor@uoguelph.ca](mailto:gwtaylor@uoguelph.ca)

Dr. Dirk Steinke Dr. Sarah Adamowicz

Associate Director – Analytics Associate Professor

Adjunct Professor Department of Integrative Biology

Centre for Biodiversity Genomics University of Guelph

Department of Integrative Biology 50 Stone Road East

University of Guelph N1G 2W1

50 Stone Road East Guelph, ON.

N1G 2W1 (519) 824-4120 ext. 53055

Guelph, ON. [sadamowi@uoguelph.ca](mailto:sadamowi@uoguelph.ca)

(519) 824-4120 ext. 53759

[dsteinke@uoguelph.ca](mailto:dsteinke@uoguelph.ca)