

# Rhiannon Cameron

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## BACHELOR OF SCIENCE MICROBIOLOGY

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## EDUCATION

- 2020-2022\* M.Sc. Bioinformatics // UNIVERSITY OF BRITISH COLUMBIA, B.C. // *\*In-Progress*  
CGPA: 4.33/4.33 (A+)
- 2019 B.Sc. Microbiology // UNIVERSITY OF VICTORIA, B.C.  
CGPA: 4.33/4.33 (A+)

## AWARDS

### NSERC-CREATE Bioinformatics Scholarship Awardee (\$24,300)

JUN 2020 | NSERC CREATE Bioinformatics Graduate Training Program Grant (2020-2021)

- Summer 2021 Internship with Dr. Athanasios Zovoilis at the Southern Alberta Genome Sciences Centre RNA Genomics Laboratory.

## SKILLS & EXPERTISE

**Research Interests** // Ontologies, Bioinformatics, Genomics, Metabolomics, Infectious diseases, Public Health, Human microbiome, Microbial ecology, Signal transduction pathways, Metabolism, UX design.

**Computation** // Python (Proficient), R (Proficient), C++ (Basics), Linux/Windows/MacOS Command line/Terminal, Version control (GitHub), BASH (RegX), Bioinformatics: BLAST, MSA, Genome Databases/Organizers (VOCS, VGO), Phylogenetics (e.g. MEGA, Dendroscope, Geneious), Nextflow, Conda.

**Math/Statistics** // Calculus, Linear Algebra, Statistics (e.g. Binomial, Poisson, and Normal distribution; ANOVA; Regression and Correlation).

**Laboratory\*** // Fundamental biochemical, microbiological and molecular biological laboratory techniques including but not limited to: SDS-PAGE, Protein Purification, ELISA, Culturing Hybridomas, Gel Electrophoresis, Molecular Cloning, Transposon Mutagenesis, and PCR.

**Other** // Markdown, LaTeX, Academic Writing, Grant Writing, Graphic Design, TESL, French (Conversational).

*\*My ability to perform laboratory bench work is severely impacted by my physical disability.*

## RESEARCH EXPERIENCE

### **CENTRE FOR INFECTIOUS DISEASE GENOMICS AND ONE HEALTH (CIDGOH) // GRADUATE STUDENT RESEARCHER**

SEP 2020 – PRESENT | DR. WILLIAM HSIAO | SIMON FRASER UNIVERSITY, B.C.

**Thesis** work involves ontology curation and development for outbreak investigation and surveillance of SARS-CoV-2 for the Canadian COVID-19 Genomics Network ([CanCOGeN](#)).

- Assist in COVID-19 DataHarmonizer tool development, quality control, user testing, as well as the development and maintenance of associated standard operating procedures.
- Collaborate with the National Microbiology Laboratory (NML) on CanCOGeN metadata initiatives.
- Contribute to, demo, and present at CanCOGeN provincial workshops/training sessions.
- Write public health reports (e.g. "CanCOGeN VirusSeq – Comparison and Analysis of Canadian Public Health SARS-CoV-2 Case Report Forms").
- Continuation of work pertaining to previous roles (see "Research Assistant / Database Curator") working on GenEpiO, FoodOn, and the Joint Food Ontology (JTO).

### **SOUTHERN ALBERTA GENOME SCIENCES CENTRE (SAGSC) // NSERC-CREATE INTERN**

MAY 2021 – AUG 2021 | DR. ATHANASIOS ZOVOILIS | UNIVERSITY OF LEHTBRIDGE, A.B.

- Updating and utilizing Liam Mitchell's A-to-I editing nextflow pipeline on nanopore human brain tissue RNA sequence data, focusing on the identification of novel RNAs with a special interest in repetitive RNAs, ncRNAs, and SINE RNAs.
- Working remotely on high-memory high-performance CPU and GPU nodes.
- Using R to apply statistical analyses to the pipeline output.

### **HSIAO PUBLIC HEALTH BIOINFORMATICS LABORATORY // RESEARCH ASSISTANT/CURATOR**

FEB 2020 – SEP 2020 | DR. WILLIAM HSIAO | CENTER FOR DISEASE CONTROL, B.C.

- Collect and curate SARS-CoV-2 and COVID-19 related metadata for the Canadian COVID Genomics Network (CanCOGeN) initiative.
- Beta tester for the CanCOGeN covid19ValidationGrid; test guidelines and functions, meticulously explore possible usage scenarios, and provide ideas for improvement.
- Produce sample data and quality control information for validating CanCOGeN tools.
- Content creator for the CanCOGeN metadata wiki.
- Manual curation and term edits for the FoodOn and GenEpiO ontology databases using Protégé and GitHub.
- Generate curator training materials and perform curator demo webinars.
- Systematic literature review on the accessibility of public health genomics.

## RESEARCH EXPERIENCE (*continued...*)

### **VIRAL BIOINFORMATICS RESEARCH CENTRE // DIRECTED STUDIES STUDENT**

SEP 2017 – APR 2018 | DR. CHRIS UPTON | UNIVERSITY OF VICTORIA, B.C.

- Produced specifications for, and managed implementation of, bioinformatics tools and interfaces developed at Dr. Chris Upton's Viral Bioinformatics Research Centre (VBRC), while performing analyses of laboratory generated data and unprocessed published data (obtained from PubMed).
- Conducted a literature review and bioinformatic analysis of the novel influenza A virus gene NEG8.
- Developed a theoretical phylogenetic Tree GUI for virus selection in Viral Orthologous Clusters (VOCs) database, and proposed solutions for tools that visualized data in a manner inaccessible to colour blind individuals.
- Studied and reviewed bioinformatics tools for the analysis of virus genomes.
- Coached VBRC programmers on the functions of the bioinformatics tools from the perspective of a researcher and regularly made optimization recommendations.
- Presented detailed reports on progress and results, as well as updated VBRC online guides and documentation.

### **UNIVERSITY OF REGINA // RESEARCH ASSISTANT**

MAY 2014 – AUG 2014 | DR. ANDREW CAMERON | REGINA, S.K.

- Generated significant and well-controlled scientific research results on Sxy toxicity in *Salmonella enterica typhimurium* in a level 2 bio-safety laboratory supervised by Dr. Andrew Cameron and Dr. Ebtihal Al-shabib. Research results added to a manuscript being prepared for [peer-reviewed publication](#).
- Performed laboratory tasks including but not limited to: aseptic pipette technique, PCR amplification and plasmid modification, cell separation, preparation of LB growth media, poured plates, inoculation of agar plates and broths, growth on selective media, colony isolation, serial dilution, plate counts, identification and characterizing of colonies, and gel electrophoresis preparation and application.
- Investigated and identified a source of contamination in the laboratory by testing all available laboratory equipment, surfaces, and media, while keeping thorough documentation of the process.
- Took accurate and comprehensive laboratory notes, now used as a positive example for future members of the laboratory, as well as provided regular slideshow presentations on progress and results.

## OTHER EXPERIENCE

### VANCOUVER BIOINFORMATICS USER GROUP // COMMUNICATIONS VOLUNTEER

JAN 2020 – PRESENT | VANCOUVER, B.C.

- Live webinar setup and facilitator/MC.
- Outreach, social media, and coordinating out-of-town speaker's activities.
- Assist with general event coordination activities and data management.

### NEIL SQUIRE SOCIETY // COMPUTER COMFORT VOLUNTEER

JUL 2019 – MAR 2020, MAY 2021 – PRESENT | BURNABY, B.C.

- Help adults with disabilities develop basic computer skills in weekly sessions.
- Recommend and develop resources for client to study and reference.

### INTERNATIONAL FAIR CONVERGENCE SYMPOSIUM // WORKSHOP PRESENTER

DEC 2020 | VANCOUVER, B.C.

- DataHarmonizer tool explanation and demonstration.

### BIOCHEMISTRY AND MICROBIOLOGY STUDENT SOCIETY // CO-PRESIDENT (Elected Position)

APR 2015 – DEC 2015 | UNIVERSITY OF VICTORIA, B.C.

- Directed and delegated to a group of my peers.
- Demonstrated communication and dispute resolution skills.
- Developed and operated within budgets, managerial accounting.
- Organized and managed BMSS events on and off campus.
- Conducted team building and training activities.

## RELEVANT TRAINING & CERTIFICATIONS

2021	<b>SAGSC //</b> Introduction to Bioinformatics <b>CIHR IRSC //</b> Core Competency Module for Sex and Gender in Biomedical Research < <i>Credential Upon Request</i> > <b>CIHR IRSC //</b> Core Competency Module for Sex and Gender in Primary Data Collection with Human Participants < <i>Credential Upon Request</i> > <b>CIHR IRSC //</b> Core Competency Module for Sex and Gender in the Analysis of Secondary Data from Human Participants < <i>Credential Upon Request</i> >
2020	<b>EPI-interactive //</b> R Shiny Masterclass Series < <i>Credential Upon Request</i> > <b>TCPS 2: CORE //</b> Tri-Council Policy Statement: Ethical Conduct for Research Involving Humans Course on Research Ethics < <i>Credential Upon Request</i> >

- 2020      **FSCI 2020 Course //** Advancing FAIR Data Stewardship: Fostering Institutional Planning and Service Development <[Details](#)>  
**FSCI 2020 Course //** Looking Beyond the Journal Article: A Guide to Building Your Biomedical Research Toolkit <[Details](#)>  
**Ladies Learning Code //** Using Data to Solve Problems: An Introduction to Artificial Intelligence and Machine Learning for Beginners <[Workshop](#)>
- 2019      **MITx (edX) 6.00.1x //** Introduction to Computer Science and Programming Using Python <[Credential](#)>  
**MITx (edX) 6.00.2x //** Introduction to Computational Thinking and Data Science <[Credential](#)>  
**Ladies Learning Code //** JavaScript for Beginners: An Introduction to the Fundamentals of Web Programming <[Workshop](#)>

## POSTERS, PRESENTATIONS, AND ABSTRACTS

- **Cameron, R.**, Dooley, D., Griffiths, E., Savić Kallesøe, S., Hsiao, W. Putting standards into practice – ontologizing the Canadian COVID genomics network specification. Poster presenting at: The Bioinformatics Open Source Conference (BOSC); Jul 27-29, 2021. Virtual. Available from: <[pending publication](#)>
- Dooley, D., Griffiths, E., **Cameron, R.**, Gosal, G., Naravane, T., Walls, R.L., Buttigieg, P., Hsiao, W. FoodOn evolution in the food ontology ecosystem. Abstract for: The Integrated Food Ontology Workshop (IFOW); Sep-Oct, 2020. Virtual. Available from: [https://github.com/FoodOntology/joint-food-ontology-wg/blob/master/presentation/IFOW\\_2020\\_sept\\_16\\_in\\_the\\_field/IFOW\\_2020\\_17\\_FoodOn\\_Evolution\\_in\\_the\\_Food\\_Ontology\\_Ecosystem.pdf](https://github.com/FoodOntology/joint-food-ontology-wg/blob/master/presentation/IFOW_2020_sept_16_in_the_field/IFOW_2020_17_FoodOn_Evolution_in_the_Food_Ontology_Ecosystem.pdf)

## REPORTS

- **Cameron, R.**, Savić Kallesøe, S., Griffiths, E., Hsiao, W. CanCOGeN VirusSeq Comparison and analysis of Canadian public health SARS-CoV-2 case report forms. CanCOGeN VirusSeq, 2020. Available from: [https://virusseq.ca/wp-content/uploads/2021/04/2020-12-10\\_CRF-Report-.pdf](https://virusseq.ca/wp-content/uploads/2021/04/2020-12-10_CRF-Report-.pdf)

## REFERENCES

**Dr. William Hsiao** | Principal Investigator & SFU Associate Professor  
The Centre for Infectious Disease Genomics and One Health  
Simon Fraser University  
Burnaby, B.C.  
[Contact Information: Omitted]

**Dr. Damion Dooley** | Scientific Programmer  
The Centre for Infectious Disease Genomics and One Health  
Simon Fraser University  
Burnaby, B.C.  
[Contact Information: Omitted]

**Dr. Emma Griffiths** | GenEpiO Project Coordinator, Curator and Research Associate  
The Centre for Infectious Disease Genomics and One Health  
Simon Fraser University  
Burnaby, B.C.  
[Contact Information: Omitted]

**Dr. Chris Upton** | Professor & VBRC Principal Investigator  
Department of Biochemistry and Microbiology  
& Viral Bioinformatics Research Centre  
University of Victoria  
Victoria, B.C.  
[Contact Information: Omitted]

**Dr. Perry Howard** | Professor and Department Chair  
Department of Biochemistry and Microbiology  
University of Victoria  
Victoria, B.C.  
[Contact Information: Omitted]

**Dr. Andrew Cameron** | Associate Professor & IMSS Co-Principal Investigator  
Department of Biology  
& Institute for Microbiology Systems and Society  
University of Regina,  
Regina, S.K.  
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