Madeleine S. Gastonguay

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Educations

University of Connecticut, Storrs, CT — May 2020
Bachelor of Science, Applied Mathematics
Summa Cum Laude with Honors
GPA: 3.98/4.00
Minor: Bioinformatics
Thesis: A Quantitative Pipeline for The Identification of Combinations of Targets for Claudin-Low Triple Negative Breast Cancer Reversion
Advisor: Dr. Paola Vera-Licona

La Sorbonne University, Paris, France — January 2018-May 2018
Course de Civilisation Française

Research Experience

The Jackson Laboratory (JAX), Bar Harbor, ME — June 2020 - present
Research Data Analyst I
Topic: A Bayesian approach to mediation analysis of complex traits with measurement noise

- Contributed to the development and validation of an R package implementation of a Bayesian model selection approach to mediation analysis that is flexible in both data inputs and potential inferences, and extended it to moderated mediation
- Diagnosed the effect of measurement noise on the inference of mediation
- Applied these tools to better understand the mechanism underly the effects of sex and diet on protein expression in the livers of genetically diverse mice

UConn Health Center for Quantitative Medicine, Farmington, CT — September 2018 - May 2020
Undergraduate Research Assistant
Topic: A quantitative pipeline for cancer reversion analysis in triple negative breast cancer

- Awarded a Summer Undergraduate Research Fund through UConn to fund work
- Constructed a static intracellular signaling network for a claudin-low triple negative breast cancer (CL TNBC) cell line with multi-omics data using Cytoscape and GeneXplain
- Applied a structure-based control method for nonlinear systems, implemented in python, to identify putative targets that steer the system to any desired attractor
- Conducted virtual screenings using the topology of the network and a signal propagation algorithm to identify concerted perturbations of control targets resulting in reversion of the CL TNBC phenotype

Metrum Research Group, Simsbury, CT — June - August 2018
Summer Intern
Topic: An open and general maternal-fetal physiologically-based pharmacokinetic model for drugs metabolized by cytochromes P450 isoenzymes

- Modeled maternal and fetal drug exposures at different gestational ages by incorporating anatomical, biochemical, and physiological changes associated with pregnancy as a system of differential equations using R and *mrgsolve*
- Performed local sensitivity analysis, optimized model parameters, and validated the model by comparing model predictions to external published data

University of Connecticut, Department of Molecular and Cellular Biology, Storrs, CT — January - September 2017
Undergraduate Research Assistant/Holster Scholar
Topic: The effect of host genetic variability on Epstein Barr Virus (EBV)-derived cancer susceptibility

- Reviewed relevant literature to identify genes that may impact EBV-derived cancer susceptibility
- Used wet lab techniques such as Polymerase Chain Reactions, Gel Electrophoresis, and Sanger Sequencing to sequence the genes of interest in several EBV-derived cancer cell lines and a non-cancerous EBV control cell line
- Aligned the genetic sequence of target genes to identify common and distinct single nucleotide polymorphisms across EBV-derived cancer cell lines using the software Geneious and BLAST

Publications and Pre-Prints

Gastonguay, M. S., Keele, G. R., & Churchill, G. A. (2021). The impact of measurement noise on mediation analysis. *Manuscript in preparation for submission*

Crouse, W. L., Keele, G. R., Gastonguay, M. S., Churchill, G. A., & Valdar, W. (2021). A Bayesian model selection approach to mediation analysis. *BioRxiv*, 2021.07.19.452969. https://doi.org/10.1101/2021.07.19.452969

Utsey, K., Gastonguay, M. S., Russell, S., Freling, R., Riggs, M. M., & Elmokadem, A. (2020). Quantification of the Impact of Partition Coefficient Prediction Methods on Physiologically Based Pharmacokinetic Model Output Using a Standardized Tissue Composition. *Drug Metabolism and Disposition*, *48*(10), 903 LP – 916. https://doi.org/10.1124/dmd.120.090498

Zuppa, A. F., Benitez, G. R., Zane, N. R., Curley, M. A. Q., Bradfield, J., Hakonarson, H., Gastonguay, M. S., Moorthy, G., Prodell, J., & Gastonguay, M. R. (2019). Morphine Dose Optimization in Critically Ill Pediatric Patients With Acute Respiratory Failure. *Critical Care Medicine*, *47*(6), e485–e494. https://doi.org/10.1097/CCM.0000000000003741

Zuppa, A. F., Conrado, D. J., Zane, N. R., Curley, M. A. Q., Bradfield, J., Hakonarson, H., Gastonguay, M. S., Moorthy, G., Prodell, J., & Gastonguay, M. R. (2019). Midazolam Dose Optimization in Critically Ill Pediatric Patients With Acute Respiratory Failure. *Critical Care Medicine*, *47*(4), e301–e309. https://doi.org/10.1097/CCM.0000000000003638

Oral Presentations

Gastonguay MS, Russell S, Freling R, Utsey K, and Elmokadem A, *Prediction of maternal-fetal exposures of CYP450-metabolized drugs using physiologic pharmacokinetic modeling implemented in R and mrgsolve.,* R/Pharma Conference, Cambridge, MA, August 23rd, 2019

Gastonguay MS, Marazzi L, Vera-Licona P, *Identification of Combinations of Targets for Claudin-Low Triple Negative Breast Cancer Reversion,* UConn Center for Quantitative Medicine, July 30th, 2019

Gastonguay MS, Marazzi L, Vera-Licona P, *Identification of Combinations of Targets for Claudin-Low Triple Negative Breast Cancer Reversion,* UConn Center for Cell Analysis and Modeling Summer Seminar, July 26th, 2019

Gastonguay MS, Russell S, Freling R, Utsey K, and Elmokadem A, *Development of an Open and General Physiologically Based Pharmacokinetic Model to Predict Maternal-Fetal Exposures for Drugs Metabolized by CYP Isoenzymes,* R/Medicine Conference, New Haven, CT, September 8th, 2018

Gastonguay MS, *The Effect of Host Genetic Variability on Epstein Barr Virus-derived cancer susceptibility,* UConn Holster Scholar Symposium, October 2017

Poster Presentations

Gastonguay MS, Marazzi L, Vera-Licona P, *Identification of Combinations of Pharmacologic Targets for Claudin-Low Triple Negative Breast Cancer Reversion,* International Society of Pharmacometrics Quantitative Systems Pharmacology Student Symposium, April 28th, 2021

Gastonguay MS, Marazzi L, Vera-Licona P, *Identification of Combinations of Targets for Claudin-Low Triple Negative Breast Cancer Reversion,* Joint Meeting in Mathematics, Denver, CO, January 15th – 18th, 2020

Gastonguay MS, Russell S, Freling R, Utsey K, and Elmokadem A, *Development of an Open-source Physiologically-Based Pharmacokinetic Model to Predict Maternal-Fetal Exposures of CYP450-Metabolized Drugs,* International Society of Pharmacometrics Regional Quantitative Systems Pharmacology Day, Princeton, NJ, July 16th, 2019

Gastonguay MS, Russell S, Freling R, Utsey K, and Elmokadem A, *Development of an Open-source Physiologically-Based Pharmacokinetic Model to Predict Maternal-Fetal Exposures of CYP450-Metabolized Drugs,* University of Connecticut Frontiers in Undergraduate Research, April, 2019

Fellowships and Grants

Summer Undergraduate Research Fund (SURF) Trimble Family Award May 2019
University of Connecticut Office of Undergraduate Research ($4,000)

Holster Scholar May 2017
University of Connecticut Honors Program ($4,000)

Honors and Awards

Blue Ribbon Poster Award, ISoP Quantitative Systems Pharmacology Student Symposium — April 2021
Dean's List, The University of Connecticut — Sept. 2016 - May 2020
Academic Excellence Scholarship, The University of Connecticut — Sept. 2016 - May 2020
Babbidge Scholar, The University of Connecticut — Dec. 2017, 2019
New England Scholar, The University of Connecticut — Dec. 2018
Global Citizenship Scholarship, The University of Connecticut Education Abroad — Jan. 2017

Skills & Certifications

Technical: R; basic Python, Matlab, SQL, Julia, and bash shell; Git; LaTeX; OpenRefine; High Performance Computing with SLURM; *mrgsolve*; *shiny*; *tidyverse;* JAGS; Bayesian Data Analysis
Wet Lab: Polymerase Chain Reactions, Gel Electrophoresis, Gel Extraction, Sanger sequencing
Language: Proficient in French conversation, reading, and writing; Certified in French level B1.2 by La Sorbonne in Paris
Teaching: Certified Instructor with The Carpentries

Professional Development and Continuing Education

Attendee

Advanced Shiny, Phil Bowsher with RStudio — Sept. 2021
Introduction to Bayesian Data Analysis, Juliacon — July 2021
Carpentries Instructor Training, The Carpentries — Mar. 2021
Shiny, RMarkdown, and RStudio Connect, Phil Bowsher with RStudio — Mar. 2021
Quantitative Trait Mapping in the Diversity Outbred, University of Wisconsin-Madison — Dec. 2020
Containerization with Singularity, JAX — Oct. 2020
Introduction to HPC, JAX — Sep. 2020
Human and Mammalian Genetics and Genomics: The 61st McKusick Short Course, JAX — July 2020
Shiny Reproducibility, Joe Cheng at R/pharma 2019 — Aug. 2019
Machine Learning, Max Kuhn at R/pharma 2019 — Aug. 2019

Instructor or Teaching Assistant

Data Carpentry Genomics Workshop, Bioinformatics Training Program at JAX (Instructor) — Nov. 2021
Introductory Statistics with R, Bioinformatics Training Program at JAX (TA) — Sept. 2021
Introduction to R and RStudio, Bioinformatics Training Program at JAX (TA) — June 2021

Professional Associations

Society for Industrial and Applied Mathematics (SIAM)
International Society of Pharmacometrics (ISoP)
American Statistics Association (ASA)

Volunteer Work and Extra-Curricular Activities

Rubyfruit A Cappella, The University of Connecticut — Sept. 2016 - May 2020

- Elected as President (2019-2020), Assistant Music Director (2018 - 2019), and Treasurer (2017- 2018)
- Coordinated funding, schedules, and rehearsals to record an album and distribute it on Spotify and Apple Music
- Communicated with other executive board members to run productive fundraisers, rehearsals, and gigs

Math Motivators, The University of Connecticut — Oct. 2017 - May 2019

- Tutored high school freshmen from underprivileged schools in Hartford, CT

iGEM Genetic Engineering Team, The University of Connecticut — Sept. 2016 - Dec. 2017

- Developed and presented a project proposal for the iGEM jamboree with a team of students