

Michael J. Geuenich

Curriculum Vitae

PROFILE

I am a PhD candidate in the Molecular Genetics Department at the University of Toronto working on building Machine Learning methods to better understand immune escape in pancreatic cancer.

Email: m.geuenich@mail.utoronto.ca Github: [Michael-Geuenich](#)

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EDUCATION

2020-present | PhD, Computational Biology in Molecular Genetics, University of Toronto.

2016–2020 | B.A.&Sc., Computational Biology, Quest University Canada.

SELECT COURSES

Mathematics	Linear Algebra, Multivariable Calculus, ODEs
CS & Statistics	Algorithm Analysis & Design, Machine Learning, Artificial Intelligence, Data Analysis in R
Biology & Chemistry	Molecular Genetics, Population Genetics, Biochemistry A, Chemistry & Organic Chemistry

SKILLS

Advanced	R, Python, Snakemake
Intermediate	Git, SQL, PyTorch/PyTorch Lightning
Basic	Shell Scripting, LaTeX, PyRosetta, Shiny, Docker

RESEARCH EXPERIENCE

Sept. 2020 – present | **PhD Student — University of Toronto & Lunenfeld Tanenbaum Research Institute**

Toronto, Canada
Developing machine learning tools to better understand immune escape in pancreatic cancer. *PI: Kieran Campbell*

May. 2020 – August 2020 | **Pre-Doctoral Summer Student — Lunenfeld Tanenbaum Research Institute**

Toronto, Canada
Analysis of imaging mass cytometry breast cancer data and benchmarking as part of the development of [Astir](#), a cell identity assignment algorithm developed for single cell proteomic approaches. *PI: Kieran Campbell*

June 2019 – May 2020 | **Undergraduate Research Assistant — Fred Hutchinson Cancer Research Center**

Seattle, USA
Analysis of ChIP-Seq and CUT&RUN NGS data pertaining to the MYC transcription factor network. Familiarity with western blotting, chromatin preparation for ChIP-Seq, fixing cells for Hi-C and Flow Cytometry. *PI: Robert N. Eisenman*

May 2018 – May 2019 | **Research Assistant & Research Fellow — Quest University Canada**

Squamish, Canada
Analysis of public sequence data for the phylogenetic analysis of ribosome maturation genes and their involvement in Shwachman-Diamond Syndrome. Homology search, sequence alignment, phylogenetic tree construction & protein visualization. Awarded paid Research Fellowship (Summer 2018) to continue this research. *PI: Marina Tourlakis*

PUBLICATIONS

* Denotes joint first author

PREPRINTS

- 2022 Hassaan Maan, Lin Zhang, Chengxin Yu, **Michael Geuenich**, Kieran R Campbell, and Bo Wang. The differential impacts of dataset imbalance in single-cell data integration. *bioRxiv*, 2022
- Sebastien Martinez, Ramona Weber, Tristan Woo, Ahmad Malik, **Michael Geuenich**, Gun Ho Jang, Dzana Dervovic, Khalid Al-Zahrani, Ricky Tsai, Nassima Fodil, et al. *In vivo CRISPR screens reveal SCAF1 and USP15 as novel drivers of pancreatic cancer*. *bioRxiv*, 2022

PUBLISHED

- 2022 Connor Yanchus, Kristen L. Drucker, Thomas M. Kollmeyer, ..., **Michael Geuenich**, ..., Robert B. Jenkins, and Daniel Schramek. *A non-coding single nucleotide polymorphism at 8q24 drives IDH1-mutant glioma formation*. *Science*, 2022
- 2021 **Michael J. Geuenich***, Jinyu Hou*, Sunyun Lee*, Shanza Ayub, Hartland W. Jackson, and Kieran R. Campbell. *Automated assignment of cell identity from single-cell multiplexed imaging and proteomic data*. *Cell Systems*, 2021
- 2020 Haritha Mathsyaraja, Jonathen Catchpole, Brian Freie, Emily Eastwood, Ekaterina Babaeva, **Michael Geuenich**, Pei Feng Cheng, Jessica Ayers, Ming Yu, Nan Wu, ..., David MacPherson, and Robert N Eisenman. *Loss of MGA repression mediated by an atypical polycomb complex promotes tumor progression and invasiveness*. *Elife*, 10:e64212, 2021
- Arnaud Augert, Haritha Mathsyaraja, Ali H Ibrahim, Brian Freie, **Michael J. Geuenich**, Pei-Feng Cheng, Sydney P Alibeckoff, Nan Wu, Joseph B Hiatt, Ryan Basom, Adi Gazdar, Lucas B Sullivan, Robert N Eisenman, and David MacPherson. *MAX Functions as a Tumor Suppressor and Rewires Metabolism in Small Cell Lung Cancer*. *Cancer Cell*, 38(1):97–114.e7, July 2020

POPULAR MEDIA

- 2018 **Michael Geuenich**. *Regarding Moustaches*. *Squamish Chief*, October 2018. Opinion about cancer awareness and funding allocation in the local newspaper

PRESENTATIONS

- 2022 **Geuenich Michael J.** *With great data come great pipelines: Creating flexible standardized pipelines for common biomedical analysis tasks using Snakemake*. In *Toronto Workshop on Reproducibility*, Feb 2022. Talk
- 2019 **Geuenich Michael J.**, Brian Freie, Patrick Carroll, Vijay Ramani, and Robert N. Eisenman. *An activating point mutation in Myc (T58A) mitigates ER stress*. In *BC Cancer Summit*, Nov 2019. Poster
- 2018 **Geuenich Michael J.**, Johanna Rommens, and Marina E. Turlakis. *Identifying differences in the Shwachman- Diamond syndrome associated gene SBDS across species*. In *BC Cancer Summit*, Nov 2018. Poster
- 2018 **Geuenich Michael J.**, Johanna Rommens, and Marina E. Turlakis. *How computation and a rare disease can teach us about Basic Biology and Cancer*. In *Quest Scholarship Symposium*, Nov 2018. Invited talk

STUDENT SUPERVISION

<i>Sept. 2021 - April 2022</i>	Anna Lai — University of Toronto BCB430 Student Supervised a student in comparing mouse and human single cell RNA-Sequencing data to infer the ability of mouse models to recapitulate human pancreatic cancer subtypes.
<i>May 2021 - Aug. 2021</i>	Sean Gong — University of Toronto BCB330 Student Supervised a student in the development <i>WhatsThatCell</i> , a shiny app for manual cell type annotation.

TEACHING

Sept. 2019	Teaching Assistant — Quest University Canada <i>Molecular Genetics</i>
<i>Squamish, Canada</i>	Created a UCSF Chimera dry lab to predict the effect of mutations on a protein interaction. The assignment was designed to ease students into using software and included usage of the command line.
April 2019	Guest Lecturer — Quest University Canada <i>Linear Algebra</i>
<i>Squamish, Canada</i>	Lecture on the mathematics of PCA, using it in R, and applications in genomics.
Oct. 2018 – Nov. 2018	Guest Lecturer & Teaching Assistant — Quest University Canada <i>Ecological Genomics</i>
<i>Squamish, Canada</i>	Lecture on Multiple Sequence Alignments using MUSCLE. Teaching assistant for the R based dry lab.

AWARDS

<i>Sept. 2022-Aug. 2023</i>	University of Toronto Open Fellowship, \$12,000 – University of Toronto
<i>Sept. 2021-Aug. 2022</i>	University of Toronto Open Fellowship, \$12,000 – University of Toronto
<i>April 2020</i>	Keystone Distinction Award – Quest University Canada – Thesis award for the top 10% of students
<i>Spring 2019</i>	Presidential Honours Award – Quest University Canada – Awarded to the top 10% of students
<i>Fall 2018</i>	Presidential Honours Award – Quest University Canada
<i>Spring 2018</i>	Presidential Honours Award – Quest University Canada
<i>2016 – 2020</i>	Presidential Scholarship, \$32,000 – Quest University Canada

OTHER EMPLOYMENT

- Oct. 2016 – **Data Analyst Coordinator — Quest University Canada**
Aug. 2018 *Communications & Development Office*
Squamish, Canada Organized the university's donation data and imported it into a single database.
- Oct. 2015 – **Business Development — EIMS**
Aug. 2016
Barcelona, Spain Worked in marketing and sales for two major IT companies (BlackBerry and Riverbed), communicating with IT executives to solve critical infrastructure problems. Responsible for generating \$775,000/year from new and existing customers.
- May 2014 – **CEO & Co-Founder — BioEnvío**
July 2015
Barcelona, Spain Built an online shop to distribute organic food across the Spanish peninsula.

OUTREACH & ACADEMIC LEADERSHIP

- June. 2021 – **Computational Biology in Molecular Genetics event organizer — University of Toronto**
Present *I regularly organize events with outside and student speakers for the students in the computational track of our department.*
- Sept. 2018 – **Co-Editor in Chief — Momenta**
May 2019 *Quest's Academic Journal*
- Sept. 2017 – **Board Member & President**
May 2019 *Quest Biomedical Journal Club*
- Dec. 2018 – **Co-Founder & Organizer — Stand-Up Science**
April. 2019 *Local Science Outreach Event*
- Mar. 2019 **Topic Editor (Biology) — Our Futures**
Academic Conference
- Dec. 2017 – **Peer-reviewer — Momenta**
Aug. 2018

LANGUAGES

German	Mothertongue
Spanish	Fluent
English	Fluent
Catalan	Basic

REFERENCES

- | | |
|-------------------------|---|
| Dr. Kieran
Campbell | Advisor & Professor, University of Toronto & Lunenfeld Tanenbaum Research Institute, Affiliate
Faculty, Vector Institute |
| Dr. Marina
Tourlakis | Undergraduate Mentor & Professor, Molecular Genetics, Quest University Canada |
| Dr. Richard
Hoshino | Professor, CS & Mathematics, Northeastern University, formerly at Quest University Canada |
| Dr. Robert
Eisenman | Professor, Fred Hutchinson Cancer Research Center |
- Contact information provided upon request*