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CONTACT INFORMATION      Email: max.turgeon@umanitoba.ca      Personal website: <https://maxturgeon.ca/>

RESEARCH INTERESTS      High-dimensional data, dimension reduction techniques, multivariate analysis, manifold learning, statistical genetics, competing risk analysis, case-base sampling

EDUCATION      **University McGill**, Montréal, QC

Ph.D., Biostatistics, May 2019

- Topic: *Principal component of explained variance: High-dimensional estimation and inference*
- Advisors: Celia M.T. Greenwood, Ph.D and Aurélie Labbe, Ph.D

M.Sc., Mathematics, September 2013

- Topic: *A relative fundamental lemma for  $U(4)$*
- Advisor: Jayce R. Getz, Ph.D

**University of Ottawa**, Ottawa, ON

B.Sc., Mathematics (Honours), April 2011

- *Summa Cum Laude*
- Topic: *Representation theory of p-adic algebraic groups*
- Advisor: Monica Nevins, Ph.D

EMPLOYMENT

### Current appointment

**Assistant Professor**

Jul 2019–

Department of Statistics,  
Department of Computer Science,  
University of Manitoba, Winnipeg MB

### Previous appointments

**Business Intelligence and HR Analytics, Lead**

Oct 2018–Jun 2019

AIMS Project,  
Saskatchewan Health Authority, Saskatoon, SK

**Senior Biostatistical & Study Design Strategies Specialist** Sep 2016–Sep 2018

Strategic Health and Information Performance Support,  
Saskatoon Health Region, Saskatoon, SK

REFEREED

JOURNAL PUBLICATIONS

1. **Turgeon, M.**, Oualkacha, K., Ciampi, A., Miftah, H., Dehghan, G., Zanke, B.W., Benedet, A.L., Rosa-Neto, P., Greenwood, C.M.T., Labbe, A., for the Alzheimer’s Disease Neuroimaging Initiative. “Principal component of explained variance: An efficient and optimal data dimension reduction framework for association studies”. *Statistical Methods in Medical Research*, 27: 2018. doi:10.1177/0962280216660128

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Name of trainees is underlined.

2. Wang, Y., Murphy, O., **Turgeon, M.**, Wang, Z.Y., Bhatnagar, S.R., Schulz, J., and Moodie, E.E.M. “The perils of quasi-likelihood information criteria”, *Stat*, 4: 2015. doi:10.1002/sta4.95
3. Ahmad, O.S., Morris, J.A., Mujammami, M., Forgetta, V., Leong, A., Li, R., **Turgeon, M.**, Greenwood, C.M.T., Thanassoulis, G., Meigs, J.B., Sladek, R., and Richards, J.B. “A Mendelian randomization study of the effect of type-2 diabetes on coronary heart disease” *Nature Communications*, 6: 2015. doi:10.1038/ncomms8060

SUBMITTED  
JOURNAL  
PUBLICATIONS

4. **Turgeon, M.**, Labbe, A., Greenwood, C.M.T. “A Tracy-Widom Empirical Estimator For Valid P-values With High-Dimensional Datasets”. Submitted to *Journal of Computational and Graphical Statistics*.
5. Bhatnagar, S., **Turgeon, M.** (co-primary author), Islam, J., Hanley, J., Saarela, O. “**casebase**: An Alternative Framework For Survival Analysis and Comparison of Event Rates”. Submitted to *The R Journal*.
6. Prystajeky, M., Basi, R., **Turgeon, M.**, Mittelstadt, M., Sparrow, C. “Improving Discharge Communication on an Internal Medicine Service”. Submitted.
7. Farkas, C., Mella, A., **Turgeon, M.**, Haig, J.J. “Generation and use of a novel SARS-CoV-2 viral sequence tracking platform has found genetic evidence that the viral 3’ untranslated region (UTR) is evolving and generating increased viral diversity”. Submitted to *Emerging Microbes & Infections*.

PAPERS IN  
PREPARATION

1. **Turgeon, M.**, Hoque, M.E., Bhatnagar, S. “Absolute risk estimation using dimension reduction and case-base sampling”.
2. Islam, J., **Turgeon, M.**, Bhatnagar, S.R. “Dynamic reversion of survival functions to estimate the original dataset.”
3. **Turgeon, M.**, Islam, J., Bhatnagar, S.R., Saarela, O. “Variable selection and competing risk analysis using case-base sampling”.
4. Czubryt, T., **Turgeon, M.** “Redundancy Analysis and Principal Component of Explained Variance: A Tale of Two Disciplines”.
5. **Turgeon, M.** “Multi-precision linear algebra for computing largest root distributions using the `rootWishart` package”.

FELLOWSHIPS AND  
GRANTS

- Submitted
- NSERC Discovery Grant  
Dimension Reduction and Complex High-Dimensional Data
  - New Frontiers in Research Fund–Exploration (co-PI)  
“Unlocking the Gate” to Targeted Drug Delivery: Understanding sulfated glycosaminoglycan binding of peptide-derivatized DNA nanostructures to predictably improve their cytosolic delivery
- Awarded
- University Research Grants Program (URGP) (\$6,000) 2020
  - Faculty of Science Pedagogy Award (\$3,000) 2020
  - Gerald Clavet Fellowship (declined) 2016–2017
  - FQRNT Doctoral Scholarship (\$20,000 per year) 2014–2016
  - FQRNT Masters Scholarship (\$15,000 per year) 2011–2013
  - Thomlinson Masters’ Fellowship (\$5,500) 2011–2012

AWARDS AND HONOURS	<ul style="list-style-type: none"> <li>• Ian C.P. Smith Integrated Science Faculty Scholar 2020–2022</li> <li>• Faculty of Medicine Graduate Mobility Award 2017               <ul style="list-style-type: none"> <li>• Research stay at Stanford University</li> </ul> </li> <li>• SSC 2017 Student Research Presentation Award 2017</li> <li>• Dr. Jim Hanley Research Day Award 2016</li> <li>• McGill Graduate Excellence Award (\$10,000) 2013–2014</li> </ul>	
PRESENTATIONS	<p>Invited presentations</p> <ul style="list-style-type: none"> <li>• Statistics Seminar, Université du Québec à Montréal, Montréal, QC Mar 2021</li> <li>• Bioinformatics-Biostatistics Research Seminar, Winnipeg MB Mar 2021</li> <li>• Science Online Public Talks, University of Manitoba, Winnipeg MB Jun 2020</li> <li>• Statistics Seminar, University of Winnipeg, Winnipeg MB Feb 2020</li> <li>• Bioinformatics-Biostatistics Research Seminar, Winnipeg MB Nov 2019</li> <li>• ICOSA-Canada Chapter 2019 Symposium, Kingston ON Aug 2019</li> <li>• Maternal Adversity, Vulnerability, and Neurodevelopment (MAVAN) Methodology group, Montréal, QC Mar 2016</li> <li>• Montreal Genomics Meeting, Montréal, QC Dec 2015</li> </ul> <p>Contributed presentations</p> <ul style="list-style-type: none"> <li>• useR! 2020, St Louis MO<sup>1</sup> Jul 2020</li> <li>• Statistical Society of Canada Annual Meeting, Montréal QC Jun 2018</li> <li>• Statistical Society of Canada Annual Meeting, Winnipeg MB Jun 2017</li> <li>• 5th Annual Canadian Statistics Student Conference, Winnipeg MB Jun 2017</li> <li>• Statistical Society of Canada Annual Meeting, St. Catherines, ON May 2016</li> <li>• EBOH Annual Research Day, Montréal, QC Apr 2016</li> <li>• 4th Annual Human and Statistical Genetics Meeting, Vancouver, BC Apr 2015</li> <li>• Biostatistics Reading Group, Montréal, QC Oct 2015</li> </ul> <p style="text-align: right;">Feb 2015, Mar 2014</p> <p>Poster presentations</p> <ul style="list-style-type: none"> <li>• EBOH 50th Anniversary Conference, Montréal, QC Apr 2015</li> <li>• 4th Annual Human and Statistical Genetics Meeting, Vancouver, BC Apr 2015</li> <li>• Statistical Society of Canada Annual Meeting, Toronto, ON May 2014</li> </ul>	
TEACHING EXPERIENCE	<p>Instructor</p> <p>SCI 2000—<i>Introduction to Data Science</i> Winter 2021 Faculty of Science University of Manitoba</p> <p>STAT 3150—<i>Statistical Computing</i> Fall 2020 Department of Statistics University of Manitoba</p> <p>STAT 7200—<i>Multivariate Statistics I</i> Winter 2020 Department of Statistics University of Manitoba</p> <p>STAT 4690—<i>Applied Multivariate Analysis</i> Fall 2019 Department of Statistics University of Manitoba</p> <p>Guest Lecturer</p> <p><i>Advanced Epidemiology</i> Summer 2020 Department of Community Health and Epidemiology University of Saskatchewan</p> <p><i>Principles of Surgery</i> Winter 2019 College of Medicine</p>	

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<sup>1</sup>Moved online due to COVID-19

University of Saskatchewan

RESEARCH  
TRAINEES  
SUPERVISED

Research Assistants

- Jay Khakhariya, Coop Student Fall 2020  
Topic: *Code Comparison Web Application*
- Erfanul Hoque Summer 2020  
Topic: *Dimension reduction and case-base sampling*

Graduate Students

- Asif Ahmed Nelay, MSc Computer Science 2021–  
Topic: *Dimension reduction and anomaly detection*

Undergraduate Students

- Jiyoung Kim, FoS Undergraduate Summer Research Award Summer 2021  
Topic: *Dimension reduction and missing data*
- Thomas Czubryt, UMSU Undergraduate Research Award Summer 2020  
Topic: *Comparison of Principal Component of Explained Variance and Redundancy Analysis*
- Wanmeng Wang, FoS Undergraduate Summer Research Award Summer 2020  
Topic: *Matrix completion for imputing missing data*
- Joshua Hamilton, Joint Honours Computer Science/Statistics Winter 2020  
Topic: *Dimension reduction and natural language processing*

Thesis Committees

- Ye Su, PhD Individual Interdisciplinary Studies 2020–  
*Advisors:* Pingzhao Hu and Hani El-Gabalawy
- Hafsa Moontari Ali, MSc Computer Science 2019–  
*Advisors:* Pingzhao Hu and Yang Wang
- Meghan Chua, MSc Computer Science 2019–  
*Advisor:* Olivier Tremblay-Savard
- Kaari Landry, PhD Computer Science 2019–  
*Advisor:* Olivier Tremblay-Savard
- Elham Alfazi, PhD Statistics 2019–  
*Advisors:* Liqun Wang and Saman Muthukumarana
- Robyn Ritchie, MSc Statistics 2019–  
*Advisor:* Alexandre Leblanc
- Jingyu Wang, MSc Statistics 2018–2020  
*Advisor:* Mohammad Jafari Jozani
- Isuru Dharmasena, MSc Statistics 2018–2020  
*Advisor:* Saman Muthukumarana
- Zimo Zhu, MSc Statistics 2018–2020  
*Advisors:* Aerambamoorthy Thavaneswaran and Ruppa Thulasiram

REVIEWING  
EXPERIENCE

Journals

- International Journal of Epidemiology
- Genetic Epidemiology
- Entropy
- Communications in Statistics—Simulation and Computation

Conferences

- Regina Qu'Appelle Health Region Annual Research Day
- Canadian Society for Epidemiology and Biostatistics Annual Meeting
- Canadian Statistics Students Conference

Poster competitions

- 2020 Undergraduate Research Poster Competition, University of Manitoba, Winnipeg MB

- Annual Faculty of Science SURE Poster Competition, University of Manitoba, Winnipeg MB
- 2019 Undergraduate Research Poster Competition, University of Manitoba, Winnipeg MB

OTHER ACADEMIC Hiring Committees

ACTIVITIES	<ul style="list-style-type: none"> <li>• Faculty of Science Indigenous Scholar University of Manitoba</li> <li>• Department of Statistics University of Manitoba</li> </ul>	<p>2021</p> <p>2021</p>
	Member of various committees	
	<ul style="list-style-type: none"> <li>• Integrated Science Program Committee University of Manitoba</li> <li>• Statistics-IT Committee University of Manitoba</li> <li>• Faculty of Science-Executive Committee University of Manitoba</li> <li>• Computer Science-Graduate Studies Committee University of Manitoba</li> <li>• Statistics-Website Ad Hoc Committee University of Manitoba</li> <li>• Statistics-Research and Seminar Committee University of Manitoba</li> <li>• Epidemiology, Biostatistics and Occupational Health Student Society, McGill University</li> <li>• Biostatistics Reading Group McGill University</li> <li>• Review Committee for the Department of Pathology McGill University</li> </ul>	<p>Jul 2020-</p> <p>Jul 2020-</p> <p>Jul 2020-</p> <p>Sep 2019-</p> <p>Sep 2019-Dec 2020</p> <p>Jul 2019-Jun 2020</p> <p>Sep 2014-2016</p> <p>Sep 2014-Apr 2016</p> <p>Mar 2013</p>
	Organised Conferences	
	<ul style="list-style-type: none"> <li>• Data Science Conference Nexus University of Manitoba, Winnipeg MB</li> <li>• 6th Annual Canadian Statistics Student Conference McGill University, Montréal QC</li> </ul>	<p>May 2021</p> <p>Jun 2018</p>
	Organised Workshops	
	<ul style="list-style-type: none"> <li>• Data Science Nexus Workshops University of Manitoba, Winnipeg MB</li> <li>• 6th Annual Canadian Statistics Student Conference McGill University, Montréal QC</li> </ul>	<p>Sep 2020-Dec 2020</p> <p>Jun 2018</p>

SOFTWARE

R packages:

- `pcev`, implementing *Principal Component of Explained Variance*.
- `casebase`, implementing case-base sampling for survival analysis.
- `covequal`, implementing a test of equality of covariance matrices valid for high-dimensional data.
- `multiKernel`, implementing multivariate prediction using kernel-machine regression.
- `rootWishart`, implementing exact computations of largest root distributions in single and double Wishart settings using arbitrary-precision linear algebra.
- `funtooNorm`, providing functions for normalization of Illumina Infinium Human Methylation 450 BeadChip (Illumina 450K) data when samples are collected from multiple tissues or cell types.

Python modules:

- `umi-bayes`, providing tools for the analysis of tagged DNA sequencing reads and implementing the parametric clustering-based deduplication algorithm.