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## Research Group

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### Current Members

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| <b>Mark Lowerison (PhD candidate, Biostatistics)</b><br>Modelling cattle movement networks<br>Co-supervised with Herman Barkema, University of Calgary                          | (Sept. 2015 – ) |
| <b>Kamso Mohammed Mujaab (PhD candidate, Biostatistics)</b><br>Bayesian clinical trials & network meta-analyses<br>Co-supervised with Glen Hazelwood, University of Calgary     | (Sept. 2018 – ) |
| <b>Tahmina Akter (PhD Candidate, Biostatistics)</b><br>Variable selection in infectious disease models  | (Sept. 2019 – ) |
| <b>Chinmoy Rahul Roy (PhD student, Biostatistics)</b><br>Non-parametric spatial infectious disease models   | (Sept. 2019 – ) |
| <b>Madeline Ward (PhD student, Biostatistics)</b><br>Behavioural-change individual-level models of disease transmission<br>Co-supervised with Lorna Deeth, University of Guelph | (Sept 2020 – )  |
| <b>Ruoyu Li (PhD student, Biostatistics)</b><br>Hospital-acquired antimicrobial resistant infections<br>Co-supervised with Jenine Leal, University of Calgary                   | (Jan 2022 – )   |
| <b>Mili Roy (PhD student, Biostatistics)</b><br>Joint Modelling of Correlated Data<br>Co-supervised with Tolu Sajobi, University of Calgary                                     | (March 2022 – ) |
| <b>Yirao Zhang (PhD student, Biostatistics)</b><br>Composite spatial individual-level models of disease transmission<br>Co-supervised with Lorna Deeth, University of Guelph    | (Sept 2022 – )  |
| <b>Haydn Hornbeck (PhD candidate, Computer Science)</b><br>Curve-fitting approaches to COVID-19 forecasting<br>Co-supervised with Usman Alim, University of Calgary             | (Sept. 2022 – ) |

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<b>Jeffrey Peitsch (MSc student, Biostatistics)</b>	(Sept 2023 – )
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Directional spatial individual-level models of disease transmission  
Co-supervised with Gyanendra Pokharel, University of Winnipeg

<b>Evans Mensah (MSc student, Biostatistics)</b>	(Sept 2023 – )
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Forecasting with behavioural-change population-level epidemic models

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### Past Members

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<b>Raja Ben Hajria (Postdoctoral Research Fellow)</b>	(Fall 2021 – Fall 2023)
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Hidden Markov Individual-level Models of Disease Transmission  
Co-supervised with Alex Schmidt, McGill University

<b>Caitlin Ward (Postdoctoral Research Fellow)</b>	(Fall 2021 – Fall 2022)
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Behavioural-change individual-level models of disease transmission  
Co-supervised with Alex Schmidt, McGill University

<b>David Vickers (Postdoctoral Research Fellow)</b>	(Fall 2021 – Fall 2022)
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Modelling COVID-19 pandemic spread  
Co-supervised with Tyler Williamson, UCalgary

<b>Leila Amiri (Postdoctoral Research Fellow)</b>	(Fall 2019 – Fall 2021)
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Spatial models for infectious disease transmission in heterogeneous systems  
Co-supervised with Mahmoud Torabi, University of Manitoba

<b>Mojtaba Pasha (Postdoctoral Research Fellow)</b>	(Summer 2019 – Summer 2021)
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Optimal design of control charts

<b>(Grace) Pui Sze Kwong (Postdoctoral Research Fellow)</b>	(Fall 2009 – Fall 2014)
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Spatio-temporal analysis of porcine respiratory and reproductive syndrome in Ontario /  
Efficient forms of individual level models for infectious disease spread

<b>Gyanendra Pokharel (Postdoctoral research fellow)</b>	(May 2015 – July 2018)
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Approximate methods of inference for spatial infectious disease models /  
Bayesian clinical trials and patient centred medicine

**Vineetha Warriyar (Postdoctoral research fellow)** (May 2016 – Aug 2018)  
Democratizing complex infectious disease data analysis

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**Lorna Deeth (PhD, Statistics)** (Fall 2007 – Winter 2013)  
Latent-conditional models of infectious disease and related topics

**Lin Zhang (PhD, Statistics)** (Fall 2009 – Fall 2013)  
Time-varying individual-level infectious disease models

**Jourdan Gold (PhD, Statistics)** (Fall 2008 – Winter 2015)  
Computational inference for network-based individual-level models of disease transmission  
Co-supervised with Zeny Feng, University of Guelph

**Rajat Malik (PhD, Statistics)** (Fall 2010 – Winter 2015)  
Sampling-based likelihood approximations for infectious disease models and related topics

**Nadia Bifulchi (PhD, Statistics)** (Fall 2010 – Winter 2015)  
Individual-level models for use with incomplete infectious disease data and related topics  
Co-supervised with Zeny Feng, University of Guelph

**Gyanendra Pokharel (PhD, Statistics)** (Fall 2011 – Winter 2015)  
Back-calculation, classification & emulation-based inference for spatial disease models

**Razvan Romanescu (PhD, Statistics)** (Fall 2012 – Summer 2016)  
Modelling heterogeneity in infectious disease systems for inference and monitoring

**Waleed Almutiry (PhD, Statistics)** (Jan 2014 – Aug 2018)  
Network uncertainty in infectious disease systems

**Justin Angevaare (PhD, Statistics)** (Fall 2014 – Fall 2020)  
Infectious disease models incorporating pathogen genomic sequence data  
Co-supervised with Zeny Feng, University of Guelph

**Carolyn Augusta (PhD, Statistics)** (Fall 2014 – Fall 2020)  
Deep learning of infectious disease systems  
Co-supervised with Graham Taylor, University of Guelph

**Syed Ali Naqvi (PhD, Biostatistics)** (Sept. 2017 – Dec. 2021)

Machine learning tools for understanding mastitis epidemiology  
Co-supervised with Herman Barkema, University of Calgary

**Md. Mahsin (PhD candidate, Statistics)** (Sept. 2015 – June 2022)

Modelling spatial heterogeneity in infectious disease data

**Babak Habibzadeh (MSc, Statistics & URA)** (Fall 2009)

Misspecification of latent and infectious periods in space-time infectious disease models

**Hau Yi (Helen) Chung (MSc, Statistics)** (Winter 2009)

Individual-level models applied to an equine-influenza outbreak

**Sanjeena Dang (née Subedi) (MSc, Statistics & URA)** (Summer 2009)

SNP selection methods: modelling the expected breeding value of Holstein Cattle

**Abbie Gardener (MSc, Statistics & URA)** (Summer 2010)

Goodness-of-fit measures for individual-level infectious disease models in a Bayesian framework

**Irene Vrbik (MSc, Statistics)** (Summer 2010)

Modelling the spatio-temporal dynamics of combustion

**Daria (Dasha) Martchenko (MSc, Statistics)** (Winter 2011)

Designing experiments to assess the spatio-temporal dynamics of crop disease.

**Mingying Fang (MSc, Statistics)** (Summer 2011)

Generalizing individual-level models of infectious disease spread

**Xuan Fang (MSc, Statistics)** (Winter 2012)

Computational gains via a discretization of the parameter space in individual-level models of infectious disease

**Angie Dobbs (MSc, Statistics)** (Winter 2012)

On computational efficiency and model approximation for spatial individual-level infectious disease models

**Longyao (Chloe) Cai (MSc, Statistics)** (Fall 2011 – Winter 2013)

Logistic growth models for estimating vaccination effects in transmission experiments

**Carolyn Augusta (MSc, Statistics)** (Sept 2012 – Summer 2014)

Fast inference for spatial infectious disease models

**Lea Enns (MSc, Statistics)** (Fall 2014 – Fall 2015)

Individual level models of infectious disease transmission for animal experiments

**Susannah Ripley (USRA)** (Summer 2014)

Random forest-based insect species identification

**Anu Stanley (MSc, Statistics & URA)** (Fall 2013 – Winter 2015)

Early prediction of seasonal influenza using school absenteeism data

**Tahsin Ferdous (MSc, Biostatistics)** (Sept. 2017 – Sept 2019)

On the effect of ignoring within-unit infectious disease dynamics when modelling spatial transmission

**Behnaz Jafari (MSc, Statistics)** (Sept. 2017 – Jan 2020)

Bias in individual-level infectious disease models

**Thet Nyein (MSc student, Statistics)** (Sept 2021 – June 2023)

Data subset-based methods of inference for spatial epidemic models  
Co-supervised with Lorna Deeth, University of Guelph

**Danika Lipman (MSc student, Statistics)** (Sept 2022 – August 2023)

A Bayesian variable selection model for semi-continuous responses using Gaussian processes  
Main supervisor: Thierry Chekouo, University of Minnesota

**Matthew Baxter (MSc student, Artificial Intelligence)** (Fall 2021 – Winter 2023)

Parameter estimation using random forests for two-stage individual-level models of infectious disease spread  
Co-supervised with Lorna Deeth, University of Guelph

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<b>William Lee (URA)</b>	(May 2016 – Dec. 2016)
Analysis of infectious disease surveillance data	
<b>Madeline Ward (URA)</b>	(May 2018 – Aug 2018)
Approaches to disease surveillance using predictive covariates Co-supervised with Lorna Deeth, University of Guelph	
<b>Arthur Novaes de Amorim (RA)</b>	(May 2019 – April 2020)
Predicting magnitude, timing, peak and duration of influenza at ER level Co-supervised with Vineet Saini, Alberta Health Services	
<b>Zeyi Liu (RA)</b>	(March 2020 – Nov. 2020)
Spatial metapopulation models of Covid-19	
<b>Emil Hodzic-Santor (URA)</b>	(May 2022 – Aug 2022 & May 2023 – Aug 2023)
Edge effects in spatial epidemic models	

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<b>Scott Hunt (PhD student – withdrew)</b>	(Fall 2012 – Spring 2014)
Bayesian optimal design of animal transmission experiments	
<b>Tulsi Paudel (PhD student – withdrew)</b>	(Fall 2012 – Jan 2016)
Identifying super-spreaders in spatial infectious disease systems	
<b>Salha Qahl (PhD/MSc student – withdrew)</b>	(July 2015 – Sept 2023)
Classification-based inference for infectious disease systems	