Research Group	
Dirk Douwes-Schultz (CANSSI Distinguished PDF)	(Sept 2024 – )
Hidden Markov Spatial Epidemic Models Co-supervised with Alex Schmidt, McGill University	
Jessica Pavani (PDF)	(Sept 2024 – )
Complex Behavioural-change in Epidemic Models Co-supervised with Alex Schmidt, McGill University	
Ruoyu Li (PhD candidate, Biostatistics)	(Jan 2022 – )
Hospital-acquired antimicrobial resistant infections Co-supervised with Jenine Leal, University of Calgary	
Mili Roy (PhD candidate, Biostatistics)	$\left( \mathrm{March}\ 2022-\right)$
Joint Modelling of Correlated Data Co-supervised with Tolu Sajobi, University of Calgary	
Yirao Zhang (PhD candidate, Biostatistics)	$(\mathrm{Sept}\ 2022-)$
Composite spatial individual-level models of disease transmission Co-supervised with Lorna Deeth, University of Guelph	
Haysn Hornbeck (PhD candidate, Computer Science)	(Sept. 2022 - )
Curve-fitting approaches to COVID-19 forecasting Co-supervised with Usman Alim, University of Calgary	
Sorif Hossein (PhD student, Biostatistics)	(Sept 2024 – )
Behavioural change epidemic models incorporating noisy alarm functions Co-supervised with Gyanendra Pokharel, University of Winnipeg	
Md Iqramul Haq (PhD student, Veterinary Medical Sciences)	(Jan. 2025 -)
Analysis of antimicrobial resistance data using modern statistical approaches Co-supervised with Diego Nobrega, University of Calgary	
Yicheng Mao (Visiting PhD student, Statistics)	$({\rm May}\ 2025-)$
Population memory mechanisms in behavioural change epidemic models Co-supervised with Lorna Deeth, University of Guelph	

# Parker Wieck (MSc student, Statistics)

(Sept 2024 - )

Fast Python-based inference for spatial disease models

#### Past Members

# Raja Ben Hajria (Postdoctoral Research Fellow)

(Fall 2021 - Fall 2023)

Hidden Markov Individual-level Models of Disease Transmission Co-supervised with Alex Schmidt, McGill University

## Caitlin Ward (CANSSI Distinguished PDF)

(Fall 2021 – Fall 2022)

Behavioural-change individual-level models of disease transmission Co-supervised with Alex Schmidt, McGill University

# David Vickers (Postdoctoral Research Fellow)

(Fall 2021 – Fall 2022)

Modelling COVID-19 pandemic spread Co-supervised with Tyler Williamson, UCalgary

#### Leila Amiri (Postdoctoral Research Fellow)

(Fall 2019 – Fall 2021)

Spatial models for infectious disease transmission in hetereogeneous systems Co-supervised with Mahmoud Torabi, University of Manitoba

#### Mojtaba Pasha (Postdoctoral Research Fellow)

(Summer 2019 – Summer 2021)

Optimal design of control charts

### (Grace) Pui Sze Kwong (Postdoctoral Research Fellow)

(Fall 2009 – Fall 2014)

Spatio-temporal analysis of porcine respiratory and reproductive syndrome in Ontario / Efficient forms of individual level models for infectious disease spread

#### Gyanendra Pokharel (Postdoctoral research fellow)

(May 2015 – July 2018)

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

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 Bifolchi (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, Statistics)

(Sept. 2015 – June 2022)

Modelling spatial heterogeneity in infectious disease data

#### Kamso Mohammed Mujaab (PhD, Biostatistics)

(Sept. 2018 - May 2024)

A novel semi-automated approach for trial identification and evaluation of the certainty of evidence from network meta-analyses Co-supervised with Glen Hazelwood, University of Calgary

# Chinmoy Rahul Roy (PhD, Biostatistics)

(Sept. 2019 – Sept. 2024)

Non-parametric spatial infectious disease models

#### Madeline Ward (PhD, Biostatistics)

(Sept 2020 – Jan 2025)

Bayesian epidemic models with mechanisms for behaviour change Co-supervised with Lorna Deeth, University of Guelph

# Tahmina Akter (PhD, Biostatistics)

(Sept. 2019 – March 2025)

Variable selection in infectious disease models

## 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, 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, 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, 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

### Jeffrey Peitsch (MSc student, Biostatistics)

(Sept 2023 – May 2025)

Directional spatial individual-level models of disease transmission Co-supervised with Gyanendra Pokharel, University of Winnipeg

## William Lee (URA)

(May 2016 - Dec. 2016)

Analysis of infectious disease surveillance data

# Madeline Ward (URA)

(May 2018 - Aug 2018)

Approaches to disease surveillance using predictive covariates Co-supervised with Lorna Deeth, University of Guelph

#### Arthur Novaes de Amorim (GRA)

(May 2019 – April 2020)

Predicting magnitude, timing, peak and duration of influenza at ER level Co-supervised with Vineet Saini, Alberta Health Services

### Zeyi Liu (GRA)

(March 2020 – Nov. 2020)

Spatial metapopulation models of Covid-19

### Emil Hodzic-Santor (URA)

(May 2022 – Aug 2022 & May 2023 – Aug 2023)

Edge effects in spatial epidemic models

#### Evans Mensah (GRA)

(Nov 2023 – June 2024)

Forecasting with behavioural-change population-level epidemic models

### Jensen MacLean (URA student)

(May 2024 – Aug. 2024 & May 2025 – Aug. 2025)

Patterns of behavioural-change in historic epidemic data