

# Ailene MacPherson

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## Positions

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<b>Assistant Professor</b> <i>Simon Fraser University</i> Infectious Disease Modelling and Estimation Canada Research Chair (Tier II) in Theoretical Evolutionary Epidemiology	2021-current
<b>PDF</b> <i>University of Toronto</i> EEB Departmental Postdoctoral fellow	2020-2021
<b>PDF</b> <i>University of British Columbia</i> Advisor: Matthew W. Pennell	2020-2020
<b>Ph.D. Zoology</b> <i>University of British Columbia</i> Thesis: Coevolutionary Epidemiology: a population genetic exploration of evolutionary interactions between hosts and their infectious pathogens Advisor: Sarah P. Otto	2015-2020
<b>MSc. Bioinformatics and Computational Biology</b> <i>University of Idaho</i> Thesis: Estimating the Strength of Natural Selection from Parallel Evolution Advisor: Scott L. Nuismer	2013-2015
<b>BSc. <i>summa cum laude</i> Mathematics</b> <i>University of Idaho</i>	2010-2013

## Publications<sup>1</sup>

### Peer-Reviewed Publications

- 1: Pennell, M.<sup>‡</sup>, **MacPherson, A.**<sup>‡</sup>, *Accepted*. Reading Yule in light of the history and present of macroevolution. *Phil. Trans. B*.
- 2: Otto, S.P., **MacPherson, A.**, Colijn, C., 2024. Endemic does not mean constant as SARS-CoV-2 continues to evolve. *Evolution*
- 3: Davies, T.J., **MacPherson, A.**, 2024. Seed masting as a mechanism for escape from pathogens. *Current Biology*
- 4: Urquhart-Cronish M.<sup>†</sup>, Angert, A.L., Otto, S.P. **MacPherson, A.**, 2023. Density-dependent selection during range expansion affects expansion load in life-history traits. *The American Naturalist*
- 5: Clancy, E., **MacPherson, A.**, Cheek, R.G., Mouton, J.C., Sillett T.S., Chalambor, C.K., Funk, W.C., Hohenlohe, P.A., 2023. Unraveling Adaptive Evolutionary Divergence at Microgeographic Scales. *The American Naturalist*
- 6: Peters, M.A.E.<sup>†</sup>, Mideo, N., **MacPherson, A.**, 2023. The maintenance of genetic diversity under host-parasite coevolution in finite, structured populations. *J. Evol. Biol.*
- 7: Liu, P., Song, Y.\*<sup>‡</sup>, Colijn, C., **MacPherson, A.**, 2022. The impact of sampling bias on viral phylogeographic reconstruction. *PLoS Global Public Health*
- 8: **MacPherson, A.**<sup>‡</sup>, Wang, S.<sup>‡</sup>, Yamaguchi, R., Rieseberg, L.H, Otto, S.P., 2022. Parental population range expansion before secondary contact promotes heterosis. *The American Naturalist*
- 9: Wang S., Nalley M.J., Chatla K., Aldaimalani R., **MacPherson A.**, Wei K., Corbett R., Mai D., Bachtrog D., 2022. Neo-sex chromosome evolution shapes sex-dependent asymmetrical introgression barrier. *PNAS*
- 10: **MacPherson, A.**, Louca, S., McLaughlin A.<sup>†</sup>, Joy, J.B., Pennell, M.W., 2021. Unifying Phylogenetic Birth-Death Models in Epidemiology and Macroevolution. *Syst. Biol.*
- 11: Louca, S., McLaughlin A.<sup>†</sup>, **MacPherson, A.**, Joy, J.B., Pennell, M.W., 2021. Fundamental identifiability limits in molecular epidemiology. *Mol. Biol. Evol.*

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<sup>1</sup>\* Indicates a trainee in my group. <sup>†</sup> Indicates a trainee in another research group. <sup>‡</sup> Indicates equal contribution authors.

- 12: MacPherson, A.,** Keeling M.J., Otto, S.P., 2021. Feedback between coevolution and epidemiology can help or hinder the maintenance of genetic variation in host-parasite models. *Evolution*
- 13: MacPherson, A.,** Keeling M.J., Otto, S.P., 2020. Coevolution fails to maintain genetic variation in a host-parasite model with constant finite population size. *Theor. Popul. Biol.*
- 14: MacPherson, A.,** Otto, S.P., Nuismer, S.L., 2018. Keeping pace with the Red Queen: Identifying the genetic basis of susceptibility to infectious disease. *Genetics*
- 15: MacPherson, A.,** Otto, S.P., 2018. Joint coevolutionary-epidemiological models dampen Red Queen cycles and alter conditions for epidemics. *Theor. Popul. Biol.*
- 16: MacPherson, A.,** Nuismer, S.L., 2017. The probability of parallel genetic evolution from standing genetic variation. *J. Evol. Biol.*
- 17: MacPherson, A.,** Hohenlohe, P.A., Nuismer, S.L., 2015. Trait dimensionality explains widespread variation in local adaptation. *Proc. R. Soc. Lond. B.*
- 18: Nuismer, S.L., MacPherson, A.,** Rosenblum, E.B., 2012. Crossing the threshold: gene-flow, dominance and the critical level of standing genetic variation required for adaptation to novel environments. *J. Evol. Biol.*
- 19: Balemba, O.B.,** Stenkamp-Strahm, C.H., Cady, J., **MacPherson, A.,** 2011. High-fat diet-induced neuropathy of enteric nervous system and the effect of Alpha-7 Nicotinic Acetylcholine receptor agonist, DMAB-Anabaseine Dihydrochloride. *Gastroenterology*

## Pre-prints and In Review

- 20: MacPherson, A.<sup>‡</sup>,** Bazzicalupo, A.<sup>‡</sup>, in review. A Unified Framework for Studying Local Adaptation.
- 21: Hall, R.A.\*,** **MacPherson, A.,** in review. Local Adaptation of Life-History Traits in a Seasonal Environment.
- 22: MacPherson, A.,** Pennell, M., in review. The Untapped Potential of Tree Size in Reconstructing Evolutionary and Epidemiological Dynamics. <https://doi.org/10.1101/2024.06.07.597929>
- 23: Salehzadeh, M.\*,** Stockie, J.M. , **MacPherson, A.,** in review. Aggregation Unveiled: A Sequential Modelling Approach to Bark Beetle Outbreaks.
- 24: Bajgai, A.\*,** Elliot, L.T. , **MacPherson, A.,** in review. Disease as a conservation threat is non-randomly distributed among animal taxa: an examination with a large language model.

## Funding and Awards

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<i>James F. Crow Early Career Award</i> Genetic Society of America (2022)	500 USD
<i>Tier II Canada Research Chair [CRC-2021-00276]</i> (2022-2027)	120,000 CAD/Year
<i>NSERC Discovery Grant [RGPIN-2022-03113]</i> (2022-2027)	29,000 CAD/Year
<i>NSERC Launch Supplement [DGEGR-2022-00326]</i> (2021)	12,000 CAD
<i>EEB Postdoctoral Fellowship</i> University of Toronto (2020)	55,000 CAD/Year
<i>Graduate Fellowship</i> American Association of University Women (2019)	20,000 USD
<i>Zoology Graduate Fellowship</i> University of British Columbia Zoology (2019)	15,000 CAD (Declined)
<i>Godfrey Hewitt Mobility Award</i> European Society of Evolutionary Biology (2018)	1,500 Euro
<i>Zoology Four-Year Fellowship</i> University of British Columbia Zoology (2015-2018)	15,000 CAD/Year
<i>BCB Graduate Research Fellowship</i> University of Idaho (2013-2015)	21,000 USD/Year
<i>Outstanding Senior in Mathematics</i> University of Idaho (2015)	500 USD
<i>College of Science Dean's Award</i> University of Idaho (2015)	200 USD

## Research Dissemination

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### Invited Talks:

*Seminar: Diversification Models for Epidemiological, Paleontological, and Macroevo-lutionary Applications.*  
ETH Zurich, Basel, Switzerland (October 2023)

*Seminar: Eco-Evo-lutionary Implications of Range Expansion.*  
Univ. Bern, Bern, Switzerland (October 2023)

*Workshop: Phylodynamic Inference across Micro and Macroevo-lutionary Scales.*  
Taming The BEAST 2023, Squamish, BC (August 2023)

*Conference: Eco-Evo-lutionary implications of range expansion.*  
ICG2023, Melbourne, Aus. (July 2023)

*Workshop: Demography, epidemiology, and geography determine the maintenance of diversity.*  
BIRS-Banff, (January 2023)

*Seminar: Re-envisioning the geographic mosaic of coevolution.*  
SFU-Biology, (November 2022)

*Workshop: Epidemiological Inference with phylogenies.*  
BIRS-CSMO, Oaxaca, Mex. (November 2022)

*Seminar: Inferring epidemiological parameters from viral phylogenies.*  
SFU Physics, (October 2022)

*Workshop: Do host-parasite interactions maintain immune diversity?.*  
IMMREP, Germany (Virtual) (July 2022)

*Workshop: Allele surfing: hybridization and the evolution of life-history traits.*  
Fields Institute Workshop on Range Dynamics, Virtual (June 2022)

*Talk: Parental population expansion promotes heterosis in secondary contact hybrid zones.*  
PEQG22, Pacific Grove, CA (June 2022)

*Workshop: Eco-evolutionary maintenance of genetic variation.*  
Eco-evolutionary modelling, Paris, France (Virtual) (March 2022)

*Talk: Limitations to inference in a general phylodynamic model.*  
AMS Special Session, Seattle, WA (Virtual) (January 2022)

*Seminar: Coevolution between hosts and their infectious pathogens.*  
Max Plank Institute, Plön, Germany (Virtual) (May, 2020)

*Workshop: Identifying the genetic basis of coevolution.*  
ESEB-STN, Freising, Germany (March, 2019)

### Conferences Talks:

*Expansion load in density-dependent life-history traits.* Evo-WIBO, (2023)

*Coevolution does not maintain genetic variation.* ASN, (2020)

*Finding disease genes in the face of the Red Queen.* Evolution, (2017)

*Epidemiological dynamics disrupt Red Queen cycles.* CSEE, (2017)

*Natural selection and probability of parallel evolution.* Evolution, (2015)

*Trait dimensionality and local adaptation.* Evo-WIBO, (2014)

## Students and Thesis Committees

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### Graduate Students/Postdoctoral Fellows:

Oliver Fijiki (Sep. 2023–current) MSc. (Coadvised with Alexander Rutherford)

Siavash Riazi (Jan. 2023–current) PDF (Coadvised with Caroline Colijn)

Mahdi Salehzadeh (Jan 2023–current ) Ph.D.(Coadvised with John Stockie)

Rebekah Hall (Sep. 2022–Aug. 2024) MSc., *Thesis*: Life-history evolution under seasonal disruptions

Yexuan Song (Sep 2022–Current) PhD. (Coadvised with Caroline Colijn)

Yexuan Song (Jan 2022–Aug 2022) MSc. (Coadvised with Caroline Colijn), *Thesis*: Accounting for sampling bias in ancestral state reconstruction

**Undergraduate Researchers/Research Assistants:**

Simon Levi Gamboa (2024–current) RA  
 Oliver Fijiki (2022–2023) USRA & RA  
 Ananga Bajgai (2023–current) USRA & RA

**Graduate Student Committees:**

Zahresh Walji (2023–current) MSc.  
 Daniel Pelletier (2022–current) MSc.  
 Mina Moeini (2021–2024) MSc.  
 Rituparna Banerjee (2022–current) PhD

**Teaching**

2024\*: *Topics in Biomathematics*: Professor. SFU.  
 2023\*: *Mathematics of Evolution*: Professor. SFU.  
 2022\*: *Mathematical Epidemiology*: Professor. SFU.  
 2021\*, 2022: *Mathematics for The Life Sciences*: Professor. SFU.

\*courses developed including syllabi, lectures, & assignments

**Professional Activities & Service**

<b>Peer Reviews:</b>	Am. Nat., Evolution, Genetics, Heredity, J. Mol. Evol. , J. Theor. Biol. , Phil. Trans. R. Soc. B, PNAS, PLoS Genet., Proc. R. Soc. Lond. B, Virus Evol., TREE, Trends Genet., Syst. Biol.
<b>Editor Boards</b>	Theor. Pop. Biol. (Associate Editor)
<b>Grant Reviews:</b>	NSERC DG
<b>Book Reviews:</b>	Springer
<b>Thesis Examiner</b>	Mario Santer, Ph.D. Thesis, Max Plank Institute, Nicola Mulberry, Ph.D. SFU
<b>Working Groups/Networks:</b>	PIMS-Maud Menton Institute Co-applicant (2024) BIRS: Modeling and Theory in Population Biology (2024) UBC Biodiversity: Linking Macroevolution and Epidemiology (2020)
<b>Society Service:</b>	SMTPB–founding member & SMTPB–Secretary (2023-2024)
<b>University Service:</b>	Chair Nominating Committee (2023) & GSC (2022–current) & TPC (2022-23, 2024–current) & Hiring Committee (2022)
<b>Outreach:</b>	Café Scientifique University of Toronto-EEB Modelling Club Managing Fisheries with Math–Grade 12 <i>Palmer H.S., Palmer, Alaska</i> Researchers Revealed Video Series: “Counting on it” Let’s Talk Science-Volunteer Idaho Women in STEM K-12 Program-Volunteer
<b>Media</b>	BCIT News 29/9/21 Citynews Vancouver 27/12/21