
Publication List
(Research Group Members Shown in Italics)

Submitted Articles

106. **C. Ward, R. Deardon** & A. Schmidt “Multivariable behavioural change modelling of epidemics in the presence of undetected infections” submitted to *Statistics in Medicine* (revision requested). <https://arxiv.org/abs/2503.00982>
 105. **M. Mahsin, W. Almutiry & R. Deardon** “Spatial modeling of infectious disease transmission using continuous time geographically-dependent individual-level models” submitted to *Statistics in Medicine* (revision requested).
 104. **D. Douwes-Schultz, R. Deardon** & A. Schmidt “Hidden Markov individual-level models of infectious disease transmission” submitted to the *Journal of the American Statistical Association*.
 103. **M. Ward, R. Deardon, L. Deeth, C. Ward** “Bayesian epidemic models with dynamic adherence to protective behaviour changes” submitted to the *Journal of the Royal Statistical Society: Series C*.
 102. **Y. Mao, R. Deardon** & L. Deeth “Identifying memory mechanisms in Bayesian models of behavioural change during epidemics” submitted to *Annals of Applied Statistics*
 101. **Y. Mao, R. Deardon** & L. Deeth “Memory mechanisms for behavioural change in Bayesian individual level spatial epidemic models” submitted to *Infectious Disease Modelling*
 100. **Y. Zhang, R. Deardon** & L. Deeth “Behavioural change model choice in spatial epidemic models” submitted to *Canadian Journal of Statistics*.
 99. **J. MacLean & R. Deardon** “Behavioural change in Canadian disease outbreaks” submitted to the *Canadian Journal of Statistics*.
 98. **R. Li, R. Deardon**, N. Li, J. Conly & J. Leal “Bayesian compartmental modelling of the transmission dynamics of methicillin-resistant *Staphylococcus aureus* (MRSA) within hospitals in Edmonton, Canada” submitted to the *Canadian Journal of Statistics* (revision requested). <http://arxiv.org/abs/2511.07353>
 97. **R. Li, R. Deardon**, N. Li, J. Conly & J. Leal “Modelling the effect of interventions to prevent the transmission of methicillin-resistant *Staphylococcus aureus* (MRSA) within hospitals in Edmonton, Canada” submitted to *Infectious Disease Modelling* (revision requested).
 96. L. G. Salazar, H. McKenzie, **R. Deardon**, K. Pepin, R. Brook, J. Bahamon, C. Neva & M. Pruvot “Interaction risk between wild pigs and livestock: implications for potential infectious disease transmission in Alberta, Canada” submitted to *Preventive Veterinary Medicine*.
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Accepted/In Press

95. V. Callier, **R. Deardon** & C. Viboud (2026) “Spatio-temporal spread of COVID-19 over three waves in the continental United States” to appear in the *Proceedings of the Royal Society B*.
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Published Articles

94. **Y. Zhang, R. Deardon** & L. Deeth (2026) “Composite method for fast computation of individual level spatial epidemic models” in *Spatial Statistics*, 72, 100957. <https://doi.org/10.1016/j.spasta.2026.100957>
93. **J. Peitsch**, G. Pokharel & **R. Deardon** (2026) “Directional spatial individual level models of infectious disease transmission” in *Spatial Statistics*, 73, 100965.
92. **M. Kamso**, S. Whittle, J. Pardo Pardo, R. Buchbinder, G. Wells, **R. Deardon**, T. Sajobi, G. Tomlinson, E. Jesse, J. Thomas, S. Kelly, G. Hazlewood (2026) “A semi-automated approach facilitated the assessment of the certainty of evidence in a network meta-analysis: Part 1 – Direct comparisons” in the *Journal of Clinical Epidemiology*, 191, 112109. <https://doi.org/10.1016/j.jclinepi.2025.112109>.
91. **M. Kamso**, S. Whittle, J. Pardo Pardo, R. Buchbinder, G. Wells, **R. Deardon**, T. Sajobi, G. Tomlinson, E. Jesse, J. Thomas, S. Kelly, G. Hazlewood (2026) “A semi-automated approach facilitated the assessment of the certainty of evidence for in a network meta-analysis: Part 2 – indirect and mixed comparisons” in the *Journal of Clinical Epidemiology*, 191, 112110. <https://doi.org/10.1016/j.jclinepi.2025.112110>.
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90. H. Qureshi, T. Hughes, E. Franco, K. Fiest, J. Gratrix, P. Smyczek, R. Read, A. Afzal, **R. Deardon**, A. Kassam & M. Fidler-Benaoudia (2026) “Risk of cancer among individuals with a history of bacterial sexually transmitted infections: a population-based study in Alberta, Canada” available online in *International Journal of Cancer*, 158(5), 1383-1395. <http://doi.org/10.1002/ijc.70215>
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89. **M. Ward, R. Deardon** & L. Deeth (2025) “A framework for incorporating behavioural change into individual-level spatial epidemic models” in the *Canadian Journal of Statistics*, 53(1), e11828. <https://doi.org/10.1002/cjs.11828>
88. **T. Akter & R. Deardon** (2025) “Conditional logistic individual-level models of spatial infectious disease dynamics” in *Infectious Disease Modelling*, 10(1), 268-286. <https://doi.org/10.1016/j.idm.2024.10.008>
87. **C. Rahul & R. Deardon** (2025) “Behavioural change piecewise constant spatial epidemic models” in *Infectious Disease Modelling*, 10(1), 302-324. <https://doi.org/10.1016/j.idm.2024.10.006>
86. **T. Akter & R. Deardon** (2025) “Variable screening methods in conditional logistic individual level models of disease spread” in *Spatial & Spatiotemporal Epidemiology*, 54, 100742. <https://doi.org/10.1016/j.sste.2025.100742>
85. M. Lewis, P. Brown, C. Colijn, L. Cowen, C. Cotton, T. Day, **R. Deardon**, D. Earn, D. Haskell, J. Heffernan, P. Leighton, K. Murty, S. Otto, E. Rafferty, C. Hughes Tuohy, J. Wu & H. Zhu “Charting a future for emerging infectious disease modelling in Canada” (2025) in *Lasting Disruption: Economic and Social Impacts of COVID-19 in Canada*, McGill-Queen’s University Press. (Ed: Christopher Cotton). <http://hdl.handle.net/1828/15042>.
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84. **C. Rahul & R. Deardon** (2024) “Individual-level models of disease transmission incorporating non-parametric spatial risk” in *Spatial & Spatiotemporal Epidemiology*, 50, 100664. <https://doi.org/10.1016/j.sste.2024.100664>
83. **E. Hodzic-Santor & R. Deardon** (2024) “Edge effects in spatial infectious disease models” in *Spatial & Spatiotemporal Epidemiology*, 50, 100673. <https://doi.org/10.1016/j.sste.2024.100673>
82. M. Biesheuvel, **C. Ward**, P. Penterman, E. van Engelen, G. Schaik, **R. Deardon** & H. Barkema (2024) “Within-herd transmission of *Mycoplasma bovis* infection in 20 Dutch dairy herds” in *Journal of Dairy Science*, 107(1), 503-516. <https://doi.org/10.3168/jds.2023-23407>
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81. **C. Ward, R. Deardon** & A. Schmidt (2023) “Bayesian modelling of dynamic behavioural change during an epidemic” *Infectious Disease Modelling*, 8(4), 947-963. <https://doi.org/10.1016/j.idm.2023.08.002>
80. **L. Amiri**, M. Torabi & **R. Deardon** (2023) “Spatial modelling of infectious diseases with covariate measurement error” in *Journal of the Royal Statistical Society: Series C*, 73(2), 460-477. <https://doi.org/10.1093/rssc/qlad104>
79. **L. Amiri**, M. Torabi & **R. Deardon** (2023) “Analyzing COVID-19 data in the Canadian Province of Manitoba: A new approach” in *Spatial Statistics*, 55:100729. doi: 10.1016/j.spasta.2023.100729.
78. **T. Akter** & **R. Deardon** (2023) “Comparison of variable screening methods in infectious disease transmission models” in *Spatial and Spatiotemporal Epidemiology*, 47, 100622.
77. **M. Kamso**, J. Pardo, S. Whittle, R. Buchbinder, G. Wells, V. Glennon, P. Tugwell, **R. Deardon**, T. Sajobi, G. Tomlinson, J. Elliot, S. Kelly & G. Hazlewood (2023). “Crowdsourcing and automation facilitated the identification and classification of randomized controlled trials in a living review” in *Journal of Clinical Epidemiology*, 164, 1-8. <https://doi.org/10.1016/j.jclinepi.2023.10.007>
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75. **M. Mahsin**, **R. Deardon** & P. Brown (2022) “Geographically-dependent individual-level models for infectious diseases transmission” in *Biostatistics*, 23(1), 1-17. <https://doi.org/10.1093/biostatistics/kxaa009>
74. **J. Angevaare**, Z. Feng & **R. Deardon** (2022) “Pathogen.jl: Infectious disease transmission network modelling with Julia” in *Journal of Statistical Software*, 104(4), 1?30.
73. **G. Pokharel** & **R. Deardon** (2022) “Emulation-based inference for spatial infectious disease transmission models incorporating event time uncertainty” in the *Scandinavian Journal of Statistics*, 49(1), 455-479. <http://doi.org/10.1111/sjos.12523>
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71. **S. A. Naqvi**, M. King, T. DeVries, H. Barkema & **R. Deardon** (2022) “Data considerations for developing deep learning models for dairy applications” in *Computers and Electronics in Agriculture*, 196: 106895. <https://doi.org/10.1016/j.compag.2022.106895>
70. **S. A. Naqvi**, M. King, R. Matson, T. DeVries, **R. Deardon** & H. Barkema (2022) “Mastitis detection with recurrent neural networks in farms using automated milking systems” in *Computers and Electronics in Agriculture*, 192: 106618. <https://doi.org/10.1016/j.compag.2021.106618>
69. **B. Jafari** & **R. Deardon** (2022) “Bias and Bias-Correction for Individual-Level Models of Infectious Disease” in *Spatial & Spatiotemporal Epidemiology*, 43, 100524.
68. J. Di Francesco, G.P.S. Kwong, **R. Deardon**, S. L. Checkley, G. F. Mastromonaco, F. Mavrot, L. Leclerc & S. Kutz (2022) “Intrinsic and extrinsic factors associated with increased qiviut cortisol in wild muskoxen (*Ovibos moschatus*)” in *Conservation Physiology*, 10(1), coab103. <https://doi.org/10.1093/conphys/coab103>
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64. **W. Almutiry** & **R. Deardon** (2021) “Contact network uncertainty in individual level models of infectious disease transmission” in *Statistical Communications in Infectious Diseases*, 13(1). DOI: <https://doi.org/10.1515/scid-2019-0012>
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60. B. Singh, **M. Lowerison**, R. Lewinson, I. Vallerand, **R. Deardon**, J. Gill, B. Singh & H. Barkema (2021) “Public health interventions slowed but did not halt the spread of COVID-19 in India” in *Transboundary and Emerging Diseases*, 68(4), 2171-2187. <https://doi.org/10.1111/tbed.13868>
59. C. Doolan, T. Louie, C. Lata, O. Larios, W. Stokes, J. Kim, K. Brown, P. Beck, **R. Deardon** & D. Pillai (2021) “Latent class analysis for the diagnosis of Clostridioides difficile infection” in *Clinical Infectious Diseases*, 73(9):e2673-e2679. <https://doi.org/10.1093/cid/ciaa1553>
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57. **W. Almutiry** & **R. Deardon** (2020) “Incorporating contact network uncertainty in individual level models of infectious disease using approximate Bayesian computation” in *The International Journal of Biostatistics*, 16(1), Article 20170092. DOI: <https://doi.org/10.1515/ijb-2017-0092>
56. **V. Warriyar**, **W. Almutiry** & **R. Deardon** (2020) “Individual level modelling of infectious disease data: EpiILM” in *The R Journal* 12(1), 199-217.
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48. **C. Augusta**, **R. Deardon** & G. Taylor (2019) “Deep learning for supervised classification of spatial epidemics” in *Spatial & Spatiotemporal Epidemiology*, 29, 187-198.
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40. **G. Pokharel & R. Deardon** (2018) “Spatially informed back-calculation for spatio-temporal infectious disease models” in *Statistical Communications in Infectious Diseases*, Vol. 10(1), Article 2.
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38. D. Toms, **R. Deardon** & M. Ungrin (2017) “Climbing the mountain: Experimental design for efficient optimization of stem cell bioprocessing” in the *Journal of Biological Engineering*, Vol. 11, No. 1
37. **R. Romanescu & R. Deardon** (2017) “Fast inference for network models of infectious disease spread” in the *Scandinavian Journal of Statistics*, 44(3), 666-683 (DOI: 10.1111/sjos.12270).
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36. **G. Pokharel & R. Deardon** (2016) “Gaussian process emulators for spatial models of infectious disease” in the *Canadian Journal of Statistics*, 44(4), 480-501.
35. **R. Romanescu & R. Deardon** (2016) “Modelling two strains of disease via aggregate-level infectivity curves” in the *Journal of Mathematical Biology*, 72(5), 1195-1224.
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16. J. Gallienne, C. Gregg, E. LeBlanc, N. Yaakob, D. Wu, K. Davies, N. Rawlings, Pierson, **R. Deardon**, & Bartlewski “Correlations between ultrasonographic characteristics of corpora lutea (CL) and systemic concentrations of progesterone (P4) during the discrete stages of CL lifespan and secretory activity in cyclic ewes” in *Experimental Biology and Medicine*, 237, 505 – 515.
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12. **R. Deardon**, S. P. Brooks, B. T. Grenfell, M. J. Keeling, M. J. Tildesley, N. J. Savill, D. J. Shaw & M. E. J. Woolhouse (2010), “Inference for individual-level models of infectious diseases in large populations” in *Statistica Sinica*, 20(1), 239-261. (Funded by: Wellcome Trust, UK).
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