

---

**Publication List**  
(*Research Group Members Shown in Italics*)

---

**Submitted Articles**

90. **C. Rahul** & **R. Deardon** “Behavioural change piecewise constant spatial epidemic models” submitted to *Infectious Disease Modelling*
  89. **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*.
  88. 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” submitted to *Facets*.  
<http://hdl.handle.net/1828/15042>.
  87. **E. Hodzic-Santor** & **R. Deardon** “Edge effects in spatial infectious disease models” submitted to *Spatial & Spatiotemporal Epidemiology* (revision requested).
  86. **M. Kamso**, S. Whittle, J. Pardo, R. Buchbinder, G. Wells, R. Johnston, V. Glennon, P. Tugwell, **R. Deardon** T. Sajobi, G. Tomlinson, J. Elliot, J. Thomas, S. Kelly & G. Hazlewood. “A semi-automated approach facilitated the assessment of the certainty of evidence for direct comparisons in network meta-analyses” submitted to the *Journal of Clinical Epidemiology*.
  85. **M. Pasha**, **R. Deardon** & A. Rahim “Multi-response and multi-cause process monitoring by applying proportional hazards models in the optimal design of  $T^2$  control charts” submitted to *Computers & Industrial Engineering*.
  84. **M. Pasha**, **R. Deardon** & A. Rahim “Multi-response process monitoring with  $T^2$  control charts under multiple assignable causes” submitted to *Quality Technology and Quantitative Management*.
- 

**Accepted/In Press**

83. **M. Ward**, **R. Deardon**, L. Deeth (2024) “A framework for incorporating behavioural change into individual-level spatial epidemic models” to appear in the *Canadian Journal of Statistics*.  
<http://arxiv.org/abs/2308.00815>
  82. **C. Rahul** & **R. Deardon** “Individual-level models of disease transmission incorporating non-parametric spatial risk” to appear in the *Spatial & Spatiotemporal Epidemiology*.  
<http://arxiv.org/abs/2405.00835>
- 

**Published Articles**

81. 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>
-

80. **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>
  79. **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*, qlad104 <https://doi.org/10.1093/jrsssc/qlad104>
  78. **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.
  77. **T. Akter** & **R. Deardon** (2023) “Comparison of variable screening methods in infectious disease transmission models” in *Spatial and Spatiotemporal Epidemiology*, 47, 100622.
  76. **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>
  75. **M. Pasha**, **R. Deardon** & A. Rahim (2023) “A study on inspection schemes in optimal design of control charts for deteriorating processes” in *Quality and Reliability Engineering International*, 39(3), 732-751. <https://doi.org/10.1002/qre.3253>
- 
74. **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>
  73. **J. Angevaere**, Z. Feng & **R. Deardon** (2022) “Pathogen.jl: Infectious disease transmission network modelling with Julia” in *Journal of Statistical Software*, 104(4), 1?30.
  72. **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>
  71. **M. Ward**, L. Deeth & **R. Deardon** (2022) “Cluster-aggregation-disaggregation methods for spatial individual level models of infectious disease transmission” in *Spatial & Spatiotemporal Epidemiology*, 41: 100497. <https://doi.org/10.1016/j.sste.2022.100497>
  70. **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>
  69. **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>
  68. **B. Jafari** & **R. Deardon** (2022) “Bias and Bias-Correction for Individual-Level Models of Infectious Disease” in *Spatial & Spatiotemporal Epidemiology*, 43, 100524.
  67. 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>
- 
66. **W. Almutiry**, **V. Warriyar** & **R. Deardon** (2021) “Continuous-time individual-level models of infectious disease: EpiILMCT” in the *Journal of Statistical Software*, 98(10), 1-44. <https://www.jstatsoft.org/article/view/v098i10>

65. **L. Amiri**, M. Torabi, **R. Deardon** & M. Pickles (2021). “Spatial modeling of individual-level infectious disease transmission: tuberculosis data in Manitoba, Canada” in *Statistics in Medicine*, 40(7), 1678-1704. <https://doi.org/10.1002/sim.8863>
  64. **J. Angevaere**, Z. Feng & **R. Deardon** (2021) “Inference of latent event times and transmission network in individual level infectious disease models” in *Spatial & Spatiotemporal Epidemiology*, 37, 100410. <https://doi.org/10.1016/j.sste.2021.100410>
  63. **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>
  62. **Z. Liu**, **R. Deardon**, Y. Fu, **T. Ferdous**, T. Ware & Q. Cheng (2021) “Estimating parameters of two-level individual-level models of the COVID-19 epidemic using ensemble learning classifiers” in *Frontiers in Physics*, 8(11), Article 602722. doi: 10.3389/fphy.2020.602722
  61. **A. Novaes de Amorim**, V. Saini & **R. Deardon** (2021) “A stacked ensemble method for forecasting influenza-like illness visit volumes at emergency departments” in *PLOS One*, 16(3): e0241725. <https://doi.org/10.1371/journal.pone.0241725>
  60. S. Andres-Lasheras, R. Ha, R. Zaheer, C. Lee, C. Booker, C. Dorin, J. Van Donkersgoed, **R. Deardon**, S. Gow, S. Hannon, S. Hendrick, M. Anholt & T. McAllister (2021) “Prevalence and risk factors associated with antimicrobial resistance in bacteria related to bovine respiratory disease - A broad cross-sectional study of beef cattle at entry into Canadian feedlots” in *Frontiers in Veterinary Science*, 8, 710. doi: 10.3389/fvets.2021.692646
  59. 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>
  58. 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>
  57. B. Singh, M. Ward, **M. Lowerison**, R. Lewinson, I. Vallerand, **R. Deardon**, J. Gill, B. Singh & H. Barkema (2021) “Meta-analysis and adjusted estimation of COVID-19 case fatality risk in India and its association with the underlying comorbidities” in *One Health*, 13:100283. <https://doi.org/10.1016/j.onehlt.2021.100283>
- 
56. **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>
  55. **V. Warriyar**, **W. Almutiry** & **R. Deardon** (2020) “Individual level modelling of infectious disease data: EpiILM” in *The R Journal* 12(1), 199-217.
  54. G. Hazelwood, **G. Pokharel**, **R. Deardon**, D. Marshall, C. Bombardier, G. Tomlinson, C. Ma, C. Seow, R. Panaccione & G. Kaplan (2020) “Patient preferences for maintenance therapy in Crohn’s disease: a discrete-choice experiment” in *PLoS One*, 15(1):e0227635.
  53. G.P.S. Kwong, **R. Deardon**, **S. Hunt** & M. Guerin (2020) “Bayesian optimal design of agricultural infectious disease transmission experiments” available online in *Statistical Communications in Infectious Diseases*, 12(1). <https://doi.org/10.1515/scid-2018-0005>
  52. **R. Romanescu** & **R. Deardon** (2020) “Implementation of power law network models of epidemic surveillance data for better evaluation of outbreak detection alarms” in *Statistical Communications in Infectious Diseases*, 12(1). <https://doi.org/10.1515/scid-2018-0004>.

51. D. Nobrega, **S. A. Naqvi**, S. Dufour, **R. Deardon**, J. Kastelic, J. de Buck & H. Barkema (2020) "Critically important antimicrobials are not needed to treat non-severe clinical mastitis in lactating dairy cows: results from a network meta-analysis" in the *Journal of Dairy Science*, 103(11), 10585-10603. <https://doi.org/10.3168/jds.2020-18365>
  50. **G. Pokharel**, **R. Deardon**, S. Johnson, G. Tomlinson, P. Hull, G. Hazelwood (2020) "Effectiveness of initial methotrexate-based treatment approaches in early rheumatoid arthritis: An elicitation of rheumatologists' beliefs" in *Rheumatology*, keaa803. <https://doi.org/10.1093/rheumatology/keaa803>
  49. A. Ogilvy, S. Collins, T. Tuokko, M. Hiltz, **R. Deardon**, W. Hare & A. Jirasek (2020) "Optimization of solid tank design for fan-beam optical CT based 3D radiation dosimetry" in *Physics in Medicine & Biology*. 65, 245012. <https://doi.org/10.1088/1361-6560/abbf98>
- 
48. **C. Augusta**, **R. Deardon** & G. Taylor (2019) "Deep learning for supervised classification of spatial epidemics" in *Spatial & Spatiotemporal Epidemiology*, 29, 187-198.
  47. **M. Ward**, **A. Stanley**, L. Deeth **R. Deardon**, Z. Feng & L. Trotz-Williams (2019) "Methods for detecting seasonal influenza epidemics using a school absenteeism surveillance system" in *BMC Public Health*, 19, Article: 1232.
  46. **C. Augusta**, G. Taylor & **R. Deardon** (2019) "Dynamic contact networks of swine movement in Manitoba, Canada: characterization and implications for infectious disease spread" in *Trans-boundary and Emerging Diseases*, 66(6), 1910 - 1919. DOI: <https://doi.org/10.1111/tbed.13220>.
  45. **G. Pokharel**, **R. Deardon**, C. Barnabe, V. Bykerk, S. Bartlett, L. Bessette, G. Boire, C. Hitchon, E. Keystone, J. Pope, O. Schieer, D. Tin, C. Thorne & G. Hazelwood (2019) "Joint estimation of remission and response for methotrexate-based DMARD options in rheumatoid arthritis: A bivariate network meta-analysis" in *ACR Open Rheumatology*, 1(8), 471-479. <https://onlinelibrary.wiley.com/doi/epdf/10.1002/acr2.11052>.
  44. **M. Lowerison**, C. Josephson, N. Jette, T. Sajobi, S. Patten, T. Williamson, **R. Deardon**, H. Barkema, & S. Wiebe (2019) "Association of levels of specialized care with risk of premature mortality in patients with epilepsy" in *JAMA Neurology*, 76(11), 1352-1358. DOI: <https://doi.org/10.1001/jamaneurol.2019.2268>
  43. S. Coward, F. Clement, E. Benchimol, C. Bernstein, J. Antonio Avina-Zubieta, A. Bitton, M. Carroll, G. Hazelwood, K. Jacobson, S. Jelinski, **R. Deardon**, J. Jones, M. Ellen Kuenzig, D. Leddin, K. McBrien, S. Murphy, G. Nguyen, A. Otley, R. Pannaccione, A. Rezaie, G. Rosenfeld, J. Pena-Sanchez, H. Singh, L. Targownik, G. Kaplan (2019) "Past and future burden of inflammatory bowel diseases based on modeling of population-based data" in *Gastroenterology*, 156(5), 1345-1353.
  42. J. Cheaveau, D. Marasinghe, S. Akakpo, **R. Deardon**, C. Naugler, A. Chin, D. R. Pillai (2019) "The impact of malaria on liver enzymes: a retrospective cohort study (2010-2017)" in *Open Forum Infectious Diseases*, 6(6).
- 
41. T. Petukhova, D. Ojkic, B. McEwen, **R. Deardon** & Z. Poljak (2018) "Assessment of ARIMA, GLARMA and random forest models for predicting Influenza A virus frequency in swine in Ontario, Canada" in *PLoS One*, 13(6): e0198313.
  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.
  39. M. Lipson, **R. Deardon**, N. Switzer, C. DeGara, C. Ball & S. Grondin (2018) "Practice and attitudes regarding double gloving among staff surgeons and surgical trainees" in the *Canadian Journal of Surgery*, 61(4), 244-250.
-

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).
- 
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.
  34. **L. Deeth** & **R. Deardon** (2016) “Spatial data aggregation for spatio-temporal individual-level models of infectious disease transmission” in *Spatial & Spatio-temporal Epidemiology*, 17, 95-104.
  33. **R. Malik**, **R. Deardon** & **G.P.S. Kwong** (2016) “Parameterizing spatial models of infectious disease spread using sampling-based likelihood approximations” in *PLoS One*, 11(1): e0146253. doi: 10.1371/journal.pone.0146253.
- 
32. **L. Deeth**, **R. Deardon** & D. Gillis (2015) “Model choice using the Deviance Information Criterion for latent conditional individual-level models of infectious disease spread” in *Epidemiologic Methods*, 4(1), 47-68.
  31. **R. Deardon**, **X. Fang** & **G.P.S. Kwong** (2015) “Statistical modelling of spatio-temporal infectious disease transmission” in *Analyzing and Modeling Spatial and Temporal Dynamics of Infectious Diseases*, 211-232, John Wiley & Sons. (Ed: D. Chen, B. Moulin, J. Wu).
- 
30. T. J. McKinley, J. Ross, **R. Deardon** & A. Cook (2014) “Simulation-based Bayesian inference for epidemic models” in *Computational Statistics & Data Analysis*, 71, 434-447.
  29. **R. Malik**, **R. Deardon**, **G.P.S. Kwong** & B. J. Cowling (2014) “Individual-level modeling of the spread of influenza within households” in *Journal of Applied Statistics*, 41(7), 1578-1592.
  28. **G. Pokharel** & **R. Deardon** (2014) “Supervised learning and prediction of spatial epidemics” in *Spatial & Spatio-Temporal Epidemiology*, 11, 59-77.
- 
27. **L. Deeth** & **R. Deardon** (2013) “Latent conditional individual level models for infectious disease modelling” in *The International Journal of Biostatistics*, 9(1), 75-93.
  26. **S. Subedi**, Z. Feng, **R. Deardon** & F. Schenkel (2013) “SNP selection for predicting a quantitative trait” in the *Journal of Applied Statistics*, 40(3), 600-613.
  25. **N. Bifulchi**, **R. Deardon** & Z. Feng (2013) “Spatial approximations of network-based individual level infectious disease models” in *Spatial & Spatio-temporal Epidemiology*, 6, 59-70.
  24. T. Agvar, **R. Deardon** & J. Fryxell (2013) “An empirically parameterized individual based model of animal movement, perception and memory” in *Ecological Modelling*, 251: 158-172.
  23. K. Bottoms, Z. Poljak, C. Dewey, **R. Deardon**, D. Holtkamp & R. Friendship (2013) “Evaluation of external biosecurity practices on southern Ontario farms” in *Preventive Veterinary Medicine*, 109(1-2):58-68.

22. **G.P.S. Kwong**, Z. Poljak, **R. Deardon** & C. Dewey (2013) “Bayesian analysis of risk factors for infection with a genotype of porcine reproductive and respiratory syndrome virus in Ontario swine herds using monitoring data” in *Preventive Veterinary Medicine*, 110(3-4):405-17.
  21. K. Bottoms, Z. Poljak, B. Friendship, J. Alsop, **R. Deardon** & C. Dewey (2013) “An assessment of external biosecurity on southern Ontario swine farms, and its application to surveillance on a geographic level” in the *Canadian Journal of Veterinary Research*, 77(4), 241 - 253.
- 
20. **I. Vrbik**, **R. Deardon**, Z. Feng, **A. Gardner** & J. Braun (2012) “Using individual-level models to model the spatio-temporal dynamics of combustion” in *Bayesian Analysis*, 7(3), 615 – 638. (Funded by: NSERC).
  19. **G.P.S. Kwong** & **R. Deardon** (2012) “Linearized forms of individual-level models for large-scale spatial infectious disease systems” in *Bulletin of Mathematical Biology*, 74(8), 1912 – 37. (Funded by: NSERC, OMAFRA).
  18. Y. Hosseinkashi, S. Chenouri, C. Small & **R. Deardon** (2012) “A Stochastic Graph Process for Epidemic Modelling” in *Canadian Journal of Statistics*, 40(1), 55 – 67. (Funded by: NSERC).
  17. **R. Deardon**, **B. Habibzadeh** & **H. Y. Chung** (2012) “Spatial measurement error in infectious disease models” in *Journal of Applied Statistics*, 39(5), 1139 – 1150. (Funded by: NSERC).
  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.
  15. H. Le, Z. Poljak, **R. Deardon** & C. Dewey (2012) “Clustering of and risk factors for the porcine high fever disease in a region of Vietnam” in *Trans-boundary and Emerging Diseases*, 59(1), 49 – 61.
  14. K. Bottoms, Z. Poljak, C. Dewey, **R. Deardon**, D. Holtkamp & R. Friendship (2012) “Investigation of strategies for the introduction and transportation of replacement gilts on southern Ontario sow farms” in *BMC Veterinary Research*, 8, 217.
- 
13. **A. Gardner**, **R. Deardon** & G. A. Darlington (2011) “Goodness-of-fit measures for individual-level infectious disease models in a Bayesian framework” in *Spatial & Spatio-temporal Epidemiology*, 2(4), 273 – 281. (Funded by: NSERC, OMAFRA).
- 
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).
  11. **B. Habibzadeh** & **R. Deardon** (2010), “The effect of misspecifying latent and infectious periods in space-time epidemic models” in *Statistical Communications in Infectious Diseases*, Vol. 2: Issue 1, Article 7. (Funded by: NSERC).
- 
10. T. J. McKinley, A. Cook & **R. Deardon** (2009) “Inference in epidemic models without likelihoods” in *The International Journal of Biostatistics*, 5(1), Article 24. (Funded by: NSERC).
  9. P.E. Caines, **R. Deardon** & H. P. Wynn (2009) “Bayes’ nets of time series: stochastic realizations and projections” in *Optimal Experimental Design and Related Areas* (Ed: L Pronzato and A Zhiglavsky), 155-166, Springer.
-

8. A. J. Grant, M. Sheppard, **R. Deardon**, S. P. Brown, G. Foster, C. E. Bryant, D. J. Maskell & P. Mastroeni (2008) "Caspase 3-dependent phagocyte death during systemic *Salmonella enterica* serovar Typhimurium infection of mice" in *Immunology*, 125(1), 28-37.
7. M. J. Tildesley, **R. Deardon**, N. J. Savill, P. Bessell, S. P Brooks, M. E. J. Woolhouse, B. T. Grenfell & M. J. Keeling (2008) "Accuracy of models for the 2001 foot-and-mouth disease epidemic" in *Proceedings of the Royal Society B*, 275(1641), 1459-1468. (Funded by: Wellcome Trust, UK).

---

6. N. J. Savill, D. J. Shaw, **R. Deardon**, M. J. Tildesley, M. J. Keeling, S. P. Brooks, M. E. J. Woolhouse & B. T. Grenfell (2007), "Effect of data quality on estimates of farm infectiousness trends in the UK 2001 foot-and-mouth disease epidemic" in *Journal of the Royal Society Interface*, 4, 235-241. (Funded by: Wellcome Trust, UK).

---

5. **R. Deardon**, S. G. Gilmour, N. A. Butler, K. Phelps & R. Kennedy (2006), "Designing field experiments which are subject to representation bias" in *Journal of Applied Statistics*, 33, 7, 665-680. (Funded by: EPSRC, UK).
4. M. J. Tildesley, N. J. Savill, D. J. Shaw, **R. Deardon**, S. P. Brooks, M. E. J. Woolhouse, B. T. Grenfell & M. J. Keeling (2006), "Optimal reactive vaccination strategies for an outbreak of foot-and-mouth disease in Great Britain" in *Nature*, 440, 1080, 83-86. (Funded by: Wellcome Trust, UK).
3. N. J. Savill, D. J. Shaw, **R. Deardon**, M. J. Tildesley, M. J. Keeling, S. P. Brooks, M. E. J. Woolhouse & B. T. Grenfell (2006), "Topographic determinants of foot and mouth disease transmission in the UK 2001 epidemic" in *BMC Veterinary Research*, Vol. 2:3. (Funded by: Wellcome Trust, UK).

---

2. **R. Deardon**, S. G. Gilmour, N. A. Butler, K. Phelps & R. Kennedy (2004), "A method for ascertaining and controlling representation bias in field trials for airborne plant pathogens" in the *Journal of Applied Statistics*, 31, 3, 2004, 329-343.

---

1. P.E. Caines, **R. Deardon** & H. P. Wynn (2002) "Conditional Orthogonality and Conditional Stochastic Realization" in *New Directions in Mathematical Systems Theory and Optimization*, Springer.

---

### Conference Proceedings

- A. Ogilvy, S. Collins, W. Hare, M. Hilt, T. Tuokko, **R. Deardon** & A. Jirasek. “Optimization of solid tank design for fan-beam optical CT based 3D radiation dosimetry.” Submitted to the International Conference on 3D and Advanced Dosimetry (IC3DDose), Quebec City, Canada.
- *M. Aghajanpoorpasha & R. Deardon* (2019) “On Minimum Cost Non-Uniform Sampling Schemes for Optimal Design of Control Charts: Application to  $\bar{X}$  and  $T^2$  Control Charts” Fourth North American International Conference on Industrial Engineering and Operations Management (IEOM).
- P.E. Caines, **R. Deardon** & H. P. Wynn (2002) “Conditional independence and general factorisations in times series graphical models” in the *2002 Proceedings of the American Statistical Association, Physical and Engineering Sciences Section* [CD-ROM], Alexandria, VA: American Statistical Association.

### Published Letters

- M. J. Keeling, M. J. Tildesley, N. J. Savill, M. E. J. Woolhouse, D. J. Shaw, **R. Deardon**, S. P. Brooks, & B. T. Grenfell (2007), “Veterinary epidemiology: Vaccination strategies for foot-and-mouth disease” (reply to Brief Communication Arising by Kitching et al.) in *Nature*, 445, E12-E13, 8 February 2007.
- M. J. Keeling, M. J. Tildesley, N. J. Savill, M. E. J. Woolhouse, D. J. Shaw, **R. Deardon**, S. P. Brooks, & B. T. Grenfell (2006), response to letter, “FMD control strategies” by Wingfield, Miller & Honhold in *The Veterinary Record*, May 20, 2006. (Funded by: Wellcome Trust, UK).

### Technical & Other Reports

- M. Lewis, P. Brown, C. Colijn, L. Cowen, C. Cotton, T. Day, **R. Deardon**, D. Earn, D. Haskell, J. Hefernan, P. Leighton, K. Murty, S. Otto, E. Rafferty, C. Hughes Tuohy, J. Wu & H. Zhu (2023) “Charting a future for emerging infectious disease modelling in Canada.” <http://hdl.handle.net/1828/15042>
- **R. Deardon** & S. P. Brooks (2007) “Bayesian modelling of the spatio-temporal dynamics of large-scale epidemics,” technical report, *Statistical Series #2007-312*, Department of Mathematics & Statistics, University of Guelph.
- P.E. Caines, **R. Deardon** & H. P. Wynn (2007) “Algebraic Methods for Conditional Independence in Time Series Graphical Models,” technical report, *Statistical Series #2007-313*, Department of Mathematics & Statistics, University of Guelph.

### Software

- **C. Ward**, **R. Deardon** & A. Schmidt (2023) “epidemicBCM: Bayesian modelling of dynamic behavioural change during an epidemic”
  - Github Repository (R Nimble): <https://github.com/ceward18/epidemicBCM>
- **W. Almutiry**, **V. Warriyar** & **R. Deardon** (2019) EpiILMCT: Continuous Time Individual-Level Models of Infectious Disease.
  - CRAN Webpage: <https://cran.r-project.org/web/packages/EpiILMCT/index.html>
  - Github Repository: <https://waleedalmutiry.github.io/EpiILMCT/>
- **J. Angevaere**, **R. Deardon**, Z. Feng (2017) PhyloTrees.jl: Phylogenetic trees in Julia.



- Github repository (Julia): <https://github.com/jangevaare/PhyloTrees.jl>
  - **J. Angevaare, R. Deardon, Z. Feng** (2017) SubstitutionModels.jl: Biological Sequence Substitution Models for Julia.
    - Github repository (Julia). <https://github.com/BioJulia/SubstitutionModels.jl>
  - **V. Warriyar & R. Deardon** (2017) EpiILM: Spatial and Network Based Individual Level Models for Epidemics (R Package).
    - CRAN Webpage: <https://CRAN.R-project.org/package=EpiILM>
  - **J. Angevaare, R. Deardon, Z. Feng** (2016) Pathogen.jl: Simulation and inference tools for phylo-dynamic individual level models.
    - Github repository (Julia): <https://github.com/jangevaa/Pathogen.jl>
  - **J. Angevaare, R. Deardon, Z. Feng** (2014) ilmtools: Tools for simulating and performing inference of individual level infectious disease transmission models.
    - Github repository (Julia): <https://github.com/jangevaa/ilmtools>
- 

### Theses

- **R. Deardon** (2001) “Representation Bias in Field Trials for Airborne Plant Pathogens” Ph.D. Thesis, School of Applied Statistics, University of Reading, UK.
- **R. Deardon** (1997) “Multiple Testing: An Investigation of the Power Properties of Members of a Family of Closed Test (including Hommel’s Test)” M.Sc. Thesis, Department of Mathematics, University of Southampton, UK.