Kylie E. C. Ainslie, PhD

Qualifications

2018	PhD Biostatistics, Emory University, Atlanta, GA
2016	MSc Biostatistics, Emory University, Atlanta, GA
2011	AB Biology and Mathematics (Double major), Ripon College, Ripon, WI
	Magna Cum Laude. Phi Beta Kappa

Employment

2022-Present

Honorary Assistant Professor

School of Public Health University of Hong Kong, Hong Kong SAR

- Led the analytical planning and development of novel methodologies for evaluating vaccine effectiveness in longitudinal studies of respiratory infections, including COVID-19 and influenza.
- Collaborated on multiple research projects, providing expert guidance, and contributing to six peer-reviewed publications, which advanced the understanding of vaccine impacts and informed subsequent public health recommendations.
- Conducted workshops and tutorials with 50+ participants on technical subjects such as using Git/GitHub and R packages for data analysis, enhancing team skills and supporting research through hands-on training.

2020-Present

Senior Researcher

Unit for Infectious Disease Modelling Centre for Infectious Diseases, Epidemiology, and Surveillance National Institute for Public Health and the Environment (RIVM), Bilthoven, The Netherlands

- Led development and implementation of a transmission model to support key decisions throughout the COVID-19 vaccine roll out in the Netherlands, resulting in an open-source R package, published peer-reviewed manuscript, and critical data-driven insights that directly influenced public health policy.
- Liaised with government advisory boards and engaged key stakeholders to translate complex scientific findings into actionable pandemic health policies through detailed presentations and reports, ensuring effective implementation of evidence-based strategies.
- Identified key gaps in the scientific literature and spearheaded the design and funding acquisition of a research project, collaborating with a multidisciplinary team and securing a €600,000 grant.

Employment

2020 - 2021

Visiting Researcher

MRC Centre for Global Infectious Disease Analysis Department of Infectious Disease Epidemiology, School of Public Health Imperial College London, London, UK

• Provided expert consultation on statistical methodologies to estimate COVID-19 vaccine effectiveness using data from a large-scale population surveillance study.

2018-2020

Research Associate

MRC Centre for Global Infectious Disease Analysis Department of Infectious Disease Epidemiology, School of Public Health Imperial College London, London, UK

- Led the development and management of the analytical pipeline for the Realtime Assessment of Community Transmission (REACT-1) study, enhancing the monitoring and analysis of SARS-CoV-2 transmission dynamics.
- Provided fortnightly situation reports on the COVID-19 epidemiological situation to the UK government, directly supporting pandemic response and informing public health strategies across England.
- Designed and implemented an agent-based model and R package to analyze individual susceptibility to repeated exposures, including infections and vaccinations, advancing understanding of population-level immunity and disease spread.

2016-2018

Research Assistant

SAS Institute Inc., Atlanta Regional Office, Atlanta, GA

2015-2016

Research Assistant

Department of Biostatistics and Bioinformatics

Emory University, Atlanta, GA

2012-2015

Research Assistant

Biostatistics and Bioinformatics Shared Resource

Winship Cancer Institute, Emory University, Atlanta, GA

Grants

2022 | Evaluation of WAning Vaccine Effectiveness (WAVE)

Role: Principal Investigator; Funder: RIVM; Amount:€600,000

Efficient and rapidly SCAlable EU-wide evidence-driven Pandemic response plans through dynamic Epidemic data assimilation (ESCAPE)

Role: Consortium member/ Work package leader; Funder: European Commission;

Amount: €3.2 million

Awards

2022	Exceptional Service Award (€1400) National Institute of Public Health and the Environment (RIVM), Bilthoven, The Netherlands
2020	Session Funding (\$1000) and complimentary registration (\$400) Society for Epidemiological Research Annual Meeting, Boston, MA
2016	Poster Competition Award, Georgia Statistics Day Georgia Institute of Technology, Atlanta, GA
2014-2016	Trainee, Broadening Experiences in Scientific Training (BEST) Program Emory University, Atlanta, GA
2016	Scholarship and Travel Award Recipient Summer Institute in Statistics and Modeling of Infectious Diseases University of Washington, Seattle, WA
2012	Travel Award Recipient, Statistical Genetics and Genomics Short Course University of Alabama at Birmingham, Birmingham, AL
2007-2011	Knop Science Scholarship (\$120,000) Ripon College, Ripon, WI

Teaching and Supervision

${\bf Workshops}$

2024	Introduction to R
2023	Introduction to Git and GitHub
2023	Using the Lexis class for time-to-event data

Teaching

2019–2020	Lecturer, Further Infectious Disease Modelling Imperial College London, London, UK
2019	Guest Lecturer, Introduction to Statistical Programming I Duke University, Durham, NC
2018	Demonstrator, Epidemiology and Control of Infectious Disease Short Course Imperial College London, London, UK
2017	Graduate Teaching Assistant, R Programming Emory University, Atlanta, GA
2013-2014	Graduate Teaching Assistant, Linear Modeling Emory University, Atlanta, GA
2012-2013	Graduate Teaching Assistant, Introductory Statistical Methods Emory University, Atlanta, GA

Supervision

2018 | David Cook, MSc of Epidemiology (with Dr. Ada Yan and Professor Steven Riley)

Service

Leadership Roles

2024	Organizer, R-Ladies Amsterdam
2020	Session Organizer and Chair, Society for Epidemiological Research Annual Meet-
	ing 2020
2018 – 2019	Postdoctoral Representative
	Department of Infectious Disease Epidemiology, Imperial College London
2018 – 2019	Chair, Steering Committee, Council for Emerging and New Statisticians
	Eastern North American Region, International Biometrics Society
2019	Session Organizer, Eastern North American Region Conference 2019
2018	Session Organizer, Eastern North American Region Conference 2018
2016-2018	Member, Steering Committee, Council for Emerging and New Statisticians
	Eastern North American Region, International Biometrics Society
2015-2018	Editor-In-Chief, Atlanta BEST Magazine
2015-2016	Professional Development Chair
	Department of Biostatistics and Bioinformatics, Emory University
2014 – 2015	Editor, Atlanta BEST Magazine

Computing

Programs	R (expert), SAS (Certified Base Programmer), C++, LaTeX
R packages	pika, serosolver, roa, morevac, vacamole, wave (in development), mitey (in development)
GitHub	https://github.com/kylieainslie
Website	https://kylieainslie.github.io

Languages

English (native)
Dutch (fluent)

Publications

Preprints

2024 Ainslie, K E C, M. Hooiveld, and J. Wallinga. On the epidemiological characteristics of scabies. (in preparation), Oct. 2024

Published

- 2024 T. K. Tsang, S. G. Sullivan, X. Huang, C. Wang, Y. Wang, J. Nealon, B. Yang, Ainslie, K E C, and B. J. Cowling. Prior infections and effectiveness of SARS-CoV-2 vaccine in test-negative studies: A systematic review and meta-analysis. Am. J. Epidemiol., June 2024
- K. Sherratt, A. Srivastava, Ainslie, K, D. E. Singh, A. Cublier, M. C. Marinescu, J. Carretero, A. C. Garcia, N. Franco, L. Willem, S. Abrams, C. Faes, P. Beutels, N. Hens, S. Müller, B. Charlton, R. Ewert, S. Paltra, C. Rakow, J. Rehmann, T. Conrad, C. Schütte, K. Nagel, S. Abbott, R. Grah, R. Niehus, B. Prasse, F. Sandmann, and S. Funk. Characterising information gains and losses when collecting multiple epidemic model outputs. Epidemics, 47(100765):100765, June 2024
- D. Chen, B. J. Cowling, **Ainslie**, **K E C**, Y. Lin, J. Y. Wong, E. H. Y. Lau, P. Wu, and J. Nealon. Association of COVID-19 vaccination with duration of hospitalization in older adults in hong kong. *Vaccine*, 42(9):2385–2393, Apr. 2024
- F. Miura, J. A. Backer, G. van Rijckevorsel, R. Bavalia, S. Raven, M. Petrignani, Ainslie, K E C, J. Wallinga, and Dutch Mpox Response Team. Time scales of human mpox transmission in the netherlands. *J. Infect. Dis.*, 229(3):800–804, Mar. 2024
- 2024 C. Zachreson, J. Savulescu, F. M. Shearer, M. J. Plank, S. Coghlan, J. C. Miller, Ainslie, K E C, and N. Geard. Ethical frameworks should be applied to computational modelling of infectious disease interventions. *PLoS Comput. Biol.*, 20 (3):e1011933, Mar. 2024
- M. Jit, K. E. C. Ainslie, C. Althaus, C. Caetano, V. Colizza, D. Paolotti, P. Beutels, et al. Reflections on epidemiological modeling to inform policy during the covid-19 pandemic in western europe, 2020–23. *Health Affairs*, 42, 2023. doi: https://doi.org/10.1377/hlthaff.2023.00688
- S. G. Sullivan, A. Khvorov, X. Huang, C. Wang, **Ainslie, K**, J. Nealon, B. Yang, B. Cowling, and T. Tsang. The need for a clinical case definition in test-negative design studies estimating vaccine effectiveness. *NPJ Vaccines*, 8, Aug. 2023
- Ainslie, K. E. C., J. A. Backer, P. T. de Boer, A. J. van Hoek, D. Klinkenberg, H. K. Altes, K. Y. Leung, H. de Melker, F. Miura, and J. Wallinga. A scenario modelling analysis to anticipate the impact of COVID-19 vaccination in adolescents and children on disease outcomes in the Netherlands, summer 2021. Eurosurveillance, 44(27):pii=2101090, 2022. doi: https://doi.org/10.2807/1560-7917.ES.2022.27.44.2101090
- M. Chadeau-Hyam, H. Wang, O. Eales, D. Haw, B. Bodinier, M. Whitaker, C. E. Walters, K. E. C. Ainslie, C. Atchison, C. Fronterre, P. J. Diggle, A. J. Page, A. J. Trotter, D. Ashby, W. Barclay, G. Taylor, G. Cooke, H. Ward, A. Darzi, S. Riley, C. A. Donnelly, P. Elliott, and COVID-19 Genomics UK consortium. SARS-CoV-2 infection and vaccine effectiveness in england (REACT-1): a series of cross-sectional random community surveys. Lancet Respir Med, 10(4):355–366, Apr. 2022

- O. Eales, **K. E.C. Ainslie**, C. E. Walters, H. Wang, C. Atchison, D. Ashby, C. A. Donnelly, G. Cooke, W. Barclay, H. Ward, A. Darzi, P. Elliott, and S. Riley. Appropriately smoothing prevalence data to inform estimates of growth rate and reproduction number. *Epidemics*, 40:100604, 2022. ISSN 1755-4365. doi: https://doi.org/10.1016/j.epidem.2022.100604
- Ainslie, K. E. C. and S. Riley. Is annual vaccination best?: a modelling study of influenza vaccination in children. *Vaccine*, 40(21):2940–2948, 2022. doi: 10.1016/j.vaccine.2022.03.065
- F. Miura, K. Y. Leung, D. Klinkenberg, K. E. C. Ainslie, and J. Wallinga. Optimal vaccine allocation for COVID-19 in the Netherlands: a data-driven prioritization. *PLoS Computational Biology*, 17(12):e1009697, 2021. doi: 10.1371/journal.pcbi.1009697
- P. Elliott, D. Haw, H. Wang, O. Eales, C. E. Walters, K. E. C. Ainslie, C. Atchison, C. Fronterre, P. J. Diggle, A. J. Page, A. J. Trotter, S. J. Prosolek, D. Ashby, C. A. Donnelly, W. Barclay, G. Taylor, G. Cooke, H. Ward, A. Darzi, and S. Riley. Exponential growth, high prevalence of SARS-CoV-2, and vaccine effectiveness associated with the delta variant. Science, 374(6574):eabl9551, 2021. doi: 10.1126/science.abl9551
- M. Haber, J. E. Tate, B. A. Lopman, W. Qia, **K. E. C. Ainslie**, and U. D. Parashar. Comparing statistical methods for detecting and estimating waning efficacy of rotavirus vaccines in developing countries. *Hum Vaccin Immunother*, 17(11):4632–4635, 2021. doi: 10.1080/21645515.2021.1968738.
- S. Riley, **K. E. C. Ainslie**, O. Eales, C. E. Walters, H. Wang, C. Atchison, C. Fronterre, P. J. Diggle, D. Ashby, C. A. Donnelly, G. Cooke, W. Barclay, H. Ward, A. Darzi, and P. Elliott. Resurgence of SARS-CoV-2: Detection by community viral surveillance. *Science*, 372(6545):990–995, 2021. ISSN 0036-8075. doi: 10.1126/science.abf0874
- P. Nouvellet, S. Bhatia, A. Cori, K. E. C. Ainslie, M. Baguelin, S. Bhatt, A. Boonyasiri, N. F. Brazeau, L. Cattarino, L. V. Cooper, H. Coupland, Z. M. Cucunuba, G. Cuomo-Dannenburg, A. Dighe, B. A. Djaafara, I. Dorigatti, O. D. Eales, S. L. van Elsland, F. F. Nascimento, R. G. FitzJohn, K. A. M. Gaythorpe, L. Geidelberg, W. D. Green, A. Hamlet, K. Hauck, W. Hinsley, N. Imai, B. Jeffrey, E. Knock, D. J. Laydon, J. A. Lees, T. Mangal, T. A. Mellan, G. Nedjati-Gilani, K. V. Parag, M. Pons-Salort, M. Ragonnet-Cronin, S. Riley, H. J. T. Unwin, R. Verity, M. A. C. Vollmer, E. Volz, P. G. T. Walker, C. E. Walters, H. Wang, O. J. Watson, C. Whittaker, L. K. Whittles, X. Xi, N. M. Ferguson, and C. A. Donnelly. Reduction in mobility and COVID-19 transmission. Nature Communications, 12:1090, 2021. doi: 10.1038/s41467-021-21358-2
- H. Fu, H. Wang, X. Xi, A. Boonyasiri, Y. Wang, W. Hinsley, R. Fraser, K. J.and McCabe, D. Olivera Mesa, J. Skarp, A. Ledda, T. Dewé, A. Dighe, P. Winskill, S. L. van Elsland, **K. E. C. Ainslie**, M. Baguelin, S. Bhatt, O. Boyd, N. F. Brazeau, L. Cattarino, G. Charles, H. Coupland, Z. M. Cucunuba, G. Cuomo-Dannenburg, C. A. Donnelly, I. Dorigatti, O. D. Eales, R. G. FitzJohn, S. Flaxman, K. A. M. Gaythorpe, A. C. Ghani, W. D. Green, A. Hamlet, K. Hauck, D. J. Haw, B. Jeffrey, D. J. Laydon, J. A. Lees, T. Mellan, S. Mishra, G. Nedjati-Gilani, P. Nouvellet, L. Okell, K. V. Parag, M. Ragonnet-Cronin, S. Riley, N. Schmit, H. A. Thompson, H. J. T. Unwin, R. Verity, M. A. C. Vollmer, E. Volz, P. G. T. Walker, C. E. Walters, O. J. Watson, C. Whittaker, L. K. Whittles, N. Imai, S. Bhatia, and N. M. Ferguson. Database of epidemic trends and control measures during the first wave of COVID-19 in mainland china. *Int. J. Infect. Dis.*, 102:463–471, Jan. 2021

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- A. Dighe, L. Cattarino, G. Cuomo-Dannenburg, J. Skarp, N. Imai, S. Bhatia, K. Gaythorpe, Ainslie, KEC, M. Baguelin, S. Bhatt, A. Boonyasiri, N. Brazeau, L. Cooper, H. Coupland, Z. Cucunuba, I. Dorigatti, O. Eales, S. van Elsland, R. FitzJohn, W. Green, D. Haw, W. Hinsley, E. Knock, D. Laydon, T. Mellan, S. Mishra, G. Nedjati-Gilani, P. Nouvellet, M. Pons-Salort, H. Thompson, H. Unwin, R. Verity, M. Vollmer, C. Walters, O. Watson, C. Whittaker, L. Whittles, A. Ghani, C. Donnelly, N. Ferguson, and S. Riley. Response to covid-19 in south korea and implications for lifting stringent interventions. BMC Med, 18(1):321, Oct 2020. doi: 10.1186/s12916-020-01791-8
- N. C. Grassly, M. Pons-Salort, E. P. K. Parker, P. J. White, N. M. Ferguson, and Imperial College COVID-19 Response Team*. Comparison of molecular testing strategies for covid-19 control: a mathematical modelling study. Lancet Infectious Diseases, 2020. doi: 10.1016/S1473-3099(20)30630-7

 * as part of the Imperial College COVID-19 Response Team
- H. Thompson, N. Imai, A. Dighe, K. E. C. Ainslie, M. Baguelin, S. Bhatia, S. Bhatt, A. Boonyasiri, O. Boyd, N. Brazeau, L. Cattarino, L. Cooper, H. Coupland, Z. Cucunuba, G. Cuomo-Dannenburg, B. Djaafara, I. Dorigatti, S. van Elsland, R. Fitzjohn, H. Fu, K. Gaythorpe, W. Green, T. Hallett, A. Hamlet, D. Haw, S. Hayes, W. Hinsley, B. Jeffrey, E. Knock, D. Laydon, J. Lees, T. Mangal, T. Mellan, S. Mishra, A. Mousa, G. Nedjati-Gilani, P. Nouvellet, L. Okell, K. Parag, M. Ragonnet-Cronin, S. R. H. Unwin, R. Verity, M. Vollmer, E. Volz, P. Walker, C. Walters, H. Wang, Y. Wang, O. Watson, C. Whittaker, L. Whittles, P. Winskill, X. Xi, C. Donnelly, and N. Ferguson. SARS-CoV-2 infection prevalence on repatriation flights from Wuhan City, China. Journal of Travel Medicine, 2020
- H. Ward, C. J. Atchison, M. Whitaker, K. E. C. Ainslie, J. Elliott, L. C. Okell,
 R. Redd, D. Ashby, C. A. Donnelly, W. Barclay, A. Darzi, G. Cooke, S. Riley, and
 P. Elliott. SARS-CoV-2 antibody prevalence in England following the first peak of the pandemic. *Nature Communications*, 12:905, 2021. doi: 10.1038/s41467-021-21237-w
- H. J. T. Unwin, S. Mishra, V. C. Bradley, A. Gandy, T. A. Mellan, H. Coupland, J. Ish-Horowicz, M. A. C. Vollmer, C. Whittaker, S. L. Filippi, X. Xi, M. Monod, O. Ratmann, M. Hutchinson, F. Valka, H. Zhu, I. Hawryluk, P. Milton, K. E. C. Ainslie, M. Baguelin, A. Boonyasiri, N. F. Brazeau, L. Cattarino, Z. M. Cucunubá, G. Cuomo-Dannenburg, I. Dorigatti, O. D. Eales, J. W. Eaton, S. L. van Elsland, R. G. FitzJohn, K. A. M. Gaythorpe, W. Green, W. Hinsley, B. Jeffrey, E. Knock, D. J. Laydon, J. Lees, G. Nedjati-Gilani, P. Nouvellet, L. C. Okell, K. V. Parag, I. Siveroni, H. A. Thompson, P. Walker, C. E. Walters, O. J. Watson, L. K. Whittles, A. Ghani, N. M. Ferguson, S. Riley, C. A. Donnelly, S. Bhatt, and S. Flaxman. State-level tracking of COVID-19 in the united states. Nature Communications, 11:6189, 2020. doi: 10.1038/s41467-020-19652-6

2020 E. Lavezzo, E. Franchin, C. Ciavarella, G. Cuomo-Dannenburg, L. Barzon, C. Del Vecchio, L. Rossi, R. Manganelli, A. Loregian, N. Navarin, D. Abate, M. Sciro, S. Merigliano, E. De Canale, M. C. Vanuzzo, V. Besutti, F. Saluzzo, F. Onelia, M. Pacenti, S. Parisi, G. Carretta, D. Donato, L. Flor, S. Cocchio, G. Masi, A. Sperduti, L. Cattarino, R. Salvador, M. Nicoletti, F. Caldart, G. Castelli, E. Nieddu, B. Labella, L. Fava, M. Drigo, K. A. M. Gaythorpe, Imperial College COVID-19 Response Team*, A. R. Brazzale, S. Toppo, M. Trevisan, V. Baldo, C. A. Donnelly, N. M. Ferguson, I. Dorigatti, and A. Crisanti. Suppression of a SARS-CoV-2 outbreak in the Italian municipality of Vo'. Nature, 584:425-429, 2020. doi: 10.1038/s41586-020-2488-1

* as part of the Imperial College COVID-19 Response Team

2020 S. Flaxman, S. Mishra, A. Gandy, H. J. T. Unwin, T. A. Mellan, H. Coupland, C. Whittaker, H. Zhu, T. Berah, J. W. Eaton, M. Monod, P. N. Perez-Guzman, N. Schmit, L. Cilloni, K. E. C. Ainslie, M. Baguelin, A. Boonyasiri, O. Boyd, L. Cattarino, L. V. Cooper, Z. Cucunubá, G. Cuomo-Dannenburg, A. Dighe, B. Djaafara, I. Dorigatti, S. L. van Elsland, R. G. FitzJohn, K. A. M. Gaythorpe, L. Geidelberg, N. C. Grassly, W. D. Green, T. Hallett, A. Hamlet, W. Hinsley, B. Jeffrey, E. Knock, D. J. Laydon, G. Nedjati-Gilani, P. Nouvellet, K. V. Parag, I. Siveroni, H. A. Thompson, R. Verity, E. Volz, C. E. Walters, H. Wang, Y. Wang, O. J. Watson, P. Winskill, X. Xi, P. G. T. Walker, A. C. Ghani, C. A. Donnelly, S. Riley, M. A. C. Vollmer, N. M. Ferguson, L. C. Okell, and S. Bhatt. Estimating the effects of non-pharmaceutical interventions on COVID-19 in europe. Nature, 584:257–261, 2020. doi: 10.1038/s41586-020-2405-7

2020 A. B. Hogan*, B. L. Jewell*, E. Sherrard-Smith*, J. F. Vesga*, O. J. Watson*, C. Whittaker*, A. Hamlet, J. A. Smith, P. Winskill, R. Verity, M. Baguelin, J. A. Lees, L. K. Whittles, K. E. C. Ainslie, S. Bhatt, A. Boonyasiri, N. F. Brazeau, L. Cattarino, L. V. Cooper, H. Coupland, G. Cuomo-Dannenburg, A. Dighe, B. A. Djaafara, C. A. Donnelly, J. W. Eaton, S. L. van Elsland, R. G. FitzJohn, H. Fu, K. A. M. Gaythorpe, W. Green, D. J. Haw, S. Hayes, W. Hinsley, N. Imai, D. J. Laydon, T. D. Mangal, T. A. Mellan, S. Mishra, G. Nedjati-Gilani, K. V. Parag, H. A. Thompson, H. J. T. Unwin, M. A. C. Vollmer, C. E. Walters, H. Wang, Y. Wang, X. Xi, N. M. Ferguson, L. C. Okell, T. S. Churcher, N. Arinaminpathy, A. C. Ghani, P. G. T. Walker, and T. B. Hallett. Potential impact of the COVID-19 pandemic on HIV, tuberculosis, and malaria in low-income and middle-income countries: a modelling study. Lancet, 2020. doi: 10.1016/S2214-109X(20)30288-6 * equal contribution

B. Jeffrey*, C. E. Walters*, K. E. C. Ainslie*, O. Eales*, C. Ciavarella, S. Bha-2020 tia, S. Hayes, M. Baguelin, A. Boonyasiri, N. F. Brazeau, G. Cuomo-Dannenburg, R. G. FitzJohn, K. Gaythorpe, W. Green, N. Imai, T. A. Mellan, S. Mishra, P. Nouvellet, H. J. T. Unwin, R. Verity, M. Vollmer, C. Whittaker, N. M. Ferguson, C. A. Donnelly, and S. Riley. Anonymised and aggregated crowd level mobility data from mobile phones suggests that initial compliance with COVID-19 social distancing interventions was high and geographically consistent across the UK. Wellcome Open Research, 5:170, 2020

* equal contribution

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- J. Hay, A. Mintor, K. E. C. Ainslie, J. Lessler, B. Yang, D. A. T. Cummings, A. Kucharski, and S. Riley. An open source tool to infer epidemiological and immunological dynamics from serological data: serosolver. *PLoS Comp Biol*, 16 (5):e1007840, 2020. doi: 10.1371/journal.pcbi.1007840
- Ainslie, K. E. C., M. Haber, and W. A. Orenstein. Challenges in estimating influenza vaccine effectiveness. *Expert Review of Vaccines*, 18(6):615–628, 2019
- 2019 **Ainslie, K. E. C.**, M. Haber, and W. A. Orenstein. Bias of influenza vaccine effectiveness estimates from test-negative studies conducted during an influenza pandemic. *Vaccine*, 37(14):1987–1993, 2019
- 2018 Ainslie, K. E. C., M. Haber, and W. A. Orenstein. A dynamic model of bias of estimates of influenza vaccine effectiveness from observational studies. *American Journal of Epidemiology*, 118(2):451–460, 2018
- 2017 Ainslie, K. E. C., M. Shi, M. Haber, and W. A. Orenstein. On the bias of estimates of influenza vaccine effectiveness from the test-negative studies. *Vaccine*, 35:7297–7301, 2017
- Ainslie, K. E. C., M. Haber, R. E. Malosh, J. G. Petrie, and A. S. Monto. Maximum likelihood estimation of influenza vaccine effectiveness against transmission from the household and from the community. *Statistics in Medicine*, 37 (6):970–982, 2017
- M. Shi, Q. An, **K. E. C. Ainslie**, M. Haber, and W. A. Orenstein. A comparison of the test-negative and the traditional case-control study designs for estimation of influenza vaccine effectiveness under nonrandom vaccination. *BMC Infectious Diseases*, 17:757–777, 2017
- J. Kowalski, B. Dwivedi, S. Newman, J. M. Switchenko, R. Pauly, D. A. Gutman, J. Aurora, K. Ghandi, K. Ainslie, G. Doho, Z. Qin, C. S. Moreno, M. R. Rossi, P. M. Vertino, S. Lonial, L. Bernal-Mizrachi, and L. H. Boise. Gene integrated set profile analysis: a context-based approach for inferring biological endpoints. Nucleic Acids Research, 44(7):e69, 2016

Reports

- Ainslie, K, J. Backer, A. J. van Hoek, D. Klinkenberg, S. McDonald, F. Miura, and J. Wallinga. The expected outcome of COVID-19 vaccination strategies. Technical report, Rijksinstituut voor Volksgezondheid en Milieu (RIVM), Aug. 2021
- N. Ferguson, D. Laydon, G. Nedjati Gilani, N. Imai, Ainslie, K, M. Baguelin, S. Bhatia, A. Boonyasiri, Z. Cucunuba Perez, G. Cuomo-Dannenburg, A. Dighe, I. Dorigatti, H. Fu, K. Gaythorpe, W. Green, A. Hamlet, W. Hinsley, L. Okell, S. Van Elsland, H. Thompson, R. Verity, E. Volz, H. Wang, Y. Wang, P. Walker, P. Winskill, C. Whittaker, C. Donnelly, S. Riley, and A. Ghani. Report 9: Impact of non-pharmaceutical interventions (NPIs) to reduce COVID19 mortality and healthcare demand. Technical report, MRC Centre for Global Infectious Disease Analysis, Imperial College London, 2020

Presentations

Invited

- 2024 "On the epidemiological characteristics of scabies."

 Applied Mathematics Seminar, Department of Mathematical Sciences
 University of Liverpool, Liverpool, UK
- Keynote: "COVID-19 modelling for policy makers: Reflections of 3 years of COVID response."
 SPARK/SPECTRUM Annual Meeting, Cape Schanck, Victoria, Australia
- "COVID-19 modelling for policy makers: Reflections of 3 years of COVID response."
 University of Melbourne, Melbourne, Victoria, Australia
- "COVID-19 modelling for policy makers."
 TRS Seminar, School of Public Health, University of Hong Kong, Hong Kong SAR
- "Is annual vaccination best?: A modeling study of influenza vaccination strategies in children."
 MRC Seminar, MRC Centre for Global Infectious Disease Analysis
 School of Public Health, Imperial College London, London, UK
- 2019 "Using longitudinal models of serological data to optimise repeated influenza vaccination."

 MRC Vaccine Symposium, MRC Centre for Global Infectious Disease Analysis
- School of Public Health, Imperial College London, London, UK

 "A dynamic model for evaluation of bias of estimates of influenza vaccine effec-
- The dynamic model for evaluation of bias of estimates of influenza vaccine effectiveness from observational studies."

 School of Public Health, Hong Kong University, Hong Kong
- 2016 "Comparing estimates of influenza vaccine effectiveness from case-control studies under non-random vaccination."
 WHO Collaborating Centre for Reference and Research on Influenza Peter Doherty Institute, Melbourne, Victoria, Australia
- 2016 "Estimation of effectiveness of influenza vaccination in household studies." Biology Seminar, Ripon College, Ripon, WI

Contributed

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2023	"A scenario modelling analysis to anticipate the impact of COVID-19 vaccination in adolescents and children on disease outcomes in the Netherlands, summer 2021." Australia and New Zealand Industrial and Applied Mathematics (ANZIAM) Conference 2023, Cairns, Queensland, Australia
2023	"Determining the trade-offs between different COVID-19 control strategies in the Netherlands: A counterfactual analysis." Australia and New Zealand Industrial and Applied Mathematics (ANZIAM) Conference 2023, Cairns, Queensland, Australia
2020	"Overview of the test-negative design and how to control for bias and confounding." Society for Epidemiological Research Annual Meeting, Boston, MA
2019	"Potential public health benefits from reduced delay in the production of pandemic influenza vaccine." Epidemics 7, Charleston, SC
2018	"A dynamic model for evaluation of bias of estimates of influenza vaccine effectiveness from observational studies." Eastern North American Region Spring Meeting, Atlanta, GA
2016	"Estimation of effectiveness of influenza vaccination in household studies." International Biometric Conference, Victoria, Canada
2016	"Estimation of effectiveness of influenza vaccination in household studies." Eastern North American Region Spring Meeting, Austin, TX
2014	"A robust statistical framework to whole-genome outlier identification for characterizing structural variants." Joint Statistical Meeting, Boston, MA
2010	"The exact distribution of the clustered Wilcoxon test." Mathematics Association of America – WI Conference, Oshkosh, WI

Posters

2022	"A scenario modelling analysis to anticipate the impact of COVID-19 vaccination in adolescents and children on disease outcomes in the Netherlands, summer 2021" Options XI, Belfast, UK
2021	"The impact of vaccinating adolescents and children on SARS-CoV-2 transmission" Epidemics 8, Online
2019	"Is annual vaccination best?: Evaluating influenza vaccination strategies in children" Epidemics 7, Charleston, SC, USA
2019	"Deaths averted from reduced delay in pandemic influenza vaccine production" Options X for the Control of Influenza, Singapore
2019	"Is annual vaccination best?: Evaluating influenza vaccination strategies in children" Options X for the Control of Influenza, Singapore
2019	"The bias of estimates of influenza vaccine effectiveness from test-negative studies during a pandemic" 40th Annual Conference of the International Society for Clinical Biostatistics, Leuven, Belgium
2017	"Bias of influenza vaccine effectiveness estimates from case-control studies" European Scientific Working Group on Influenza, Riga, Latvia
2016	"Estimation of effectiveness of influenza vaccination in household studies" Georgia Statistics Day, Atlanta, GA
2015	"A robust statistical framework for whole genome outlier identification" Winship Symposium, Emory University, Atlanta, GA, USA
2012	"Outlier analysis to filter translocations from next generation sequence data" IEEE International Conference on Bioinformatics and Biomedicine, Philadelphia, PA, USA