

KENNETH BAILLIE

QUALIFICATIONS AND FELLOWSHIP AWARDS

- Academy of Medical Sciences, 2022: **FMedSci**
Royal Society of Edinburgh, 2022: **FRSE**
Faculty of Intensive Care Medicine, 2016: **FFICM**
Royal College of Physicians, 2016: **FRCP(Ed)**
University of Edinburgh, 2012: **PhD** “Statistical Genetics in Infectious Disease”
Royal College of Anaesthetists 2009: **FRCA**
Royal College of Physicians, 2005: **MRCP(UK)**
University of Edinburgh, 2002: **MB,ChB**
University of Edinburgh, 1999: **BSc(Hons)(Med Sci)** in Physiology

SELECTED PRIZES AND AWARDS

- 2023 | **Chancellor's award for Research.** An annual prize awarded by the Principal and Chancellor of the University of Edinburgh.
- 2022 | **Tenovus Scotland Lady Margaret McLellan Award.** A biennial award for outstanding contributions to medical science in Scotland.
- 2021 | **Pride of Scotland “Special Recognition Award”.** One of two public awards given each year to recognise outstanding public service in any field.
- 2021 | **RCPE Ballantyne Prize.** The Royal College of Physicians of Edinburgh awards this prize, usually once every 10-15 years, for studies in the inheritance of disease.
- 2018 | **International Sepsis Forum Lowry-Fink Fellowship.** The International Sepsis Forum selects one “rising star” in sepsis research worldwide each year to appoint to this position.
- 2014 | **Wellcome-Beit Prize Fellowship.** Awarded to four intermediate fellowship candidates each year “to provide additional recognition for outstanding biomedical researchers.”
- 2013 | **Intensive Care Society Gold Medal.** Awarded annually for research related to critical care medicine.
- 2013 | **ANZICS Global Rising Star Award.** An annual award from the Australia and New Zealand Intensive Care Society.
- 2006 | **Nuffield Medal for the FRCA Primary.** For obtaining the highest marks in the UK in this sitting.
- 2001 | **Sir William Darling Memorial Prize.** Awarded by the Principal to the students “who have done most in any one year to advance the reputation of Edinburgh University”.

CURRENT EMPLOYMENT

2022-present	Co-Director, Pandemic Science Hub, University of Edinburgh
2021-present	Professor of Experimental Medicine, University of Edinburgh
2021-present	Wellcome Trust Senior Fellow, University of Edinburgh
2020-present	Director of Research, UK Intensive Care Society
2014-present	Honorary Consultant in Critical Care Medicine, Royal Infirmary of Edinburgh

PREVIOUS EMPLOYMENT

2019-2022	Visiting Professor, Peking University, Beijing, China
2014-21	Group Leader, Roslin Institute, University of Edinburgh
2014-20	Deputy Director of Research, UK Intensive Care Society
2014-20	Wellcome Trust Intermediate Clinical Fellow, University of Edinburgh
2014-15	Visiting Scientist, Broad Institute of Harvard and MIT, Cambridge, MA
Nov 2014	Consultant to World Health Organisation on Ebola, WHO country HQ, Sierra Leone.
2013-14	Honorary Advanced Specialist Trainee, Critical Care, Royal Infirmary Edinburgh.
May 2013	Consultant to World Health Organisation on MERS/Influenza, WHO HQ, Geneva.
2008-14	Clinical Lecturer, University of Edinburgh & Honorary Specialist Trainee in Anaesthesia & Critical Care, South East Scotland School of Anaesthesia.
2007-08	Specialist Trainee, Anaesthesia & Critical Care, Royal Infirmary, Edinburgh.
2005-07	Senior House Officer, Anaesthesia & Critical Care, Royal Infirmary Edinburgh
2003-05	Senior House Officer, General Medicine, Western Infirmary, Glasgow.
2002-03	House Officer, Anaesthesia, Medicine & Surgery, Royal Infirmary, Edinburgh.

RESEARCH GRANTS

FUNDING BODY & TITLE	START	AMOUNT	ROLE
MRC: Edinburgh Molecular Mechanisms Cluster	01/03/24 (4y)	£5,313,306	CI
EPSRC: Causality in Healthcare AI with Real Data	01/02/24 (5y)	£6,000,000	Coapplicant
ISARIC CCP activation for acute hepatitis of unknown cause	01/10/22 (1y)	£1,227,465	CI
Interim support for ISARIC4C CCP activation for acute severe hepatitis of unknown cause	01/06/22 (1y)	£74,267	CI
Baillie Gifford Pandemic Science Hub	01/06/22 (5y)	£14,700,000	Co-CI
CSO: Time critical precision medicine for acute critical illness (TRAITS)	01/04/22 (4y)	£3,610,000	Coapplicant
Wellcome Senior Research Fellowship : Translational Genomics in Critical Care Medicine	01/11/21 (5y)	£1,659,251	Applicant
UKRI: National Core Studies - Outbreak Data Analysis Platform	1/08/21 (1y)	£958,915	CI
UKRI: UK Coronavirus Immunology consortium (UK-CIC)	25/08/20 (1y)	£6,541,000	WP lead
UKRI: Post-hospitalisation for COVID-19 follow-up (PHOSP)	21/07/20 (1y)	£9,700,000	WP lead
DHSC: GenOMICC-Genomics England Partnership for whole genome sequencing	01/06/20 (1.5y)	£28,000,000	CI
UKRI: ISARIC4C (Coronavirus Clinical Characterisation Consortium)	18/03/20 (1.5y)	£5,958,870	CI
CSO: Coronavirus Clinical Characterisation (Scotland)	01/04/20 (1y)	£296,773.90	Coapplicant
NIHR: Coronavirus Clinical Information Network (CO-CIN)	03/03/20 (2y)	£8,900,000	Coapplicant
Technology Strategy Board: Establishing a Pipeline for African Swine Fever Virus Genetic Targets	01/10/19 (1.5y)	£105,767	Coapplicant

Tropical Animal Genetics (UK) Ltd: Cell library for candidate genes involved in Newcastle Disease Virus infection in vitro	01/09/19 (3y)	£100,000	Coapplicant
BBSRC: Host factors controlling foot-and-mouth disease virus (FMDV) replication: towards genetic control of FMD in pigs	01/08/19 (3y)	£901,522	Coapplicant
Fiona Elizabeth Agnew Trust (FEAT): Genetics of Mortality in Critical Care (GenOMICC)	01/04/19 (3y)	£120,000	CI
BBSRC 18ALERT Equipment Initiative: Large memory HPC infrastructure to underpin world-class biological research	01/04/19 (3y)	£600,000	Coapplicant
Health Technology Accelerator Fund: Guiding Therapy in Septic Shock	08/01/19 (3y)	£53,000	CI
Wellcome: Clinical PhD Fellowship for Dr Clark Russell	01/08/18 (3y)	£650,000	Co-supervisor
BBSRC: Identification of interferon stimulated genes that restrict cross-species transmission of influenza A virus.	01/03/19 (3y)	£756,000	Coapplicant
Wellcome: (Institution Strategic Support Fund) Generation of a constitutive Cas9 pig for gene-level pharmaceutical target validation.	01/06/18 (1y)	£50,000	PI
Wellcome: Clinical PhD Fellowship for Dr Nick Parkinson	01/08/17 (3y)	£650,000	CI
MRC: The SHIELD Consortium for Host Defence Therapeutics	01/11/16 (5y)	£4,500,000	WP lead
Wellcome: Establishing causal relationships from phenotypic, genotypic, and proteomic data.	01/08/16 (3y)	£650,000	Co-supervisor
BBSRC: Tools and resources: Generation of genome wide CRISPR libraries for pig and chicken	01/10/16 (1y)	£88,000	Coapplicant
Wellcome Trust: A clinical trial of convalescent plasma for treatment of Ebolavirus disease	01/03/15 (1y)	£120,000	Coapplicant
US Center for Disease Control: HostFlu observational study of influenza in Malawi	01/08/14 (5y)	£160,000	Co-CI
Wellcome Intermediate Clinical Fellowship : Identification and characterisation of host factors underlying susceptibility to influenza	01/08/14 (4y)	£1,011,082	Applicant
Wellcome Trust: (Institution Strategic Support Fund) The International Severe Influenza Genetics (ISIG) Programme	01/02/12 (1y)	£47,531	CI
Wellcome: Mechanisms of Severe Acute Influenza Consortium (MOSAIC)	01/09/09 (5y)	£3,645,239	WP lead
UK Embassy to United States of America UK/US Collaboration Development Award	01/08/10 (1mth)	£1500	Applicant
Chief Scientist Office: (Project Grant) GenISIS: Genetics of Influenza Susceptibility in Scotland	01/11/09 (2y)	£89,742	CI
Wellcome Trust: (Clinical Fellowship) Comparative genomics of the inflammatory response	01/08/09 (3y)	£267,208	Applicant
British Heart Foundation: (Project Grant) Oxidative Stress, erythrocyte anion channel damage, and antioxidant function in healthy lowlanders at high altitude.	01/04/01 (1y)	£10,525	CI
The Carnegie Trust for the Universities of Scotland: Apex Bolivia 2001	01/03/01 (1y)	£2000	CI

SELECTED PUBLICATIONS (*JOINT AUTHORSHIP) ([†]CORRESPONDING)

(h-index: 95 — citations: 69413 — ORCID:0000-0001-5258-793X)

Baillie JK, Angus D, Burnham K, Calandra T, Calfee C, Gutteridge A, Hacohen N, Khatri P, Langley R, Ma'ayan A, Marshall J, ..., Randolph AG. Causal inference can lead us to modifiable mechanisms and informative archetypes in sepsis. *Intensive Care Medicine*(2024); doi:10.1007/s00134-024-07665-4

Tan CCS, Kelly G, Cregan J, Wilson JD, James T, Chand M, Hopkins S, Swets M, **Baillie JK**, Jeffery K, Walker AS, Eyre DW, Stoesser N, Matthews PC. Retrospective analysis of hospital electronic health records reveals unseen cases of acute hepatitis with unknown aetiology in adults in Oxfordshire. *BMC Public Health*(2024); 24: 1890. **PMC11251388**

Sidhu JK, Siggins MK, Liew F, Russell CD, Uruchurtu ASS, Davis C, Turtle L, Moore SC, Hardwick HE, Oosthuyzen W, ..., **Baillie JK**, Openshaw PJM, Thwaites RS. Delayed Mucosal Antiviral Responses Despite Robust Peripheral Inflammation in Fatal COVID-19. *The Journal of Infectious Diseases*(2024); 230: e17-e29. doi:10.1093/infdis/jiad590

Boos J, van der Made CI, Ramakrishnan G, Coughlan E, Asselta R, Löscher B, Valenti LVC, de Cid R, Bujanda L, Julià A, Pairo-Castineira E, **Baillie JK**, ..., Ludwig KU. Stratified analyses refine association between TLR7 rare variants and severe COVID-19. *HGG Advances*(2024); 100323. doi:10.1016/j.xhgg.2024.100323

Chu JY, McCormick B, Sundaram K, Hardisty G, Karmakar U, Pumpe C, Krull E, Lucas CD, Amado-Azevedo J, Hordijk PL, ..., **Baillie JK**, Rossi AG, Vermeren S. ARAP3 protects from excessive formylated peptide-induced microvascular leakage by acting on endothelial cells and neutrophils. *The Journal of Pathology*(2024); doi:10.1002/path.6288

Turtle L, Elliot S, Drake TM, Thorpe M, Khouri EG, Greenhalf W, Hardwick HE, Leeming G, Law A, Oosthuyzen W, ..., **Baillie JK**, Openshaw PJM, Docherty AB, Semple MG, Harrison EM, Palmieri C. Changes in hospital mortality in patients with cancer during the COVID-19 pandemic (ISARIC-CCP-UK): a prospective, multicentre cohort study. *The Lancet. Oncology*(2024); doi:10.1016/S1470-2045(24)00107-4

Michael BD, Dunai C, Needham EJ, Tharmaratnam K, Williams R, Huang Y, Boardman SA, Clark JJ, Sharma P, Subramaniam K, ..., **Baillie JK**, Openshaw PJ, Semple MG, Taams LS, Menon DK. Author Correction: Para-infectious brain injury in COVID-19 persists at follow-up despite attenuated cytokine and autoantibody responses. *Nature Communications*(2024); 15: 2918. **PMC10995197**

Liew F, Efstatihou C, Fontanella S, Richardson M, Saunders R, Swieboda D, Sidhu JK, Ascough S, Moore SC, Mohamed N, ..., **Baillie JK**, Semple MG, Turtle L, Evans RA, Wain LV, Brightling C, Thwaites RS, Openshaw PJM. Large-scale phenotyping of patients with long COVID post-hospitalization reveals mechanistic subtypes of disease. *Nature Immunology*(2024); 25: 607-621. **PMC11003868**

Shankar-Hari M, Calandra T, Soares MP, Bauer M, Wiersinga WJ, Prescott HC, Knight JC, **Baillie JK**, Bos LDJ, Derde LPG, Finfer S, ..., van der Poll T. Reframing sepsis immunobiology for translation: towards informative subtyping and targeted immunomodulatory therapies. *The Lancet. Respiratory Medicine*(2024); doi:10.1016/S2213-2600(23)00468-X

Horby PW, Peto L, Staplin N, Campbell M, Pessoa-Amorim G, Mafham M, Emberson JR, Stewart R, Prudon B, Uriel A, ..., **Baillie JK**, Buch MH, Day J, Faust SN, Jaki T, Jeffery K, Juszczak E, Knight M, Lim WS, Montgomery A, Mumford A, Rowan K, Thwaites G, Haynes R, Landray MJ. Dimethyl fumarate in patients admitted to hospital with COVID-19 (RECOVERY): a randomised, controlled, open-label, platform trial. *Nature Communications*(2024); 15: 924. **PMC10831058**

Devine K, Russell CD, Blanco GR, Walker BR, Homer NZM, Denham SG, Simpson JP, Leavy OC, Elneima O, McAuley HJC, ..., **Kenneth Baillie J**, Reynolds RM. Plasma steroid concentrations reflect acute disease severity and normalise during recovery in people hospitalised with COVID-19. *Clinical Endocrinology*(2024); 100: 317-327. doi:10.1111/cen.15012

Cho S, Premraj L, Battaglini D, Fanning JP, Suen J, Bassi GL, Fraser J, Robba C, Griffe M, Solomon T, ..., **Baillie JK**, Sigfrid L, Scott JT, Citarella BW, Merson L, Arora RC, Whitman G, Thomson D, White N. Sex differences in post-acute neurological sequelae of SARS-CoV-2 and symptom resolution in adults after coronavirus disease 2019 hospitalization: an international multi-centre prospective observational study. *Brain Communications*(2024); 6: fcae036. **PMC10914448**

Swets MC, Kerr S, Scott-Brown J, Brown AB, Gupta R, Millar JE, Spata E, McCurrach F, Bretherick AD, Docherty A, ..., **Baillie JK**. Evaluation of pragmatic oxygenation measurement as a proxy for Covid-19 severity. *Nature Communications*(2023); 14: 7374. **PMC10651917**

Sidhu JK, Siggins MK, Liew F, Russell CD, Uruchurtu ASS, Davis C, Turtle L, Moore SC, Hardwick HE, Oosthuizen W, ..., **Baillie JK**, Openshaw PJM, Thwaites RS. Delayed mucosal anti-viral responses despite robust peripheral inflammation in fatal COVID-19. *The Journal of Infectious Diseases*(2023); doi:10.1093/infdis/jiad590

Michael BD, Dunai C, Needham EJ, Tharmaratnam K, Williams R, Huang Y, Boardman SA, Clark JJ, Sharma P, Subramaniam K, ..., **Baillie JK**, Openshaw PJ, Semple MG, Taams LS, Menon DK. Para-infectious brain injury in COVID-19 persists at follow-up despite attenuated cytokine and autoantibody responses. *Nature Communications*(2023); 14: 8487. [PMC10746705](#)

Costigan D, Fenn J, Yen S, Ilott N, Bullers S, Hale J, Greenhalf W, Conibear E, Koycheva A, Madon K, ..., **Baillie JK**, Semple MG, Openshaw PJM, Pearson C, Johnson J, Lalvani A, Thornton EE. A pro-inflammatory gut mucosal cytokine response is associated with mild COVID-19 disease and superior induction of serum antibodies. *Mucosal Immunology*(2023); doi:10.1016/j.mucimm.2023.11.005

Burke J, Pugh D, Farrah T, Hamid C, Godden E, MacGillivray TJ, Dhaun N, **Baillie JK**, King S, MacCormick IJC. Evaluation of an Automated Choroid Segmentation Algorithm in a Longitudinal Kidney Donor and Recipient Cohort. *Translational Vision Science and Technology*(2023); 12: 19. [PMC10668611](#)

Zhu Y, Almeida FJ, **Baillie JK**, Bowen AC, Britton PN, Brizuela ME, Buonsenso D, Burgner D, Chew KY, Chokephaibulkit K, Cohen C, ..., Short KR. International Pediatric COVID-19 Severity Over the Course of the Pandemic. *JAMA Pediatrics*(2023); [PMC10442787](#)

Pannhorst K, Carlson J, Hölder JE, Grey F, **Baillie JK**, Hölder D, Wöhnke E, Franzke K, Karger A, Fuchs W, Mettenleiter TC. The non-classical major histocompatibility complex II protein SLA-DM is crucial for African swine fever virus replication. *Scientific Reports*(2023); 13: 10342. [PMC10442341](#)

Hayes AJ, Zheng X, O'Kelly J, Neyton LPA, Bochkina NA, Uings I, Liddle J, **Baillie JK**, Just G, Binnie M, Homer NZM, ..., Mole DJ. Kynurenine monooxygenase regulates inflammation during critical illness and recovery in experimental acute pancreatitis. *Cell Reports*(2023); 42: 112763. doi:10.1016/j.celrep.2023.112763

Pairo-Castineira E, Rawlik K, Bretherick AD, Qi T, Wu Y, Nassiri I, McConkey GA, Zechner M, Klaric L, Griffiths F, ..., **Baillie JK**. Author Correction: GWAS and meta-analysis identifies 49 genetic variants underlying critical COVID-19. *Nature*(2023); doi:10.1038/s41586-023-06383-z

Yang H, Sun H, Brackenridge S, Zhuang X, Wing PAC, Quastel M, Walters L, Garner L, Wang B, Yao X, ..., **Baillie JK**, Mentzer AJ, Klenerman P, Borrow P, Dong T, McKeating JA, Gillespie GM, McMichael AJ. HLA-E-restricted SARS-CoV-2-specific T cells from convalescent COVID-19 patients suppress virus replication despite HLA class Ia down-regulation. *Science Immunology*(2023); 8: eabl8881. doi:10.1126/sciimmunol.abl8881

Docherty AB, Farrell J, Thorpe M, Egan C, Dunn S, Norman L, Shaw CA, Law A, Leeming G, Norris L, ..., **Baillie JK**, Pius R, Seth S, Harrison EM, Lone NI. Patient emergency health-care use before hospital admission for COVID-19 and long-term outcomes in Scotland: a national cohort study. *The Lancet. Digital Health*(2023); 5: e446-e457. doi:10.1016/S2589-7500(23)00051-1

Pauley E, Drake TM, Griffith DM, Sigrid L, Lone NI, Harrison EM, **Baillie JK**, Scott JT, Walsh TS, Semple MG, Docherty AB. Recovery from Covid-19 critical illness: A secondary analysis of the ISARIC4C CCP-UK cohort study and the RECOVER trial. *Journal of The Intensive Care Society*(2023); 24: 162-169. [PMC10225805](#)

Gillan JL, Chokshi M, Hardisty GR, Clohisey Hendry S, Prasca-Chamorro D, Robinson NJ, Lasota B, Clark R, Murphy L, Whyte MKB, **Baillie JK**, Davidson DJ, Bao G, Gray RD. CAGE sequencing reveals CFTR-dependent dysregulation of type I IFN signaling in activated cystic fibrosis macrophages. *Science Advances*(2023); 9: eadg5128. doi:10.1126/sciadv.adg5128

Pairo-Castineira E, Rawlik K, Bretherick AD, Qi T, Wu Y, Nassiri I, McConkey GA, Zechner M, Klaric L, Griffiths F, ..., **Baillie JK**. GWAS and meta-analysis identifies 49 genetic variants underlying critical COVID-19. *Nature*(2023); doi:10.1038/s41586-023-06034-3

Morris KM, Mishra A, Raut AA, Gaunt ER, Borowska D, Kuo RI, Wang B, Vijayakumar P, Chingtham S, Dutta R, **Baillie JK**, Digard P, Vervelde L, Burt DW, Smith J. Corrigendum: The molecular basis of differential host responses to avian influenza viruses in avian species with differing susceptibility. *Frontiers In Cellular and Infection Microbiology*(2023); 13: 1194878. [PMC10112380](#)

Parker E, Thomas J, Roper KJ, Ijaz S, Edwards T, Marchesin F, Katsanovskaja K, Lett L, Jones C, Hardwick HE, ..., **Baillie JK**, Openshaw PJM, Turtle L, Ho A, Semple MG, Paxton WA, Tedder RS, Pollakis G. SARS-CoV-2 antibody responses associate with sex, age and disease severity in previously uninfected people admitted to hospital with COVID-19: An ISARIC4C prospective study. *Frontiers In Immunology*(2023); 14: 1146702. [PMC10087108](#)

Morfopoulou S, Buddle S, Montaguth OET, Atkinson L, Guerra-Assunção JA, Marjaneh MM, Chiozzi RZ, Storey N, Campos L, Hutchinson JC, ..., **Baillie JK**, Semple MG, Martin J, Ushiro-Lumb I, Noursadeghi M, Deheragoda M, Hadzic N, Grammatikopoulos T, Brown R, Kelgeri C, Thalassinos K, Waddington SN, Jacques TS, Thomson E, Levin M, Brown JR, Breuer J. Genomic investigations of unexplained acute hepatitis in children. *Nature*(2023); doi:10.1038/s41586-023-06003-w

Ho A, Orton R, Tayler R, Asamaphan P, Herder V, Davis C, Tong L, Smollett K, Manali M, Allan J, ..., **Baillie JK**, Thomson EC. Adeno-associated virus 2 infection in children with non-A-E hepatitis. *Nature*(2023); doi:10.1038/s41586-023-05948-2

Morris KM, Mishra A, Raut AA, Gaunt ER, Borowska D, Kuo RI, Wang B, Vijayakumar P, Chingtham S, Dutta R, **Baillie JK**, Digard P, Vervelde L, Burt DW, Smith J. The molecular basis of differential host responses to avian influenza viruses in avian species with differing susceptibility. *Frontiers In Cellular and Infection Microbiology*(2023); 13: 1067993. PMC10011077

Goldswain H, Dong X, Penrice-Randal R, Alruwaili M, Shawli GT, Prince T, Williamson MK, Raghwani J, Randle N, Jones B, ..., **Baillie JK**, Emmett SR, Digard P, Matthews DA, Turtle L, Darby AC, Davidson AD, Carroll MW, Hiscox JA. The P323L substitution in the SARS-CoV-2 polymerase (NSP12) confers a selective advantage during infection. *Genome Biology*(2023); 24: 47. PMC10009825

Hamilton F, Mentzer AJ, Parks T, **Baillie JK**, Smith GD, Ghazal P, Timpson NJ. Variation in ERAP2 has opposing effects on severe respiratory infection and autoimmune disease. *American Journal of Human Genetics*(2023); doi:10.1016/j.ajhg.2023.02.008

Kartsonaki C, **Baillie JK**, Barrio NG, Baruch J, Beane A, Blumberg L, Bozza F, Broadley T, Burrell A, Carson G, Citarella BW, ..., Olliario PL. Characteristics and outcomes of an international cohort of 600,000 hospitalized patients with COVID-19. *International Journal of Epidemiology*(2023); doi:10.1093/ije/dyad012

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Ryan EM, Sadiku P, Coelho P, Watts ER, Zhang A, Howden AJM, Sanchez-Garcia MA, Bewley M, Cole J, McHugh BJ, ..., **Baillie JK**, Cantrell DA, McCafferty J, Choudhury G, Singh D, Dockrell DH, Whyte MKB, Walmsley SR. NRF2 Activation Reprogrammes Defects in Oxidative Metabolism to Restore Macrophage Function in COPD. *American Journal of Respiratory and Critical Care Medicine*(2023); doi:10.1164/rccm.202203-0482OC

Liew F, Talwar S, Cross A, Willett BJ, Scott S, Logan N, Siggins MK, Swieboda D, Sidhu JK, Efsthathiou C, ..., **Baillie JK**, Semple MG, Brightling C, Thwaites RS, Turtle L, Openshaw PJM. SARS-CoV-2-specific nasal IgA wanes 9 months after hospitalisation with COVID-19 and is not induced by subsequent vaccination. *Ebiomedicine*(2023); 87: 104402. PMC9762734

Hamilton FW, Thomas M, Arnold D, Palmer T, Moran E, Mentzer AJ, Maskell N, **Baillie JK**, Summers C, Hingorani A, MacGowan A, ..., Timpson NJ. Therapeutic potential of IL6R blockade for the treatment of sepsis and sepsis-related death: A Mendelian randomisation study. *PLoS Medicine*(2023); 20: e1004174. PMC9925069

Vink E, Davis C, MacLean A, Pascall D, McDonald SE, Gunson R, Hardwick HE, Oosthuyzen W, Openshaw PJM, **Baillie JK**, Semple MG, Ho A. Viral Coinfections in Hospitalized Coronavirus Disease 2019 Patients Recruited to the International Severe Acute Respiratory and Emerging Infections Consortium WHO Clinical Characterisation Protocol UK Study. *Open Forum Infectious Diseases*(2022); 9: ofac531. PMC9619746

Swann OV, Lone NI, Harrison EM, Tomlinson LA, Walker AJ, Seaborne MJ, Pollock L, Farrell J, Hall PS, Seth S, ..., **Baillie JK**, Katikireddi SV, Akbari A, Lyons R, Simpson CR, Semple MG, Goldacre B, Brophy S, Sheikh A, Docherty AB. Studying the Long-term Impact of COVID-19 in Kids (SLICK). Healthcare use and costs in children and young people following community-acquired SARS-CoV-2 infection: protocol for an observational study using linked primary and secondary routinely collected healthcare data from England, Scotland and Wales. *BMJ Open*(2022); 12: e063271. PMC9659708

Stewart I, Jacob J, George PM, Molyneaux PL, Porter JC, Allen RJ, Aslani S, **Baillie JK**, Barratt SL, Beirne P, Bianchi SM, ..., Jenkins RG. Residual Lung Abnormalities Following COVID-19 Hospitalization: Interim Analysis of the UKILD Post-COVID Study. *American Journal of Respiratory and Critical Care Medicine*(2022); doi:10.1164/rccm.202203-0564OC

Jefferson E, Cole C, Mumtaz S, Cox S, Giles T, Adejumo S, Urwin E, Lea D, McDonald C, Best J, ..., **Baillie JK**, Tang M, Leeming G, Power L, Breeze T, Gilson N, Murray DJ, Orton C, Pierce I, Hall I, Ladhani S, Whitaker M, Shallcross L, Seymour D, Varma S, Reilly G, Morris A, Hopkins S, Sheikh A, Quinlan P. CO-CONNECT: A hybrid architecture to facilitate rapid discovery and access to UK wide data in the response to the COVID-19 pandemic. *Journal of Medical Internet Research*(2022); doi:10.2196/40035

Grundmann A, Wu C, Hardwick M, **Baillie JK**, Openshaw PJM, Semple MG, Böhning D, Pett S, Michael BD, Thomas RH, Galea I. Fewer COVID-19 Neurological Complications with Dexamethasone and Remdesivir. *Annals of Neurology*(2022); doi:10.1002/ana.26536

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Butler-Laporte G, Povysil G, Kosmicki JA, Cirulli ET, Drivas T, Furini S, Saad C, Schmidt A, Olszewski P, Korotko U, Quinodoz M, Çelik E, Kundu K, Walter K, Jung J, Stockwell AD, Sloofman LG, Jordan DM, Thompson RC, Del Valle D, Simons N, Cheng E, Sebra R, Schadt EE, Kim-Schulze S, Gnjatic S, Merad M, Buxbaum JD, Beckmann ND, Charney AW, Przychodzen B, Chang T, Pottinger TD, Shang N, Brand F, Fava F, Mari F, Chwialkowska K, Niemira M, Pula S, **Baillie JK**, ..., Richards JB. Exome-wide association study to identify rare variants influencing COVID-19 outcomes: Results from the Host Genetics Initiative. *PLoS Genetics*(2022); 18: e1010367. PMC9632827

Siggins MK, Davies K, Fellows R, Thwaites RS, **Baillie JK**, Semple MG, Openshaw PJM, Zelek WM, Harris CL, Morgan BP. Alternative pathway dysregulation in tissues drives sustained complement activation and predicts outcome across the disease course in COVID-19. *Immunology*(2022); doi:10.1111/imm.13585

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SELECTED INVITED LECTURES

"RECOVERY: Dexamethasone is an effective treatment for Covid-19" Opening plenary, European Society for Intensive Care Medicine Annual Meeting, November 2020

"Optimisation and meta-analysis of genome-scale signals" CRISPR and beyond: perturbations at scale to understand genomes, Wellcome Trust Genome Campus, Hinxton, UK, September 2020

"Genetic effects on outcomes in sepsis: What mechanisms are amenable to treatment" World Congress of Intensive Care, Melbourne, October 2019

"EBOLA / MERS / influenza pandemics" World Congress of Intensive Care, Melbourne, October 2019

"Distinguishing adaptive from maladaptive processes in sepsis." Sepsis 2019, Rio de Janeiro, May 2019

"Extremes of human physiology - sorting out adaptive and maladaptive processes" Futuristic Medicine, Royal College of Physicians of Edinburgh, Feb 2019

“Transcriptomic biomarkers of critical illness” British Pharmacology Society Annual Meeting, London, December 2018

“Genomic stratification in critical illness” Intensive Care Society State of the Art Meeting, London, December 2018

“Genome editing studies of severe influenza” British Thoracic Society Annual Meeting, London, December 2018

“Gene-level target validation in large animals” One Health Symposium, University of Edinburgh, October 2018

“Understanding disease pathogenesis through functional genomics” RIKEN institute seminar, Yokohama, October 2018

“Biological endotypes in sepsis” Sepsis 2018, Bangkok, October 2018

“Sepsis: a syndrome simultaneously too broad and too narrow” Sepsis update meeting, Glasgow, September 2018

“Preparedness and management of global pandemics.” European Society of Intensive Care Medicine NEXT Webinar, September 2018

“Viral host genomic interactions.” Cambridge Centre for Cardiovascular and Respiratory Research Seminar, Cambridge, September 2018

“Applications of functional genomics and genome editing in post-GWAS analyses” BRIC institute seminar, Copenhagen, May 2018

“Precision medicine through genome-scale technologies” China-UK Life Sciences Summit, Beijing, September 2017

“Data-driven identification of syndromes at high altitude” 10th Kno2wledge meeting, London, May 2017

Public lecture: *“Starved of Oxygen”* Pint of Science, Edinburgh, May 2017

“Sepsis - what underlies individual susceptibility?” Singapore-ANZICS Research Meeting, Singapore, March 2015

“International collaboration in research during outbreaks” Singapore-ANZICS Research Meeting, Singapore, March 2015

“Host genetic susceptibility to influenza - moving towards therapeutic targets” Broad Institute Seminar Series on Infectious Diseases, Cambridge (MA), January 2015

“Why do some people get sepsis while others don’t?” Intensive Care Society State of the Art Meeting, London, December 2014.

“Host genetics of influenza susceptibility” Keystone Symposium on Pathogenesis of Respiratory Viruses, Keystone, Colorado, January 2014.

“Redefining Acute Mountain Sickness” Session chair and speaker, World Congress on Mountain Medicine, Bolzano 2014.

“Virus:host interactions: time to focus on the host?” ANZICS Global Rising Star Fellowship Session, Hobart, Tasmania, October 2013

“Genomics in Anaesthesia and Critical Care” Royal College of Anaesthetists Jubilee Meeting, *“Rising star”* Session. October 2013

“Large scale biological research in critical care” World Federation of Intensive and Critical Care Societies 11th Congress, Durban, South Africa, August 2013

“Using coexpression to maximise statistical power in genome-wide studies” Computational Genomics Analysis and Training Programme, Oxford, Feb 2012

“Challenges in Severe Acute Respiratory Illness” European Society of Clinical Microbiology and Infectious Diseases, Barcelona, Jan 2012

Co-chair, Biological Studies group. InFACT/Li Ka Shing Knowledge Institute/ISARIC colloquium on SARI research, Toronto, June 2011.

“Host genetic studies” NIH Flu002/Flu003 Analysis planning meeting, Boston, Oct 2011.