James M Mccaw

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

116
papers2,219
citations25
h-index42
g-index130
ext. papers2,740
ext. citations5.4
avg, IF4.92
L-index

#	Paper	IF	Citations
116	Hypnozoite dynamics for Plasmodium vivax malaria: the epidemiological effects of radical cure <i>Journal of Theoretical Biology</i> , 2022 , 537, 111014	2.3	O
115	Estimation of the probability of epidemic fade-out from multiple outbreak data <i>Epidemics</i> , 2022 , 38, 100539	5.1	О
114	Rapid assessment of the risk of SARS-CoV-2 importation: case study and lessons learned <i>Epidemics</i> , 2022 , 38, 100549	5.1	O
113	COVID-19 in low-tolerance border quarantine systems: Impact of the Delta variant of SARS-CoV-2 <i>Science Advances</i> , 2022 , 8, eabm3624	14.3	1
112	Modelling the Effect of MUC1 on Influenza Virus Infection Kinetics and Macrophage Dynamics. <i>Viruses</i> , 2021 , 13,	6.2	2
111	Modelling within-host macrophage dynamics in influenza virus infection. <i>Journal of Theoretical Biology</i> , 2021 , 508, 110492	2.3	5
110	Antibody Dynamics for Plasmodium vivax Malaria: A Mathematical Model. <i>Bulletin of Mathematical Biology</i> , 2021 , 83, 6	2.1	3
109	Constructing an ethical framework for priority allocation of pandemic vaccines. <i>Vaccine</i> , 2021 , 39, 797-8	8 0 µ41	3
108	Development of an influenza pandemic decision support tool linking situational analytics to national response policy. <i>Epidemics</i> , 2021 , 36, 100478	5.1	O
107	An Activation-Clearance Model for Plasmodium vivax Malaria. <i>Bulletin of Mathematical Biology</i> , 2020 , 82, 32	2.1	3
106	Coordinating the real-time use of global influenza activity data for better public health planning. <i>Influenza and Other Respiratory Viruses</i> , 2020 , 14, 105-110	5.6	2
105	Early analysis of the Australian COVID-19 epidemic. <i>ELife</i> , 2020 , 9,	8.9	31
104	Author response: Early analysis of the Australian COVID-19 epidemic 2020 ,		3
103	Infectious disease pandemic planning and response: Incorporating decision analysis. <i>PLoS Medicine</i> , 2020 , 17, e1003018	11.6	37
102	Estimation of the force of infection and infectious period of skin sores in remote Australian communities using interval-censored data. <i>PLoS Computational Biology</i> , 2020 , 16, e1007838	5	1
101	Influencing public health policy with data-informed mathematical models of infectious diseases: Recent developments and new challenges. <i>Epidemics</i> , 2020 , 32, 100393	5.1	14
100	Coronavirus Disease Model to Inform Transmission-Reducing Measures and Health System Preparedness, Australia. <i>Emerging Infectious Diseases</i> , 2020 , 26, 2844-2853	10.2	12

(2018-2020)

99	Key questions for modelling COVID-19 exit strategies. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2020 , 287, 20201405	4.4	65
98	Estimation of the force of infection and infectious period of skin sores in remote Australian communities using interval-censored data 2020 , 16, e1007838		
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96	Estimation of the force of infection and infectious period of skin sores in remote Australian communities using interval-censored data 2020 , 16, e1007838		
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93	Estimation of the force of infection and infectious period of skin sores in remote Australian communities using interval-censored data 2020 , 16, e1007838		
92	Sequential infection experiments for quantifying innate and adaptive immunity during influenza infection. <i>PLoS Computational Biology</i> , 2019 , 15, e1006568	5	5
91	Accounting for Healthcare-Seeking Behaviours and Testing Practices in Real-Time Influenza Forecasts. <i>Tropical Medicine and Infectious Disease</i> , 2019 , 4,	3.5	8
90	Modeling the dynamics of gametocytes in humans during malaria infection. ELife, 2019, 8,	8.9	15
89	A simple influenza model with complicated dynamics. <i>Journal of Mathematical Biology</i> , 2019 , 78, 607-62	24	4
88	A biological model of scabies infection dynamics and treatment informs mass drug administration strategies to increase the likelihood of elimination. <i>Mathematical Biosciences</i> , 2019 , 309, 163-173	3.9	10
87	Anatomy of a seasonal influenza epidemic forecast. <i>Communicable Diseases Intelligence (2018)</i> , 2019 , 43,	1.9	3
86	Clonally diverse CD38HLA-DRCD8 T cells persist during fatal H7N9 disease. <i>Nature Communications</i> , 2018 , 9, 824	17.4	69
85	Evidence for Viral Interference and Cross-reactive Protective Immunity Between Influenza B Virus Lineages. <i>Journal of Infectious Diseases</i> , 2018 , 217, 548-559	7	36
84	Epidemic forecasts as a tool for public health: interpretation and (re)calibration. <i>Australian and New Zealand Journal of Public Health</i> , 2018 , 42, 69-76	2.3	13
83	The distribution of the time taken for an epidemic to spread between two communities. <i>Mathematical Biosciences</i> , 2018 , 303, 139-147	3.9	3
82	Investigating Viral Interference Between Influenza A Virus and Human Respiratory Syncytial Virus in a Ferret Model of Infection. <i>Journal of Infectious Diseases</i> , 2018 , 218, 406-417	7	41

81	Within-host modeling of blood-stage malaria. <i>Immunological Reviews</i> , 2018 , 285, 168-193	11.3	17
80	New Mathematical Models of Antimalarial Drug Action to Improve Drug Dosing Regimens. <i>Mathematics for Industry</i> , 2018 , 7-11	0.1	
79	Investigation of the Decline in Clinical Efficacy of Artemisinin Combination Therapies Due to Increasing Artemisinin and Partner Drug Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2018 , 62,	5.9	3
78	Predicting the Outcomes of New Short-Course Regimens for Multidrug-Resistant Tuberculosis Using Intrahost and Pharmacokinetic-Pharmacodynamic Modeling. <i>Antimicrobial Agents and Chemotherapy</i> , 2018 , 62,	5.9	6
77	Characterization of Influenza B Virus Variants with Reduced Neuraminidase Inhibitor Susceptibility. <i>Antimicrobial Agents and Chemotherapy</i> , 2018 , 62,	5.9	11
76	Investigating the Efficacy of Triple Artemisinin-Based Combination Therapies for Treating Plasmodium falciparum Malaria Patients Using Mathematical Modeling. <i>Antimicrobial Agents and Chemotherapy</i> , 2018 , 62,	5.9	32
75	Infection-acquired versus vaccine-acquired immunity in an SIRWS model. <i>Infectious Disease Modelling</i> , 2018 , 3, 118-135	15.7	4
74	Calculation of the age of the first infection for skin sores and scabies in five remote communities in northern Australia. <i>Epidemiology and Infection</i> , 2018 , 146, 1194-1201	4.3	8
73	Turnover of Village Chickens Undermines Vaccine Coverage to Control HPAI H5N1. <i>Zoonoses and Public Health</i> , 2017 , 64, 53-62	2.9	1
72	Characterising pandemic severity and transmissibility from data collected during first few hundred studies. <i>Epidemics</i> , 2017 , 19, 61-73	5.1	23
71	Influenza as a trigger for cardiovascular disease: An investigation of serotype, subtype and geographic location. <i>Environmental Research</i> , 2017 , 156, 688-696	7.9	11
70	Model selection for seasonal influenza forecasting. <i>Infectious Disease Modelling</i> , 2017 , 2, 56-70	15.7	13
69	Retrospective forecasting of the 2010-2014 Melbourne influenza seasons using multiple surveillance systems. <i>Epidemiology and Infection</i> , 2017 , 145, 156-169	4.3	18
68	A Dynamic Stress Model Explains the Delayed Drug Effect in Artemisinin Treatment of Plasmodium falciparum. <i>Antimicrobial Agents and Chemotherapy</i> , 2017 , 61,	5.9	5
67	A mechanistic model quantifies artemisinin-induced parasite growth retardation in blood-stage Plasmodium falciparum infection. <i>Journal of Theoretical Biology</i> , 2017 , 430, 117-127	2.3	6
66	Modelling cross-reactivity and memory in the cellular adaptive immune response to influenza infection in the host. <i>Journal of Theoretical Biology</i> , 2017 , 413, 34-49	2.3	15
65	The Mechanisms for Within-Host Influenza Virus Control Affect Model-Based Assessment and Prediction of Antiviral Treatment. <i>Viruses</i> , 2017 , 9,	6.2	20
64	Periodic solutions in an SIRWS model with immune boosting and cross-immunity. <i>Journal of Theoretical Biology</i> , 2016 , 410, 55-64	2.3	4

(2015-2016)

63	Reducing disease burden in an influenza pandemic by targeted delivery of neuraminidase inhibitors: mathematical models in the Australian context. <i>BMC Infectious Diseases</i> , 2016 , 16, 552	4	10
62	High conservation level of CD8(+) T cell immunogenic regions within an unusual H1N2 human influenza variant. <i>Journal of Medical Virology</i> , 2016 , 88, 1725-32	19.7	3
61	Heightened self-reactivity associated with selective survival, but not expansion, of nalle virus-specific CD8+ T cells in aged mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 1333-8	11.5	40
60	Parasite Strain, Host Immunity, and Circulating Blood Cells with Dead Parasites: Why Predicting Malaria Parasite Clearance Is Not a Simple Task. <i>Antimicrobial Agents and Chemotherapy</i> , 2016 , 60, 1172	5.9	2
59	On the extinction probability in models of within-host infection: the role of latency and immunity. Journal of Mathematical Biology, 2016 , 73, 787-813	2	13
58	Model-Informed Risk Assessment and Decision Making for an Emerging Infectious Disease in the Asia-Pacific Region. <i>PLoS Neglected Tropical Diseases</i> , 2016 , 10, e0005018	4.8	8
57	On the Role of CD8 T Cells in Determining Recovery Time from Influenza Virus Infection. <i>Frontiers in Immunology</i> , 2016 , 7, 611	8.4	20
56	Comparison of the Exposure Time Dependence of the Activities of Synthetic Ozonide Antimalarials and Dihydroartemisinin against K13 Wild-Type and Mutant Plasmodium falciparum Strains. Antimicrobial Agents and Chemotherapy, 2016, 60, 4501-10	5.9	38
55	Forecasting influenza outbreak dynamics in Melbourne from Internet search query surveillance data. <i>Influenza and Other Respiratory Viruses</i> , 2016 , 10, 314-23	5.6	24
54	Quantifying differences in the epidemic curves from three influenza surveillance systems: a nonlinear regression analysis. <i>Epidemiology and Infection</i> , 2015 , 143, 427-39	4.3	4
53	Quantifying relative within-host replication fitness in influenza virus competition experiments. <i>Journal of Theoretical Biology</i> , 2015 , 382, 259-71	2.3	14
52	Targeting the cell stress response of Plasmodium falciparum to overcome artemisinin resistance. <i>PLoS Biology</i> , 2015 , 13, e1002132	9.7	193
51	A new approach to estimating trends in chlamydia incidence. <i>Sexually Transmitted Infections</i> , 2015 , 91, 513-9	2.8	10
50	Interval Between Infections and Viral Hierarchy Are Determinants of Viral Interference Following Influenza Virus Infection in a Ferret Model. <i>Journal of Infectious Diseases</i> , 2015 , 212, 1701-10	7	62
49	The effects of demographic change on disease transmission and vaccine impact in a household structured population. <i>Epidemics</i> , 2015 , 13, 56-64	5.1	36
48	Defining long-term drivers of pertussis resurgence, and optimal vaccine control strategies. <i>Vaccine</i> , 2015 , 33, 5794-5800	4.1	19
47	Factors associated with transmission of influenza-like illness in a cohort of households containing multiple children. <i>Influenza and Other Respiratory Viruses</i> , 2015 , 9, 247-54	5.6	8
46	Social encounter profiles of greater Melbourne residents, by locationa telephone survey. <i>BMC Infectious Diseases</i> , 2015 , 15, 494	4	14

45	Prior population immunity reduces the expected impact of CTL-inducing vaccines for pandemic influenza control. <i>PLoS ONE</i> , 2015 , 10, e0120138	3.7	9
44	Pertussis models to inform vaccine policy. <i>Human Vaccines and Immunotherapeutics</i> , 2015 , 11, 669-78	4.4	8
43	Innate Immunity and the Inter-exposure Interval Determine the Dynamics of Secondary Influenza Virus Infection and Explain Observed Viral Hierarchies. <i>PLoS Computational Biology</i> , 2015 , 11, e1004334	, 5	38
42	The dynamical consequences of seasonal forcing, immune boosting and demographic change in a model of disease transmission. <i>Journal of Theoretical Biology</i> , 2014 , 361, 124-32	2.3	8
41	The influence of changing host immunity on 1918-19 pandemic dynamics. <i>Epidemics</i> , 2014 , 8, 18-27	5.1	7
40	Making the most of clinical data: reviewing the role of pharmacokinetic-pharmacodynamic models of anti-malarial drugs. <i>AAPS Journal</i> , 2014 , 16, 962-74	3.7	19
39	Population pharmacokinetics of intravenous artesunate: a pooled analysis of individual data from patients with severe malaria. <i>CPT: Pharmacometrics and Systems Pharmacology</i> , 2014 , 3, e145	4.5	12
38	Evaluation of oseltamivir prophylaxis regimens for reducing influenza virus infection, transmission and disease severity in a ferret model of household contact. <i>Journal of Antimicrobial Chemotherapy</i> , 2014 , 69, 2458-69	5.1	24
37	Pandemic controllability: a concept to guide a proportionate and flexible operational response to future influenza pandemics. <i>Journal of Public Health</i> , 2014 , 36, 5-12	3.5	13
36	Estimating the fitness advantage conferred by permissive neuraminidase mutations in recent oseltamivir-resistant A(H1N1)pdm09 influenza viruses. <i>PLoS Pathogens</i> , 2014 , 10, e1004065	7.6	97
35	Dynamical crises, multistability and the influence of the duration of immunity in a seasonally-forced model of disease transmission. <i>Theoretical Biology and Medical Modelling</i> , 2014 , 11, 43	2.3	6
34	Virus detection and its association with symptoms during influenza-like illness in a sample of healthy adults enrolled in a randomised controlled vaccine trial. <i>Influenza and Other Respiratory Viruses</i> , 2013 , 7, 330-9	5.6	14
33	Antigenic drift of the pandemic 2009 A(H1N1) influenza virus in A ferret model. <i>PLoS Pathogens</i> , 2013 , 9, e1003354	7.6	52
32	Altered temporal response of malaria parasites determines differential sensitivity to artemisinin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 5157-62	11.5	139
31	Reducing uncertainty in within-host parameter estimates of influenza infection by measuring both infectious and total viral load. <i>PLoS ONE</i> , 2013 , 8, e64098	3.7	25
30	Synthetic Population Dynamics: A Model of Household Demography. <i>Jasss</i> , 2013 , 16,	4.8	36
29	Drivers and consequences of influenza antiviral resistant-strain emergence in a capacity-constrained pandemic response. <i>Epidemics</i> , 2012 , 4, 219-26	5.1	4
28	Household transmission of respiratory viruses - assessment of viral, individual and household characteristics in a population study of healthy Australian adults. <i>BMC Infectious Diseases</i> , 2012 , 12, 345	4	14

(2009-2012)

27	Application of a case-control study design to investigate genotypic signatures of HIV-1 transmission. <i>Retrovirology</i> , 2012 , 9, 54	3.6	5
26	Assessing the utility of an anti-malarial pharmacokinetic-pharmacodynamic model for aiding drug clinical development. <i>Malaria Journal</i> , 2012 , 11, 303	3.6	33
25	Influence of contact definitions in assessment of the relative importance of social settings in disease transmission risk. <i>PLoS ONE</i> , 2012 , 7, e30893	3.7	11
24	Likely effectiveness of pharmaceutical and non-pharmaceutical interventions for mitigating influenza virus transmission in Mongolia. <i>Bulletin of the World Health Organization</i> , 2012 , 90, 264-71	8.2	21
23	THE INFLUENCE OF INCREASING LIFE EXPECTANCY ON THE DYNAMICS OF SIRS SYSTEMS WITH IMMUNE BOOSTING. <i>ANZIAM Journal</i> , 2012 , 54, 50-63	0.5	14
22	H1N1 influenza and the Australian macroeconomy. <i>Journal of the Asia Pacific Economy</i> , 2012 , 17, 22-51	1	16
21	Incorporating population dynamics into household models of infectious disease transmission. <i>Epidemics</i> , 2011 , 3, 152-8	5.1	15
20	Diagnosis and antiviral intervention strategies for mitigating an influenza epidemic. <i>PLoS ONE</i> , 2011 , 6, e14505	3.7	19
19	Understanding mortality in the 1918-1919 influenza pandemic in England and Wales. <i>Influenza and Other Respiratory Viruses</i> , 2011 , 5, 89-98	5.6	11
18	A mathematical framework for estimating pathogen transmission fitness and inoculum size using data from a competitive mixtures animal model. <i>PLoS Computational Biology</i> , 2011 , 7, e1002026	5	15
17	Modelling strategic use of the national antiviral stockpile during the CONTAIN and SUSTAIN phases of an Australian pandemic influenza response. <i>Australian and New Zealand Journal of Public Health</i> , 2010 , 34, 113-9	2.3	17
16	Assessing the viral fitness of oseltamivir-resistant influenza viruses in ferrets, using a competitive-mixtures model. <i>Journal of Virology</i> , 2010 , 84, 9427-38	6.6	56
15	Prior immunity helps to explain wave-like behaviour of pandemic influenza in 1918-9. <i>BMC Infectious Diseases</i> , 2010 , 10, 128	4	37
14	Comparison of three methods for ascertainment of contact information relevant to respiratory pathogen transmission in encounter networks. <i>BMC Infectious Diseases</i> , 2010 , 10, 166	4	42
13	Influenza: accounting for prior immunity. <i>Science</i> , 2009 , 325, 1071; author reply 1072-3	33.3	24
12	Optimal dosing and dynamic distribution of vaccines in an influenza pandemic. <i>American Journal of Epidemiology</i> , 2009 , 169, 1517-24	3.8	11
11	Understanding influenza transmission, immunity and pandemic threats. <i>Influenza and Other Respiratory Viruses</i> , 2009 , 3, 143-9	5.6	58
10	Understanding Australias influenza pandemic policy on the strategic use of the antiviral drug stockpile. <i>Medical Journal of Australia</i> , 2009 , 191, 136-7	4	4

9	Impact of emerging antiviral drug resistance on influenza containment and spread: influence of subclinical infection and strategic use of a stockpile containing one or two drugs. <i>PLoS ONE</i> , 2008 , 3, e2362	3.7	43	
8	A biological model for influenza transmission: pandemic planning implications of asymptomatic infection and immunity. <i>PLoS ONE</i> , 2007 , 2, e1220	3.7	62	
7	Prophylaxis or treatment? Optimal use of an antiviral stockpile during an influenza pandemic. <i>Mathematical Biosciences</i> , 2007 , 209, 336-60	3.9	55	
6	Pure point spectrum for the time evolution of a periodically rank-N kicked Hamiltonian. <i>Journal of Mathematical Physics</i> , 2005 , 46, 032108	1.2	4	
5	On the continuous spectral component of the Floquet operator for a periodically kicked quantum system. <i>Journal of Mathematical Physics</i> , 2005 , 46, 103503	1.2	2	
4	Anatomy of a seasonal influenza epidemic forecast. Communicable Diseases Intelligence (2018),43,	1.9	4	
3	Modelling within-host macrophage dynamics in influenza virus infection		1	
2	Assessing the risk of spread of COVID-19 to the Asia Pacific region		4	
1	Estimation of the force of infection and infectious period of skin sores in remote Australian communities using interval-censored data		1	