

James R Price

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

550
citations

1040056

9
h-index

1281871

11
g-index

20
all docs

20
docs citations

20
times ranked

1078
citing authors

#	ARTICLE	IF	CITATIONS
1	SARS-CoV-2 lineage B.1.1.7 is associated with greater disease severity among hospitalised women but not men: multicentre cohort study. <i>BMJ Open Respiratory Research</i> , 2021, 8, e001029.	3.0	22
2	The Alpha variant was not associated with excess nosocomial SARS-CoV-2 infection in a multi-centre UK hospital study. <i>Journal of Infection</i> , 2021, 83, 693-700.	3.3	11
3	Joint Healthcare Infection Society (HIS) and Infection Prevention Society (IPS) guidelines for the prevention and control of methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) in healthcare facilities. <i>Journal of Hospital Infection</i> , 2021, 118, S1-S39.	2.9	14
4	Antimicrobial resistance determinants are associated with <i>Staphylococcus aureus</i> bacteraemia and adaptation to the healthcare environment: a bacterial genome-wide association study. <i>Microbial Genomics</i> , 2021, 7, .	2.0	15
5	Development and Delivery of a Real-time Hospital-onset COVID-19 Surveillance System Using Network Analysis. <i>Clinical Infectious Diseases</i> , 2020, 72, 82-89.	5.8	14
6	The TACTIC experience: establishing an international, interdisciplinary network to tackle antimicrobial resistance. <i>Journal of Medical Microbiology</i> , 2020, 69, 1213-1220.	1.8	1
7	Survival following <i>Staphylococcus aureus</i> bloodstream infection: A prospective multinational cohort study assessing the impact of place of care. <i>Journal of Infection</i> , 2018, 77, 516-525.	3.3	48
8	Re-emergence of methicillin susceptibility in a resistant lineage of <i>Staphylococcus aureus</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, dkw570.	3.0	22
9	<i>Staphylococcus aureus</i> in critical care – Authors' reply. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 580-581.	9.1	0
10	Transmission of <i>Staphylococcus aureus</i> between health-care workers, the environment, and patients in an intensive care unit: a longitudinal cohort study based on whole-genome sequencing. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 207-214.	9.1	155
11	Severe infections emerge from commensal bacteria by adaptive evolution. <i>ELife</i> , 2017, 6, .	6.0	93
12	Reply to Mills and Linkin. <i>Clinical Infectious Diseases</i> , 2014, 59, 752-753.	5.8	0
13	Whole-Genome Sequencing Shows That Patient-to-Patient Transmission Rarely Accounts for Acquisition of <i>Staphylococcus aureus</i> in an Intensive Care Unit. <i>Clinical Infectious Diseases</i> , 2014, 58, 609-618.	5.8	142