

Tjibbe Donker

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1927244/publications.pdf>

Version: 2024-02-01

21
papers

715
citations

686830

13
h-index

676716

22
g-index

28
all docs

28
docs citations

28
times ranked

1069
citing authors

#	ARTICLE	IF	CITATIONS
1	Impact of antibiotic use on patient-level risk of death in 36 million hospital admissions in England. <i>Journal of Infection</i> , 2022, 84, 311-320.	1.7	7
2	Navigating hospitals safely through the COVID-19 epidemic tide: Predicting case load for adjusting bed capacity. <i>Infection Control and Hospital Epidemiology</i> , 2021, 42, 653-658.	1.0	12
3	Modelling how antimicrobial resistance spreads between wards. <i>ELife</i> , 2020, 9, .	2.8	2
4	Using hospital network-based surveillance for antimicrobial resistance as a more robust alternative to self-reporting. <i>PLoS ONE</i> , 2019, 14, e0219994.	1.1	3
5	Mathematical modelling for antibiotic resistance control policy: do we know enough?. <i>BMC Infectious Diseases</i> , 2019, 19, 1011.	1.3	37
6	The relative importance of large problems far away versus small problems closer to home: insights into limiting the spread of antimicrobial resistance in England. <i>BMC Medicine</i> , 2017, 15, 86.	2.3	30
7	Measuring distance through dense weighted networks: The case of hospital-associated pathogens. <i>PLoS Computational Biology</i> , 2017, 13, e1005622.	1.5	8
8	Population genetic structuring of methicillin-resistant <i>Staphylococcus aureus</i> clone EMRSA-15 within UK reflects patient referral patterns. <i>Microbial Genomics</i> , 2017, 3, e000113.	1.0	19
9	Building a genomic framework for prospective MRSA surveillance in the United Kingdom and the Republic of Ireland. <i>Genome Research</i> , 2016, 26, 263-270.	2.4	63
10	Identifying the effect of patient sharing on between-hospital genetic differentiation of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Genome Medicine</i> , 2016, 8, 18.	3.6	20
11	Monitoring the spread of methicillin-resistant <i>Staphylococcus aureus</i> in The Netherlands from a reference laboratory perspective. <i>Journal of Hospital Infection</i> , 2016, 93, 366-374.	1.4	13
12	Dispersal of antibiotic-resistant high-risk clones by hospital networks: changing the patient direction can make all the difference. <i>Journal of Hospital Infection</i> , 2014, 86, 34-41.	1.4	31
13	Efficient surveillance for healthcare-associated infections spreading between hospitals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 2271-2276.	3.3	46
14	Infection prevention in a connected world: The case for a regional approach. <i>International Journal of Medical Microbiology</i> , 2013, 303, 380-387.	1.5	66
15	Estimating the Generation Interval of Influenza A (H1N1) in a Range of Social Settings. <i>Epidemiology</i> , 2013, 24, 244-250.	1.2	19
16	Finding Evidence for Local Transmission of Contagious Disease in Molecular Epidemiological Datasets. <i>PLoS ONE</i> , 2013, 8, e69875.	1.1	14
17	Mumps outbreak among vaccinated university students associated with a large party, the Netherlands, 2010. <i>Vaccine</i> , 2012, 30, 4676-4680.	1.7	58
18	Hospital Networks and the Dispersal of Hospital-Acquired Pathogens by Patient Transfer. <i>PLoS ONE</i> , 2012, 7, e35002.	1.1	97

#	ARTICLE	IF	CITATIONS
19	Nowcasting pandemic influenza A/H1N1 2009 hospitalizations in the Netherlands. <i>European Journal of Epidemiology</i> , 2011, 26, 195-201.	2.5	33
20	Transmission of Novel Influenza A(H1N1) in Households with Post-Exposure Antiviral Prophylaxis. <i>PLoS ONE</i> , 2010, 5, e11442.	1.1	29
21	Patient Referral Patterns and the Spread of Hospital-Acquired Infections through National Health Care Networks. <i>PLoS Computational Biology</i> , 2010, 6, e1000715.	1.5	102