

T Phuong Quan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1708693/publications.pdf>

Version: 2024-02-01

20
papers

1,404
citations

623574

14
h-index

752573

20
g-index

22
all docs

22
docs citations

22
times ranked

2603
citing authors

#	ARTICLE	IF	CITATIONS
1	Combining Charlson and Elixhauser scores with varying lookback predicated mortality better than using individual scores. <i>Journal of Clinical Epidemiology</i> , 2021, 130, 32-41.	2.4	3
2	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, .	1.0	15
3	Antimicrobial resistance in commensal opportunistic pathogens isolated from non-sterile sites can be an effective proxy for surveillance in bloodstream infections. <i>Scientific Reports</i> , 2021, 11, 23359.	1.6	2
4	Reconciling the Potentially Irreconcilable? Genotypic and Phenotypic Amoxicillin-Clavulanate Resistance in <i>Escherichia coli</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	1.4	33
5	Investigation of the impact of the NICE guidelines regarding antibiotic prophylaxis during invasive dental procedures on the incidence of infective endocarditis in England: an electronic health records study. <i>BMC Medicine</i> , 2020, 18, 84.	2.3	39
6	Genomic surveillance of <i>Escherichia coli</i> and <i>Klebsiella</i> spp. in hospital sink drains and patients. <i>Microbial Genomics</i> , 2020, 6, .	1.0	26
7	“Caveat emptor”: the cautionary tale of endocarditis and the potential pitfalls of clinical coding data—an electronic health records study. <i>BMC Medicine</i> , 2019, 17, 169.	2.3	25
8	A Role for Tetracycline Selection in Recent Evolution of Agriculture-Associated <i>Clostridium difficile</i> PCR Ribotype 078. <i>MBio</i> , 2019, 10, .	1.8	46
9	Using linked electronic health records to report healthcare-associated infections. <i>PLoS ONE</i> , 2018, 13, e0206860.	1.1	3
10	A <i>Candida auris</i> Outbreak and Its Control in an Intensive Care Setting. <i>New England Journal of Medicine</i> , 2018, 379, 1322-1331.	13.9	318
11	Trends over time in <i>Escherichia coli</i> bloodstream infections, urinary tract infections, and antibiotic susceptibilities in Oxfordshire, UK, 1998–2016: a study of electronic health records. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 1138-1149.	4.6	121
12	Evaluation of Whole-Genome Sequencing for Mycobacterial Species Identification and Drug Susceptibility Testing in a Clinical Setting: a Large-Scale Prospective Assessment of Performance against Line Probe Assays and Phenotyping. <i>Journal of Clinical Microbiology</i> , 2018, 56, .	1.8	72
13	Effects of control interventions on <i>Clostridium difficile</i> infection in England: an observational study. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 411-421.	4.6	269
14	Mortality risks associated with emergency admissions during weekends and public holidays: an analysis of electronic health records. <i>Lancet</i> , The, 2017, 390, 62-72.	6.3	114
15	<i>Clostridium difficile</i> in England: can we stop washing our hands? “ Authors' reply. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 478-479.	4.6	1
16	Contribution to <i>Clostridium Difficile</i> Transmission of Symptomatic Patients With Toxigenic Strains Who Are Fecal Toxin Negative. <i>Clinical Infectious Diseases</i> , 2017, 64, 1163-1170.	2.9	45
17	Epidemiology of <i>Clostridium difficile</i> in infants in Oxfordshire, UK: Risk factors for colonization and carriage, and genetic overlap with regional <i>C. difficile</i> infection strains. <i>PLoS ONE</i> , 2017, 12, e0182307.	1.1	82
18	Antibiotic use and clinical outcomes in the acute setting under management by an infectious diseases acute physician versus other clinical teams: a cohort study. <i>BMJ Open</i> , 2016, 6, e010969.	0.8	8

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19	Increasing burden of community-acquired pneumonia leading to hospitalisation, 1998â€“2014. <i>Thorax</i> , 2016, 71, 535-542.	2.7	80
20	Mycobacterial DNA Extraction for Whole-Genome Sequencing from Early Positive Liquid (MGIT) Cultures. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1137-1143.	1.8	90