

Nicole Stoesser

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4554113/publications.pdf>

Version: 2024-02-01

138
papers

11,235
citations

47409

49
h-index

46524

93
g-index

206
all docs

206
docs citations

206
times ranked

16878
citing authors

#	ARTICLE	IF	CITATIONS
1	Symptoms and Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Positivity in the General Population in the United Kingdom. <i>Clinical Infectious Diseases</i> , 2022, 75, e329-e337.	2.9	20
2	Tocilizumab for treatment of SARS-CoV-2 infection at home: A case report. <i>Acute Medicine</i> , 2022, 21, 53-55.	0.1	1
3	Antibody responses and correlates of protection in the general population after two doses of the ChAdOx1 or BNT162b2 vaccines. <i>Nature Medicine</i> , 2022, 28, 1072-1082.	15.2	147
4	Systematic review of wastewater surveillance of antimicrobial resistance in human populations. <i>Environment International</i> , 2022, 162, 107171.	4.8	29
5	Hospital outbreak of carbapenem-resistant Enterobacterales associated with a bla OXA-48 plasmid carried mostly by <i>Escherichia coli</i> ST399. <i>Microbial Genomics</i> , 2022, 8, .	1.0	3
6	Prediction of Antibiotic Resistance Evolution by Growth Measurement of All Proximal Mutants of Beta-Lactamase. <i>Molecular Biology and Evolution</i> , 2022, 39, .	3.5	3
7	Whole genome sequencing reveals hidden transmission of carbapenemase-producing Enterobacterales. <i>Nature Communications</i> , 2022, 13, .	5.8	16
8	SARS-CoV-2 antibody trajectories after a single COVID-19 vaccination with and without prior infection. <i>Nature Communications</i> , 2022, 13, .	5.8	6
9	Antibody Status and Incidence of SARS-CoV-2 Infection in Health Care Workers. <i>New England Journal of Medicine</i> , 2021, 384, 533-540.	13.9	803
10	Community prevalence of SARS-CoV-2 in England from April to November, 2020: results from the ONS Coronavirus Infection Survey. <i>Lancet Public Health</i> , The, 2021, 6, e30-e38.	4.7	147
11	Ten Years of Population-Level Genomic <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> Serotype Surveillance Informs Vaccine Development for Invasive Infections. <i>Clinical Infectious Diseases</i> , 2021, 73, 2276-2282.	2.9	16
12	Stringent thresholds in SARS-CoV-2 IgG assays lead to under-detection of mild infections. <i>BMC Infectious Diseases</i> , 2021, 21, 187.	1.3	23
13	Amplification-Free Detection of Viruses in Minutes using Single-Particle Imaging and Machine Learning. <i>Biophysical Journal</i> , 2021, 120, 195a.	0.2	1
14	A haemagglutination test for rapid detection of antibodies to SARS-CoV-2. <i>Nature Communications</i> , 2021, 12, 1951.	5.8	54
15	Genomic network analysis of environmental and livestock F-type plasmid populations. <i>ISME Journal</i> , 2021, 15, 2322-2335.	4.4	24
16	Niche and local geography shape the pangenome of wastewater- and livestock-associated Enterobacteriaceae. <i>Science Advances</i> , 2021, 7, .	4.7	47
17	Diagnosis of SARS-CoV-2 Infection with LamPORE, a High-Throughput Platform Combining Loop-Mediated Isothermal Amplification and Nanopore Sequencing. <i>Journal of Clinical Microbiology</i> , 2021, 59, .	1.8	30
18	Quantitative SARS-CoV-2 anti-spike responses to Pfizerâ€BioNTech and Oxfordâ€AstraZeneca vaccines by previous infection status. <i>Clinical Microbiology and Infection</i> , 2021, 27, 1516.e7-1516.e14.	2.8	100

#	ARTICLE	IF	CITATIONS
19	Population-level faecal metagenomic profiling as a tool to predict antimicrobial resistance in Enterobacterales isolates causing invasive infections: An exploratory study across Cambodia, Kenya, and the UK. <i>EClinicalMedicine</i> , 2021, 36, 100910.	3.2	10
20	COVID-19: Rapid antigen detection for SARS-CoV-2 by lateral flow assay: A national systematic evaluation of sensitivity and specificity for mass-testing. <i>EClinicalMedicine</i> , 2021, 36, 100924.	3.2	162
21	Impact of vaccination on new SARS-CoV-2 infections in the United Kingdom. <i>Nature Medicine</i> , 2021, 27, 1370-1378.	15.2	260
22	Ct threshold values, a proxy for viral load in community SARS-CoV-2 cases, demonstrate wide variation across populations and over time. <i>ELife</i> , 2021, 10, .	2.8	91
23	Antibody responses to SARS-CoV-2 vaccines in 45,965 adults from the general population of the United Kingdom. <i>Nature Microbiology</i> , 2021, 6, 1140-1149.	5.9	254
24	Flanker: a tool for comparative genomics of gene flanking regions. <i>Microbial Genomics</i> , 2021, 7, .	1.0	12
25	Ten-year longitudinal molecular epidemiology study of <i>Escherichia coli</i> and <i>Klebsiella</i> species bloodstream infections in Oxfordshire, UK. <i>Genome Medicine</i> , 2021, 13, 144.	3.6	35
26	Antimicrobial resistance genes and clonal success in <i>Escherichia coli</i> isolates causing bloodstream infection. <i>Lancet Microbe</i> , The, 2021, 2, e492.	3.4	2
27	Viral detection and identification in 20Âmin by rapid single-particle fluorescence in-situ hybridization of viral RNA. <i>Scientific Reports</i> , 2021, 11, 19579.	1.6	16
28	Epidemiological data and genome sequencing reveals that nosocomial transmission of SARS-CoV-2 is underestimated and mostly mediated by a small number of highly infectious individuals. <i>Journal of Infection</i> , 2021, 83, 473-482.	1.7	55
29	The Duration, Dynamics, and Determinants of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Antibody Responses in Individual Healthcare Workers. <i>Clinical Infectious Diseases</i> , 2021, 73, e699-e709.	2.9	235
30	A genomic epidemiological study shows that prevalence of antimicrobial resistance in Enterobacterales is associated with the livestock host, as well as antimicrobial usage. <i>Microbial Genomics</i> , 2021, 7, .	1.0	20
31	Effect of Delta variant on viral burden and vaccine effectiveness against new SARS-CoV-2 infections in the UK. <i>Nature Medicine</i> , 2021, 27, 2127-2135.	15.2	450
32	Anti-spike antibody response to natural SARS-CoV-2 infection in the general population. <i>Nature Communications</i> , 2021, 12, 6250.	5.8	88
33	Frequencies and patterns of microbiology test requests from primary care in Oxfordshire, UK, 2008â€”2018: a retrospective cohort study of electronic health records to inform point-of-care testing. <i>BMJ Open</i> , 2021, 11, e048527.	0.8	2
34	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
35	Tracking the Emergence of SARS-CoV-2 Alpha Variant in the United Kingdom. <i>New England Journal of Medicine</i> , 2021, 385, 2582-2585.	13.9	49
36	Performance characteristics of five immunoassays for SARS-CoV-2: a head-to-head benchmark comparison. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 1390-1400.	4.6	336

#	ARTICLE	IF	CITATIONS
37	Risk Factors Associated with Carbapenemase-Producing <i>Enterobacterales</i> (CPE) Positivity in the Hospital Wastewater Environment. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	1.4	11
38	Pathogenicity of <i>mcr-1</i> -positive <i>Escherichia coli</i> from human infections. <i>Lancet Microbe</i> , The, 2020, 1, e195.	3.4	0
39	Genomic Epidemiology of Complex, Multispecies, Plasmid-Borne <i>bla</i> _{KPC} Carbapenemase in <i>Enterobacterales</i> in the United Kingdom from 2009 to 2014. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	1.4	26
40	Dynamics of <i>mcr-1</i> prevalence and <i>mcr-1</i> -positive <i>Escherichia coli</i> after the cessation of colistin use as a feed additive for animals in China: a prospective cross-sectional and whole genome sequencing-based molecular epidemiological study. <i>Lancet Microbe</i> , The, 2020, 1, e34-e43.	3.4	85
41	Genomic diversity affects the accuracy of bacterial single-nucleotide polymorphism calling pipelines. <i>GigaScience</i> , 2020, 9, .	3.3	92
42	Risk factors for <i>Klebsiella pneumoniae</i> carbapenemase (KPC) gene acquisition and clinical outcomes across multiple bacterial species. <i>Journal of Hospital Infection</i> , 2020, 104, 456-468.	1.4	24
43	Genomic dynamics of species and mobile genetic elements in a prolonged <i>bla</i> MP-4-associated carbapenemase outbreak in an Australian hospital. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 873-882.	1.3	31
44	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
45	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
46	Optimized use of Oxford Nanopore flowcells for hybrid assemblies. <i>Microbial Genomics</i> , 2020, 6, .	1.0	14
47	SARS-CoV-2 RNA detected in blood products from patients with COVID-19 is not associated with infectious virus. <i>Wellcome Open Research</i> , 2020, 5, 181.	0.9	81
48	SARS-CoV-2 RNA detected in blood products from patients with COVID-19 is not associated with infectious virus. <i>Wellcome Open Research</i> , 2020, 5, 181.	0.9	122
49	SARS-CoV-2 antibody prevalence, titres and neutralising activity in an antenatal cohort, United Kingdom, 14 April to 15 June 2020. <i>Eurosurveillance</i> , 2020, 25, .	3.9	17
50	Differential occupational risks to healthcare workers from SARS-CoV-2 observed during a prospective observational study. <i>ELife</i> , 2020, 9, .	2.8	196
51	The Role of <i>fosA</i> in Challenges with Fosfomycin Susceptibility Testing of Multispecies <i>Klebsiella pneumoniae</i> Carbapenemase-Producing Clinical Isolates. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	1.8	26
52	Molecular characterization of carbapenem-resistant <i>Escherichia coli</i> and <i>Acinetobacter baumannii</i> in the Lao People's Democratic Republic. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 2810-2821.	1.3	8
53	<i>Klebsiella quasipneumoniae</i> Provides a Window into Carbapenemase Gene Transfer, Plasmid Rearrangements, and Patient Interactions with the Hospital Environment. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	1.4	44
54	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

#	ARTICLE	IF	CITATIONS
55	Epidemiology of paediatric gastrointestinal colonisation by extended spectrum cephalosporin-resistant <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> isolates in north-west Cambodia. <i>BMC Microbiology</i> , 2019, 19, 59.	1.3	17
56	Don't overlook the little guy: An evaluation of the frequency of small plasmids co-conjugating with larger carbapenemase gene containing plasmids. <i>Plasmid</i> , 2019, 103, 1-8.	0.4	36
57	Occurrence and characterization of <i>Escherichia coli</i> ST410 co-harboring blaNDM-5, blaCMY-42 and blaTEM-190 in a dog from the UK. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 1207-1211.	1.3	31
58	The impact of sequencing depth on the inferred taxonomic composition and AMR gene content of metagenomic samples. <i>Environmental Microbiomes</i> , 2019, 14, 7.	2.2	69
59	Carbapenem-resistant Enterobacteriaceae dispersal from sinks is linked to drain position and drainage rates in a laboratory model system. <i>Journal of Hospital Infection</i> , 2019, 102, 63-69.	1.4	46
60	Comparison of long-read sequencing technologies in the hybrid assembly of complex bacterial genomes. <i>Microbial Genomics</i> , 2019, 5, .	1.0	171
61	Panton-Valentine leucocidin is the key determinant of <i>Staphylococcus aureus</i> pyomyositis in a bacterial GWAS. <i>ELife</i> , 2019, 8, .	2.8	56
62	Transmission dynamics and control of multidrug-resistant <i>Klebsiella pneumoniae</i> in neonates in a developing country. <i>ELife</i> , 2019, 8, .	2.8	17
63	Intensive Care Unit Wastewater Interventions to Prevent Transmission of Multispecies <i>Klebsiella pneumoniae</i> Carbapenemase-Producing Organisms. <i>Clinical Infectious Diseases</i> , 2018, 67, 171-178.	2.9	74
64	Illumina short-read and MinION long-read WGS to characterize the molecular epidemiology of an NDM-1 <i>Serratia marcescens</i> outbreak in Romania. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 672-679.	1.3	29
65	High Rates of Human Fecal Carriage of mcr-1-Positive Multidrug-Resistant Enterobacteriaceae Emerge in China in Association With Successful Plasmid Families. <i>Clinical Infectious Diseases</i> , 2018, 66, 676-685.	2.9	68
66	TETyper: a bioinformatic pipeline for classifying variation and genetic contexts of transposable elements from short-read whole-genome sequencing data. <i>Microbial Genomics</i> , 2018, 4, .	1.0	33
67	A Large, Refractory Nosocomial Outbreak of <i>Klebsiella pneumoniae</i> Carbapenemase-Producing <i>Escherichia coli</i> Demonstrates Carbapenemase Gene Outbreaks Involving Sink Sites Require Novel Approaches to Infection Control. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	1.4	99
68	Identification of Biomarkers for Differentiation of Hypervirulent <i>Klebsiella pneumoniae</i> from Classical <i>K. pneumoniae</i> . <i>Journal of Clinical Microbiology</i> , 2018, 56, .	1.8	378
69	Transmission Dynamics of Hyper-Endemic Multi-Drug Resistant <i>Klebsiella pneumoniae</i> in a Southeast Asian Neonatal Unit: A Longitudinal Study With Whole Genome Sequencing. <i>Frontiers in Microbiology</i> , 2018, 9, 1197.	1.5	24
70	Antimicrobial Resistance in Invasive Bacterial Infections in Hospitalized Children, Cambodia, 2007-2016. <i>Emerging Infectious Diseases</i> , 2018, 24, 841-851.	2.0	50
71	Bayesian reconstruction of transmission within outbreaks using genomic variants. <i>PLoS Computational Biology</i> , 2018, 14, e1006117.	1.5	69
72	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> , 2018, 18, 1138-1149.	4.6	121

#	ARTICLE	IF	CITATIONS
73	Carriage of β -lactamase-producing Enterobacteriaceae by Chinese travellers. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 138-139.	4.6	7
74	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
75	IncX2 and IncX1-X2 Hybrid Plasmids Coexisting in a FosA6-Producing <i>Escherichia coli</i> Strain. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	1.4	14
76	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
77	The Hospital Water Environment as a Reservoir for Carbapenem-Resistant Organisms Causing Hospital-Acquired Infections—A Systematic Review of the Literature. <i>Clinical Infectious Diseases</i> , 2017, 64, 1435-1444.	2.9	242
78	Enhanced <i>Klebsiella pneumoniae</i> Carbapenemase Expression from a Novel Tn <i>4401</i> Deletion. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	1.4	64
79	Ordering the mob: Insights into replicon and MOB typing schemes from analysis of a curated dataset of publicly available plasmids. <i>Plasmid</i> , 2017, 91, 42-52.	0.4	89
80	Chromosomal Integration of the <i>Klebsiella pneumoniae</i> Carbapenemase Gene, <i>bla</i> _{KPC} , in <i>Klebsiella</i> Species Is Elusive but Not Rare. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	1.4	46
81	A curated dataset of complete Enterobacteriaceae plasmids compiled from the NCBI nucleotide database. <i>Data in Brief</i> , 2017, 12, 423-426.	0.5	58
82	Genomic epidemiology of global <i>Klebsiella pneumoniae</i> carbapenemase (KPC)-producing <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2017, 7, 5917.	1.6	108
83	Covert dissemination of carbapenemase-producing <i>Klebsiella pneumoniae</i> (KPC) in a successfully controlled outbreak: long- and short-read whole-genome sequencing demonstrate multiple genetic modes of transmission. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 3025-3034.	1.3	73
84	Severity of illness and the weekend effect — Authors' reply. <i>Lancet</i> , The, 2017, 390, 1735.	6.3	11
85	Mortality Risks Associated With Emergency Admissions During Weekends and Public Holidays: An Analysis of Electronic Health Records. <i>Obstetrical and Gynecological Survey</i> , 2017, 72, 699-701.	0.2	0
86	Plasmid Classification in an Era of Whole-Genome Sequencing: Application in Studies of Antibiotic Resistance Epidemiology. <i>Frontiers in Microbiology</i> , 2017, 8, 182.	1.5	191
87	Resolving plasmid structures in Enterobacteriaceae using the MinION nanopore sequencer: assessment of MinION and MinION/Illumina hybrid data assembly approaches. <i>Microbial Genomics</i> , 2017, 3, e000118.	1.0	74
88	Supporting surveillance capacity for antimicrobial resistance: Laboratory capacity strengthening for drug resistant infections in low and middle income countries. <i>Wellcome Open Research</i> , 2017, 2, 91.	0.9	48
89	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
90	Complete Sequencing of Plasmids Containing <i>bla</i> _{OXA-163} and <i>bla</i> _{OXA-48} in <i>Escherichia coli</i> Sequence Type 131. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 6948-6951.	1.4	19

#	ARTICLE	IF	CITATIONS
91	Characterization of a Novel IncHI2 Plasmid Carrying Tandem Copies of <i>bla</i> _{CTX-M-2} in a <i>fosA6</i> -Harboring <i>Escherichia coli</i> Sequence Type 410 Strain. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 6742-6747.	1.4	12
92	First Report of <i>bla</i> _{IMP-14} on a Plasmid Harboring Multiple Drug Resistance Genes in <i>Escherichia coli</i> Sequence Type 131. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 5068-5071.	1.4	15
93	Oral fosfomycin for treatment of urinary tract infection: a retrospective cohort study. <i>BMC Infectious Diseases</i> , 2016, 16, 556.	1.3	53
94	Identifying lineage effects when controlling for population structure improves power in bacterial association studies. <i>Nature Microbiology</i> , 2016, 1, 16041.	5.9	247
95	Separate F-Type Plasmids Have Shaped the Evolution of the <i>H</i> 30 Subclone of <i>Escherichia coli</i> Sequence Type 131. <i>MSphere</i> , 2016, 1, .	1.3	98
96	Nested Russian Doll-Like Genetic Mobility Drives Rapid Dissemination of the Carbapenem Resistance Gene <i>bla</i> _{KPC} . <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 3767-3778.	1.4	255
97	Glutathione-S-transferase FosA6 of <i>Klebsiella pneumoniae</i> origin conferring fosfomycin resistance in ESBL-producing <i>Escherichia coli</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 2460-2465.	1.3	49
98	Antimicrobial susceptibility of uropathogens isolated from Cambodian children. <i>Paediatrics and International Child Health</i> , 2016, 36, 113-117.	0.3	24
99	Colistin resistance gene <i>mcr-1</i> and pHNSHP45 plasmid in human isolates of <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> . <i>Lancet Infectious Diseases</i> , The, 2016, 16, 285-286.	4.6	119
100	Evolutionary History of the Global Emergence of the <i>Escherichia coli</i> Epidemic Clone ST131. <i>MBio</i> , 2016, 7, e02162.	1.8	289
101	Complete Genome Sequence of KPC-Producing <i>Klebsiella pneumoniae</i> Strain CAV1193. <i>Genome Announcements</i> , 2016, 4, .	0.8	20
102	Characterisation of Invasive <i>Streptococcus pneumoniae</i> Isolated from Cambodian Children between 2007 – 2012. <i>PLoS ONE</i> , 2016, 11, e0159358.	1.1	15
103	Dynamics of MDR <i>Enterobacter cloacae</i> outbreaks in a neonatal unit in Nepal: insights using wider sampling frames and next-generation sequencing. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 1008-1015.	1.3	45
104	<i>Klebsiella pneumoniae</i> Carbapenemase (KPC)-Producing <i>K. pneumoniae</i> at a Single Institution: Insights into Endemicity from Whole-Genome Sequencing. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 1656-1663.	1.4	140
105	Colonization with Enterobacteriaceae producing ESBLs in children attending pre-school childcare facilities in the Lao People's Democratic Republic. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 1893-1897.	1.3	62
106	Extensive Within-Host Diversity in Fecally Carried Extended-Spectrum-Beta-Lactamase-Producing <i>Escherichia coli</i> Isolates: Implications for Transmission Analyses. <i>Journal of Clinical Microbiology</i> , 2015, 53, 2122-2131.	1.8	84
107	Rapid Diagnostic Tests for Dengue Virus Infection in Febrile Cambodian Children: Diagnostic Accuracy and Incorporation into Diagnostic Algorithms. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003424.	1.3	24
108	Multidrug-resistant <i>Escherichia coli</i> soft tissue infection investigated with bacterial whole genome sequencing. <i>BMJ Case Reports</i> , 2014, 2014, bcr2014207200-bcr2014207200.	0.2	4

#	ARTICLE	IF	CITATIONS
109	Ophthalmic infections in children presenting to Angkor Hospital for Children, Siem Reap, Cambodia. <i>BMC Research Notes</i> , 2014, 7, 784.	0.6	9
110	Genome Sequencing of an Extended Series of NDM-Producing <i>Klebsiella pneumoniae</i> Isolates from Neonatal Infections in a Nepali Hospital Characterizes the Extent of Community- versus Hospital-Associated Transmission in an Endemic Setting. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 7347-7357.	1.4	142
111	Septic arthritis of the hip in a Cambodian child caused by multidrug-resistant <i>Salmonella enterica</i> serovar Typhi with intermediate susceptibility to ciprofloxacin treated with ceftriaxone and azithromycin. <i>Paediatrics and International Child Health</i> , 2014, 34, 227-229.	0.3	4
112	Evolutionary History of the <i>Clostridium difficile</i> Pathogenicity Locus. <i>Genome Biology and Evolution</i> , 2014, 6, 36-52.	1.1	190
113	Treatment of Suspected Hyper-Reactive Malarial Splenomegaly (HMS) in Pregnancy with Mefloquine. <i>American Journal of Tropical Medicine and Hygiene</i> , 2014, 90, 609-611.	0.6	6
114	Predicting antimicrobial susceptibilities for <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> isolates using whole genomic sequence data. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 2234-2244.	1.3	314
115	Comparison of Multilocus Variable-Number Tandem-Repeat Analysis and Whole-Genome Sequencing for Investigation of <i>Clostridium difficile</i> Transmission. <i>Journal of Clinical Microbiology</i> , 2013, 51, 4141-4149.	1.8	69
116	The Epidemiology of Pediatric Bone and Joint Infections in Cambodia, 2007-11. <i>Journal of Tropical Pediatrics</i> , 2013, 59, 36-42.	0.7	22
117	Risk Factors for <i>Clostridium difficile</i> Acquisition in Infants: Importance of Study Design. <i>Clinical Infectious Diseases</i> , 2013, 56, 1680-1681.	2.9	5
118	The value of intermittent point-prevalence surveys of healthcare-associated infections for evaluating infection control interventions at Angkor Hospital for Children, Siem Reap, Cambodia. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2013, 107, 248-253.	0.7	29
119	Pediatric Bloodstream Infections in Cambodia, 2007 to 2011. <i>Pediatric Infectious Disease Journal</i> , 2013, 32, e272-e276.	1.1	34
120	A Prospective Study of the Causes of Febrile Illness Requiring Hospitalization in Children in Cambodia. <i>PLoS ONE</i> , 2013, 8, e60634.	1.1	88
121	Increasing incidence of <i>Escherichia coli</i> bacteraemia is driven by an increase in antibiotic-resistant isolates: electronic database study in Oxfordshire 1999-2011. <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 1514-1524.	1.3	45
122	Fidaxomicin Versus Vancomycin for <i>Clostridium difficile</i> Infection: Meta-analysis of Pivotal Randomized Controlled Trials. <i>Clinical Infectious Diseases</i> , 2012, 55, S93-S103.	2.9	228
123	Changing Patterns of Gastrointestinal Parasite Infections in Cambodian Children: 2006-2011. <i>Journal of Tropical Pediatrics</i> , 2012, 58, 509-512.	0.7	14
124	Characteristics of CTX-M ESBL-producing <i>Escherichia coli</i> isolates from the Lao People's Democratic Republic, 2004-09. <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 240-242.	1.3	25
125	Paediatric <i>Chromobacterium violaceum</i> in Cambodia: the first documented case. <i>Tropical Doctor</i> , 2012, 42, 178-179.	0.2	8
126	Pediatric Suppurative Parotitis in Cambodia Between 2007 and 2011. <i>Pediatric Infectious Disease Journal</i> , 2012, 31, 865-868.	1.1	32

#	ARTICLE	IF	CITATIONS
127	Surveillance of healthcare-associated infection at Angkor Hospital for Children, Siem Reap, Cambodia. <i>International Journal of Infectious Diseases</i> , 2012, 16, e375.	1.5	0
128	Enteric fever in Cambodian children is dominated by multidrug resistant H58 <i>Salmonella enterica</i> serovar Typhi with decreased susceptibility to ciprofloxacin. <i>International Journal of Infectious Diseases</i> , 2012, 16, e427.	1.5	0
129	Enteric fever in Cambodian children is dominated by multidrug-resistant H58 <i>Salmonella enterica</i> serovar Typhi with intermediate susceptibility to ciprofloxacin. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2012, 106, 718-724.	0.7	38
130	Clinical <i>Clostridium difficile</i> : Clonality and Pathogenicity Locus Diversity. <i>PLoS ONE</i> , 2011, 6, e19993.	1.1	150
131	Molecular Epidemiology of <i>Clostridium difficile</i> Strains in Children Compared with That of Strains Circulating in Adults with <i>Clostridium difficile</i> -Associated Infection : Fig. 1.. <i>Journal of Clinical Microbiology</i> , 2011, 49, 3994-3996.	1.8	32
132	Multilocus Sequence Typing of <i>Clostridium difficile</i> . <i>Journal of Clinical Microbiology</i> , 2010, 48, 770-778.	1.8	399
133	Re-visiting <i>Clostridium difficile</i> in children: Reservoir, victims, both or none?. <i>Journal of Infection</i> , 2009, 59, S429-S430.	1.7	0
134	Delayed parasite elimination in human infections treated with clindamycin parallels "delayed death" of <i>Plasmodium falciparum</i> in vitro. <i>International Journal for Parasitology</i> , 2007, 37, 777-785.	1.3	27
135	Treatment of enteric fever (typhoid and paratyphoid fever) with third and fourth generation cephalosporins. <i>The Cochrane Library</i> , 0, , .	1.5	3
136	Antimicrobial susceptibility of uropathogens isolated from Cambodian children. <i>Paediatrics and International Child Health</i> , 0, , 1-5.	0.3	1
137	Risk Factors Associated with Carbapenemase-Producing Enterobacterales (CPE) Positivity in the Hospital Wastewater Environment. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0
138	Mortality risks associated with empirical antibiotic activity in <i>Escherichia coli</i> bacteraemia: an analysis of electronic health records. <i>Journal of Antimicrobial Chemotherapy</i> , 0, , .	1.3	3