

# Paul Turner

## List of Publications by Year in descending order

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145  
papers

14,070  
citations

81900

39  
h-index

26613

107  
g-index

176  
all docs

176  
docs citations

176  
times ranked

14288  
citing authors

#	ARTICLE	IF	CITATIONS
1	Global burden of bacterial antimicrobial resistance in 2019: a systematic analysis. <i>Lancet</i> , The, 2022, 399, 629-655.	13.7	4,915
2	Reagent and laboratory contamination can critically impact sequence-based microbiome analyses. <i>BMC Biology</i> , 2014, 12, 87.	3.8	2,677
3	Phylogeographical analysis of the dominant multidrug-resistant H58 clade of <i>Salmonella</i> Typhi identifies inter- and intracontinental transmission events. <i>Nature Genetics</i> , 2015, 47, 632-639.	21.4	403
4	Standard method for detecting upper respiratory carriage of <i>Streptococcus pneumoniae</i> : Updated recommendations from the World Health Organization Pneumococcal Carriage Working Group. <i>Vaccine</i> , 2013, 32, 165-179.	3.8	374
5	Dense genomic sampling identifies highways of pneumococcal recombination. <i>Nature Genetics</i> , 2014, 46, 305-309.	21.4	371
6	Evolutionary History of the Global Emergence of the <i>Escherichia coli</i> Epidemic Clone ST131. <i>MBio</i> , 2016, 7, e02162.	4.1	289
7	Global patterns in monthly activity of influenza virus, respiratory syncytial virus, parainfluenza virus, and metapneumovirus: a systematic analysis. <i>The Lancet Global Health</i> , 2019, 7, e1031-e1045.	6.3	266
8	A New Pneumococcal Capsule Type, 10D, is the 100th Serotype and Has a Large <i>cps</i> Fragment from an Oral <i>Streptococcus</i> . <i>MBio</i> , 2020, 11, .	4.1	219
9	Comprehensive Identification of Single Nucleotide Polymorphisms Associated with Beta-lactam Resistance within Pneumococcal Mosaic Genes. <i>PLoS Genetics</i> , 2014, 10, e1004547.	3.5	205
10	Genomic surveillance for hypervirulence and multi-drug resistance in invasive <i>Klebsiella pneumoniae</i> from South and Southeast Asia. <i>Genome Medicine</i> , 2020, 12, 11.	8.2	178
11	Pneumococcal lineages associated with serotype replacement and antibiotic resistance in childhood invasive pneumococcal disease in the post-PCV13 era: an international whole-genome sequencing study. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 759-769.	9.1	165
12	An extended genotyping framework for <i>Salmonella enterica</i> serovar Typhi, the cause of human typhoid. <i>Nature Communications</i> , 2016, 7, 12827.	12.8	145
13	A current perspective on antimicrobial resistance in Southeast Asia. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 2963-2972.	3.0	139
14	Improved Detection of Nasopharyngeal Cocolonization by Multiple Pneumococcal Serotypes by Use of Latex Agglutination or Molecular Serotyping by Microarray. <i>Journal of Clinical Microbiology</i> , 2011, 49, 1784-1789.	3.9	134
15	Evolution of antibiotic resistance is linked to any genetic mechanism affecting bacterial duration of carriage. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 1075-1080.	7.1	133
16	A Longitudinal Study of <i>Streptococcus pneumoniae</i> Carriage in a Cohort of Infants and Their Mothers on the Thailand-Myanmar Border. <i>PLoS ONE</i> , 2012, 7, e38271.	2.5	129
17	Horizontal DNA Transfer Mechanisms of Bacteria as Weapons of Intragenomic Conflict. <i>PLoS Biology</i> , 2016, 14, e1002394.	5.6	127
18	Evidence for Soft Selective Sweeps in the Evolution of Pneumococcal Multidrug Resistance and Vaccine Escape. <i>Genome Biology and Evolution</i> , 2014, 6, 1589-1602.	2.5	112

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19	Antimicrobial Resistance Surveillance in Low- and Middle-Income Countries: Progress and Challenges in Eight South Asian and Southeast Asian Countries. <i>Clinical Microbiology Reviews</i> , 2020, 33, .	13.6	105
20	Performance of C-reactive protein and procalcitonin to distinguish viral from bacterial and malarial causes of fever in Southeast Asia. <i>BMC Infectious Diseases</i> , 2015, 15, 511.	2.9	103
21	Genome-wide identification of lineage and locus specific variation associated with pneumococcal carriage duration. <i>ELife</i> , 2017, 6, .	6.0	95
22	Interacting networks of resistance, virulence and core machinery genes identified by genome-wide epistasis analysis. <i>PLoS Genetics</i> , 2017, 13, e1006508.	3.5	85
23	South Asia as a Reservoir for the Global Spread of Ciprofloxacin-Resistant <i>Shigella sonnei</i> : A Cross-Sectional Study. <i>PLoS Medicine</i> , 2016, 13, e1002055.	8.4	84
24	Antimicrobial susceptibility of bacterial isolates from community acquired infections in Sub-Saharan Africa and Asian low and middle income countries. <i>Tropical Medicine and International Health</i> , 2011, 16, 1167-1179.	2.3	67
25	Pneumococcal Capsule Synthesis Locus <i>cps</i> as Evolutionary Hotspot with Potential to Generate Novel Serotypes by Recombination. <i>Molecular Biology and Evolution</i> , 2017, 34, 2537-2554.	8.9	65
26	Predicting the severity of dengue fever in children on admission based on clinical features and laboratory indicators: application of classification tree analysis. <i>BMC Pediatrics</i> , 2018, 18, 109.	1.7	65
27	Global Phylogenomic Analysis of Nonencapsulated <i>Streptococcus pneumoniae</i> Reveals a Deep-Branching Classic Lineage That Is Distinct from Multiple Sporadic Lineages. <i>Genome Biology and Evolution</i> , 2014, 6, 3281-3294.	2.5	63
28	A longitudinal study of the infant nasopharyngeal microbiota: The effects of age, illness and antibiotic use in a cohort of South East Asian children. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005975.	3.0	62
29	Towards understanding global patterns of antimicrobial use and resistance in neonatal sepsis: insights from the NeoAMR network. <i>Archives of Disease in Childhood</i> , 2020, 105, 26-31.	1.9	56
30	Panton-Valentine leucocidin is the key determinant of <i>Staphylococcus aureus</i> pyomyositis in a bacterial GWAS. <i>ELife</i> , 2019, 8, .	6.0	56
31	Serum antibody responses to pneumococcal colonization in the first 2 years of life: results from an SE Asian longitudinal cohort study. <i>Clinical Microbiology and Infection</i> , 2013, 19, E551-E558.	6.0	55
32	Microbiology Investigation Criteria for Reporting Objectively (MICRO): a framework for the reporting and interpretation of clinical microbiology data. <i>BMC Medicine</i> , 2019, 17, 70.	5.5	55
33	Dermal absorption of isopropyl alcohol from a commercial hand rub: implications for its use in hand decontamination. <i>Journal of Hospital Infection</i> , 2004, 56, 287-290.	2.9	52
34	One hypervirulent clone, sequence type 283, accounts for a large proportion of invasive <i>Streptococcus agalactiae</i> isolated from humans and diseased tilapia in Southeast Asia. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007421.	3.0	51
35	Antimicrobial Resistance in Invasive Bacterial Infections in Hospitalized Children, Cambodia, 2007-2016. <i>Emerging Infectious Diseases</i> , 2018, 24, 841-851.	4.3	50
36	Using machine learning to guide targeted and locally-tailored empiric antibiotic prescribing in a children's hospital in Cambodia. <i>Wellcome Open Research</i> , 2018, 3, 131.	1.8	48

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37	Impact of CLSI and EUCAST breakpoint discrepancies on reporting of antimicrobial susceptibility and AMR surveillance. <i>Clinical Microbiology and Infection</i> , 2019, 25, 910-911.	6.0	48
38	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.	1.8	48
39	High Prevalence of Antimicrobial-resistant Gram-negative Colonization in Hospitalized Cambodian Infants. <i>Pediatric Infectious Disease Journal</i> , 2016, 35, 856-861.	2.0	47
40	An inventory of supranational antimicrobial resistance surveillance networks involving low- and middle-income countries since 2000. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 1737-1749.	3.0	47
41	Molecular Surveillance Identifies Multiple Transmissions of Typhoid in West Africa. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004781.	3.0	46
42	Clinically and Microbiologically Derived Azithromycin Susceptibility Breakpoints for <i>Salmonella enterica</i> Serovars Typhi and Paratyphi A. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 2756-2764.	3.2	44
43	Neonatal Intensive Care in a Karen Refugee Camp: A 4 Year Descriptive Study. <i>PLoS ONE</i> , 2013, 8, e72721.	2.5	43
44	Climate induces seasonality in pneumococcal transmission. <i>Scientific Reports</i> , 2015, 5, 11344.	3.3	41
45	Dissecting the molecular evolution of fluoroquinolone-resistant <i>Shigella sonnei</i> . <i>Nature Communications</i> , 2019, 10, 4828.	12.8	41
46	A Prospective Evaluation of Real-Time PCR Assays for the Detection of <i>Orientia tsutsugamushi</i> and <i>Rickettsia</i> spp. for Early Diagnosis of Rickettsial Infections during the Acute Phase of Undifferentiated Febrile Illness. <i>American Journal of Tropical Medicine and Hygiene</i> , 2013, 89, 308-310.	1.4	40
47	The Molecular and Spatial Epidemiology of Typhoid Fever in Rural Cambodia. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004785.	3.0	40
48	Susceptibility of community-acquired pathogens to antibiotics in Africa and Asia in neonates – an alarmingly short review. <i>Tropical Medicine and International Health</i> , 2011, 16, 145-151.	2.3	36
49	Group B streptococcal carriage, serotype distribution and antibiotic susceptibilities in pregnant women at the time of delivery in a refugee population on the Thai-Myanmar border. <i>BMC Infectious Diseases</i> , 2012, 12, 34.	2.9	36
50	High Rates of Pneumonia in Children under Two Years of Age in a South East Asian Refugee Population. <i>PLoS ONE</i> , 2013, 8, e54026.	2.5	36
51	Matrix-assisted laser desorption/ionization time-of-flight mass spectrometry for the identification of <i>Burkholderia pseudomallei</i> from Asia and Australia and differentiation between <i>Burkholderia</i> species. <i>PLoS ONE</i> , 2017, 12, e0175294.	2.5	36
52	Duplex Real-Time PCR Assay for Detection of <i>Streptococcus pneumoniae</i> in Clinical Samples and Determination of Penicillin Susceptibility. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2751-2758.	3.9	35
53	Capsular Typing Method for <i>Streptococcus agalactiae</i> Using Whole-Genome Sequence Data. <i>Journal of Clinical Microbiology</i> , 2016, 54, 1388-1390.	3.9	35
54	Leapfrogging laboratories: the promise and pitfalls of high-tech solutions for antimicrobial resistance surveillance in low-income settings. <i>BMJ Global Health</i> , 2020, 5, e003622.	4.7	30

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55	A retrospective analysis of melioidosis in Cambodian children, 2009–2013. BMC Infectious Diseases, 2016, 16, 688.	2.9	29
56	Respiratory virus surveillance in hospitalised pneumonia patients on the Thailand-Myanmar border. BMC Infectious Diseases, 2013, 13, 434.	2.9	27
57	Antimicrobial resistance in Cambodia: a review. International Journal of Infectious Diseases, 2019, 85, 98-107.	3.3	27
58	The aetiologies of central nervous system infections in hospitalised Cambodian children. BMC Infectious Diseases, 2017, 17, 806.	2.9	26
59	A prospective evaluation of diagnostic methodologies for the acute diagnosis of dengue virus infection on the Thailand-Myanmar border. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2011, 105, 32-37.	1.8	25
60	Trimethoprim/sulfamethoxazole resistance in Burkholderia pseudomallei. International Journal of Antimicrobial Agents, 2014, 44, 368-369.	2.5	24
61	Transmission Dynamics of Hyper-Endemic Multi-Drug Resistant Klebsiella pneumoniae in a Southeast Asian Neonatal Unit: A Longitudinal Study With Whole Genome Sequencing. Frontiers in Microbiology, 2018, 9, 1197.	3.5	24
62	Climatic drivers of melioidosis in Laos and Cambodia: a 16-year case series analysis. Lancet Planetary Health, The, 2018, 2, e334-e343.	11.4	23
63	Two-phase importance sampling for inference about transmission trees. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20141324.	2.6	22
64	Understanding pneumococcal serotype 1 biology through population genomic analysis. BMC Infectious Diseases, 2016, 16, 649.	2.9	22
65	Assessment of Streptococcus pneumoniae pilus islet-1 prevalence in carried and transmitted isolates from mother–infant pairs on the Thailand–Burma border. Clinical Microbiology and Infection, 2012, 18, 970-975.	6.0	21
66	Field Evaluation of Culture plus Latex Sweep Serotyping for Detection of Multiple Pneumococcal Serotype Colonisation in Infants and Young Children. PLoS ONE, 2013, 8, e67933.	2.5	21
67	Impact of 13-Valent Pneumococcal Conjugate Vaccine on Colonization and Invasive Disease in Cambodian Children. Clinical Infectious Diseases, 2020, 70, 1580-1588.	5.8	21
68	The role of interspecies recombination in the evolution of antibiotic-resistant pneumococci. ELife, 2021, 10, .	6.0	21
69	Surveillance strategies using routine microbiology for antimicrobial resistance in low- and middle-income countries. Clinical Microbiology and Infection, 2021, 27, 1391-1399.	6.0	20
70	Pneumococcal Infection among Children before Introduction of 13-Valent Pneumococcal Conjugate Vaccine, Cambodia. Emerging Infectious Diseases, 2015, 21, 2080-2083.	4.3	19
71	Horizontal gene transfer rate is not the primary determinant of observed antibiotic resistance frequencies in <i>Streptococcus pneumoniae</i> . Science Advances, 2020, 6, eaaz6137.	10.3	19
72	Time to switch from CLSI to EUCAST? A Southeast Asian perspective. Clinical Microbiology and Infection, 2019, 25, 782-785.	6.0	18

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73	ACORN (A Clinically-Oriented Antimicrobial Resistance Surveillance Network): a pilot protocol for case based antimicrobial resistance surveillance. Wellcome Open Research, 2020, 5, 13.	1.8	18
74	<i>Elizabethkingia anophelis</i> Infection in Infants, Cambodia, 2012–2018. Emerging Infectious Diseases, 2020, 26, 320-322.	4.3	17
75	Evolutionary histories and antimicrobial resistance in <i>Shigella flexneri</i> and <i>Shigella sonnei</i> in Southeast Asia. Communications Biology, 2021, 4, 353.	4.4	17
76	Transmission dynamics and control of multidrug-resistant <i>Klebsiella pneumoniae</i> in neonates in a developing country. ELife, 2019, 8, .	6.0	17
77	Development of a TaqMan Array Card for Pneumococcal Serotyping on Isolates and Nasopharyngeal Samples. Journal of Clinical Microbiology, 2016, 54, 1842-1850.	3.9	16
78	'Candidatus <i>Ornithobacterium hominis</i> ': insights gained from draft genomes obtained from nasopharyngeal swabs. Microbial Genomics, 2019, 5, .	2.0	16
79	Susceptibility of bacterial isolates from community-acquired infections in sub-Saharan Africa and Asia to macrolide antibiotics. Tropical Medicine and International Health, 2011, 16, 1192-1205.	2.3	15
80	Characterisation of Invasive <i>Streptococcus pneumoniae</i> Isolated from Cambodian Children between 2007 – 2012. PLoS ONE, 2016, 11, e0159358.	2.5	15
81	Utilization of a clinical microbiology service at a Cambodian paediatric hospital and its impact on appropriate antimicrobial prescribing. Journal of Antimicrobial Chemotherapy, 2018, 73, 509-516.	3.0	14
82	Automating the Generation of Antimicrobial Resistance Surveillance Reports: Proof-of-Concept Study Involving Seven Hospitals in Seven Countries. Journal of Medical Internet Research, 2020, 22, e19762.	4.3	14
83	A prospective study of urinary pneumococcal antigen detection in healthy Karen mothers with high rates of pneumococcal nasopharyngeal carriage. BMC Infectious Diseases, 2011, 11, 108.	2.9	13
84	A High Burden of Respiratory Syncytial Virus Associated Pneumonia in Children Less than Two Years of Age in a South East Asian Refugee Population. PLoS ONE, 2012, 7, e50100.	2.5	13
85	Impact of delays to incubation and storage temperature on blood culture results: a multi-centre study. BMC Infectious Diseases, 2021, 21, 173.	2.9	13
86	Laboratory informatics capacity for effective antimicrobial resistance surveillance in resource-limited settings. Lancet Infectious Diseases, The, 2021, 21, e170-e174.	9.1	13
87	Predictors of disease severity in children presenting from the community with febrile illnesses: a systematic review of prognostic studies. BMJ Global Health, 2021, 6, e003451.	4.7	13
88	ACORN (A Clinically-Oriented Antimicrobial Resistance Surveillance Network): a pilot protocol for case based antimicrobial resistance surveillance. Wellcome Open Research, 2020, 5, 13.	1.8	13
89	Detection of Respiratory Viruses by PCR Assay of Nasopharyngeal Swabs Stored in Skim Milk-Tryptone-Glucose-Glycerol Transport Medium. Journal of Clinical Microbiology, 2011, 49, 2311-2313.	3.9	12
90	A three year descriptive study of early onset neonatal sepsis in a refugee population on the Thailand Myanmar border. BMC Infectious Diseases, 2013, 13, 601.	2.9	12

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91	Diagnostic Accuracy Assessment of Immunochromatographic Tests for the Rapid Detection of Antibodies Against <i>Orientia tsutsugamushi</i> Using Paired Acute and Convalescent Specimens. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 93, 1168-1171.	1.4	12
92	Prospective surveillance of healthcare associated infections in a Cambodian pediatric hospital. <i>Antimicrobial Resistance and Infection Control</i> , 2017, 6, 16.	4.1	12
93	Grading antimicrobial susceptibility data quality: room for improvement. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 603-604.	9.1	12
94	A mosaic tetracycline resistance gene <i>tet(S/M)</i> detected in an MDR pneumococcal CC230 lineage that underwent capsular switching in South Africa. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 512-520.	3.0	12
95	Impact of low blood culture usage on rates of antimicrobial resistance. <i>Journal of Infection</i> , 2021, 82, 355-362.	3.3	12
96	Prediction of disease severity in young children presenting with acute febrile illness in resource-limited settings: a protocol for a prospective observational study. <i>BMJ Open</i> , 2021, 11, e045826.	1.9	12
97	Evaluation of the Coverage of 3 Antibiotic Regimens for Neonatal Sepsis in the Hospital Setting Across Asian Countries. <i>JAMA Network Open</i> , 2020, 3, e1921124.	5.9	11
98	Introduction and establishment of fluoroquinolone-resistant <i>Shigella sonnei</i> into Bhutan. <i>Microbial Genomics</i> , 2015, 1, e000042.	2.0	11
99	Antimicrobial use and resistance data in human and animal sectors in the Lao PDR: evidence to inform policy. <i>BMJ Global Health</i> , 2021, 6, e007009.	4.7	11
100	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.	7.1	10
101	Ophthalmic infections in children presenting to Angkor Hospital for Children, Siem Reap, Cambodia. <i>BMC Research Notes</i> , 2014, 7, 784.	1.4	9
102	MALDI-TOF mass spectrometry for sub-typing of <i>Streptococcus pneumoniae</i> . <i>BMC Microbiology</i> , 2020, 20, 367.	3.3	9
103	Influenza in Refugees on the Thailand–Myanmar Border, May–October 2009. <i>Emerging Infectious Diseases</i> , 2010, 16, 1366-1372.	4.3	8
104	Rectal pH in Well and Unwell Infants. <i>Journal of Tropical Pediatrics</i> , 2012, 58, 311-313.	1.5	8
105	Standardising the reporting of microbiology and antimicrobial susceptibility data. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 1163-1164.	9.1	8
106	Antimicrobial-resistant Gram-negative colonization in infants from a neonatal intensive care unit in Thailand. <i>Journal of Hospital Infection</i> , 2019, 103, 151-155.	2.9	8
107	Improving Treatment and Outcomes for Melioidosis in Children, Northern Cambodia, 2009–2018. <i>Emerging Infectious Diseases</i> , 2021, 27, 1169-1172.	4.3	8
108	Causes of fever in primary care in Southeast Asia and the performance of C-reactive protein in discriminating bacterial from viral pathogens. <i>International Journal of Infectious Diseases</i> , 2020, 96, 334-342.	3.3	8



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109	Seroprevalence of Dengue Virus and Rickettsial Infections in Cambodian Children. American Journal of Tropical Medicine and Hygiene, 2019, 100, 635-638.	1.4	8
110	A Novel Technique for Detecting Antibiotic-Resistant Typhoid from Rapid Diagnostic Tests. Journal of Clinical Microbiology, 2015, 53, 1758-1760.	3.9	7
111	Salmonella Typhi and Paratyphi A infections in Cambodian children, 2012â€“2016. International Journal of Infectious Diseases, 2020, 97, 334-336.	3.3	7
112	Boosting heritability: estimating the genetic component of phenotypic variation with multiple sample splitting. BMC Bioinformatics, 2021, 22, 164.	2.6	7
113	A multi-country study using MALDI-TOF mass spectrometry for rapid identification of Burkholderia pseudomallei. BMC Microbiology, 2021, 21, 213.	3.3	7
114	Two fatal cases of melioidosis on the Thai-Myanmar border. F1000Research, 2014, 3, 4.	1.6	7
115	Serotype Distribution of Clinical Streptococcus pneumoniae Isolates before the Introduction of the 13-Valent Pneumococcal Conjugate Vaccine in Cambodia. American Journal of Tropical Medicine and Hygiene, 2018, 98, 791-796.	1.4	7
116	Epidemiology of Extended-Spectrum Beta-Lactamase and Carbapenemase-Producing Enterobacterales in the Greater Mekong Subregion: A Systematic-Review and Meta-Analysis of Risk Factors Associated With Extended-Spectrum Beta-Lactamase and Carbapenemase Isolation. Frontiers in Microbiology, 2021, 12, 695027.	3.5	7
117	Use of Blood Smears and Dried Blood Spots for Polymerase Chain Reactionâ€“Based Detection and Quantification of Bacterial Infection and Plasmodium falciparum in Severely Ill Febrile African Children. American Journal of Tropical Medicine and Hygiene, 2016, 94, 322-326.	1.4	6
118	The cost-effectiveness of the use of selective media for the diagnosis of melioidosis in different settings. PLoS Neglected Tropical Diseases, 2019, 13, e0007598.	3.0	6
119	The spread of chloramphenicol-resistant Neisseria meningitidis in Southeast Asia. International Journal of Infectious Diseases, 2020, 95, 198-203.	3.3	6
120	Antimicrobial resistance detection in Southeast Asian hospitals is critically important from both patient and societal perspectives, but what is its cost?. PLOS Global Public Health, 2021, 1, e0000018.	1.6	6
121	Antimicrobial resistance patterns in bacteria causing febrile illness in Africa, South Asia, and Southeast Asia: a systematic review of published etiological studies from 1980-2015. International Journal of Infectious Diseases, 2022, 122, 612-621.	3.3	6
122	Aeromonas spp. Bacteremia in Pregnant Women, Thailandâ€“Myanmar Border, 2011. Emerging Infectious Diseases, 2012, 18, 1522-3.	4.3	5
123	Prevalence of MDR organism (MDRO) carriage in children and their household members in Siem Reap Province, Cambodia. JAC-Antimicrobial Resistance, 2020, 2, dlaa097.	2.1	5
124	Genome-wide association, prediction and heritability in bacteria with application to Streptococcus pneumoniae. NAR Genomics and Bioinformatics, 2022, 4, lqac011.	3.2	5
125	Impact of delayed processing of positive blood cultures on organism detection: a prospective multi-centre study. BMC Infectious Diseases, 2022, 22, .	2.9	5
126	Antibody-Mediated Complement C3b/iC3b Binding to Group B Streptococcus in Paired Mother and Baby Serum Samples in a Refugee Population on the Thailand-Myanmar Border. Vaccine Journal, 2015, 22, 319-326.	3.1	4



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127	Case-based surveillance of antimicrobial resistance in the ACORN (A Clinically Oriented Antimicrobial) Tj ETQq1 1 0,784314 rgBT /Overd	2.1	4
128	Setting priorities for patient-centered surveillance of drug-resistant infections. International Journal of Infectious Diseases, 2020, 97, 60-65.	3.3	4
129	Two fatal cases of melioidosis on the Thai-Myanmar border. F1000Research, 2014, 3, 4.	1.6	4
130	Facilitating Safe Discharge Through Predicting Disease Progression in Moderate Coronavirus Disease 2019 (COVID-19): A Prospective Cohort Study to Develop and Validate a Clinical Prediction Model in Resource-Limited Settings. Clinical Infectious Diseases, 2022, 75, e368-e379.	5.8	4
131	World Health Organization (WHO) standard methods for pneumococcal carriage studies. Clinical Infectious Diseases, 2022, , .	5.8	4
132	A Streptococcus pneumoniae lineage usually associated with pneumococcal conjugate vaccine (PCV) serotypes is the most common cause of serotype 35B invasive disease in South Africa, following routine use of PCV. Microbial Genomics, 2022, 8, .	2.0	4
133	Evaluation of trends in hospital antimicrobial use in the Lao PDR using repeated point-prevalence surveys-evidence to improve treatment guideline use. The Lancet Regional Health - Western Pacific, 2022, 27, 100531.	2.9	4
134	Molecular Epidemiology of Group A Streptococcus Infections in Cambodian Children, 2007â€“2012. Pediatric Infectious Disease Journal, 2015, 34, 1414-1415.	2.0	3
135	Necrotizing fasciitis complicating snakebite in Cambodia. IDCases, 2015, 2, 86-87.	0.9	3
136	The evolutionary history of Shigella flexneri serotype 6 in Asia. Microbial Genomics, 2021, 7, .	2.0	3
137	A case of human syngamosis. Travel Medicine and Infectious Disease, 2003, 1, 231-233.	3.0	2
138	Get the Basics Right: A Description of the Key Priorities for Establishing a Neonatal Service in a Resource-Limited Setting in Cambodia. Journal of Tropical Pediatrics, 2019, 65, 160-168.	1.5	2
139	Antimicrobial resistance in commensal opportunistic pathogens isolated from non-sterile sites can be an effective proxy for surveillance in bloodstream infections. Scientific Reports, 2021, 11, 23359.	3.3	2
140	Practical Methods to Permit the Analysis of Host Biomarkers in Resource-Limited Settings. American Journal of Tropical Medicine and Hygiene, 2022, 106, 1765-1769.	1.4	2
141	Detection of colonisation by extended-spectrum beta-lactamase or carbapenemase producing Enterobacterales from frozen stool specimens. BMC Research Notes, 2020, 13, 429.	1.4	1
142	Rectal pH in Well and Unwell Infants. Journal of Tropical Pediatrics, 2013, 59, 162-162.	1.5	0
143	Lack of Utility of Nasopharyngeal Swabs for Diagnosis ofBurkholderia pseudomalleiPneumonia in Paediatric Patients. Journal of Tropical Pediatrics, 2016, 62, 328-330.	1.5	0
144	Evolutionary histories and antimicrobial resistance inShigella flexneri and Shigella sonnei in Southeast Asia. Access Microbiology, 2022, 4, .	0.5	0

#	ARTICLE	IF	CITATIONS
145	Genetic background of Cambodian pneumococcal carriage isolates following pneumococcal conjugate vaccine 13. Microbial Genomics, 2022, 8, .	2.0	0