

# Paul Turner

## List of Publications by Year in descending order

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145  
papers

14,070  
citations

81900  
39  
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26613  
107  
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176  
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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	Genome-wide association, prediction and heritability in bacteria with application to <i>Streptococcus pneumoniae</i> . <i>NAR Genomics and Bioinformatics</i> , 2022, 4, lqac011.	3.2	5
3	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. <i>Clinical Infectious Diseases</i> , 2022, 75, e368-e379.	5.8	4
4	World Health Organization (WHO) standard methods for pneumococcal carriage studies. <i>Clinical Infectious Diseases</i> , 2022, , .	5.8	4
5	A <i>Streptococcus pneumoniae</i> 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. <i>Microbial Genomics</i> , 2022, 8, .	2.0	4
6	Practical Methods to Permit the Analysis of Host Biomarkers in Resource-Limited Settings. <i>American Journal of Tropical Medicine and Hygiene</i> , 2022, 106, 1765-1769.	1.4	2
7	Evolutionary histories and antimicrobial resistance in <i>Shigella flexneri</i> and <i>Shigella sonnei</i> in Southeast Asia. <i>Access Microbiology</i> , 2022, 4, .	0.5	0
8	Impact of delayed processing of positive blood cultures on organism detection: a prospective multi-centre study. <i>BMC Infectious Diseases</i> , 2022, 22, .	2.9	5
9	Genetic background of Cambodian pneumococcal carriage isolates following pneumococcal conjugate vaccine 13. <i>Microbial Genomics</i> , 2022, 8, .	2.0	0
10	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. <i>International Journal of Infectious Diseases</i> , 2022, 122, 612-621.	3.3	6
11	Evaluation of trends in hospital antimicrobial use in the Lao PDR using repeated point-prevalence surveys-evidence to improve treatment guideline use. <i>The Lancet Regional Health - Western Pacific</i> , 2022, 27, 100531.	2.9	4
12	Impact of low blood culture usage on rates of antimicrobial resistance. <i>Journal of Infection</i> , 2021, 82, 355-362.	3.3	12
13	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
14	Impact of delays to incubation and storage temperature on blood culture results: a multi-centre study. <i>BMC Infectious Diseases</i> , 2021, 21, 173.	2.9	13
15	Boosting heritability: estimating the genetic component of phenotypic variation with multiple sample splitting. <i>BMC Bioinformatics</i> , 2021, 22, 164.	2.6	7
16	Evolutionary histories and antimicrobial resistance in <i>Shigella flexneri</i> and <i>Shigella sonnei</i> in Southeast Asia. <i>Communications Biology</i> , 2021, 4, 353.	4.4	17
17	Improving Treatment and Outcomes for Melioidosis in Children, Northern Cambodia, 2009–2018. <i>Emerging Infectious Diseases</i> , 2021, 27, 1169-1172.	4.3	8
18	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

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19	Surveillance strategies using routine microbiology for antimicrobial resistance in low- and middle-income countries. <i>Clinical Microbiology and Infection</i> , 2021, 27, 1391-1399.	6.0	20
20	Laboratory informatics capacity for effective antimicrobial resistance surveillance in resource-limited settings. <i>Lancet Infectious Diseases</i> , The, 2021, 21, e170-e174.	9.1	13
21	A multi-country study using MALDI-TOF mass spectrometry for rapid identification of <i>Burkholderia pseudomallei</i> . <i>BMC Microbiology</i> , 2021, 21, 213.	3.3	7
22	The role of interspecies recombination in the evolution of antibiotic-resistant pneumococci. <i>ELife</i> , 2021, 10, .	6.0	21
23	Predictors of disease severity in children presenting from the community with febrile illnesses: a systematic review of prognostic studies. <i>BMJ Global Health</i> , 2021, 6, e003451.	4.7	13
24	Antimicrobial resistance detection in Southeast Asian hospitals is critically important from both patient and societal perspectives, but what is its cost?. <i>PLOS Global Public Health</i> , 2021, 1, e0000018.	1.6	6
25	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
26	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. <i>Frontiers in Microbiology</i> , 2021, 12, 695027.	3.5	7
27	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.	3.3	2
28	The evolutionary history of <i>Shigella flexneri</i> serotype 6 in Asia. <i>Microbial Genomics</i> , 2021, 7, .	2.0	3
29	Impact of 13-Valent Pneumococcal Conjugate Vaccine on Colonization and Invasive Disease in Cambodian Children. <i>Clinical Infectious Diseases</i> , 2020, 70, 1580-1588.	5.8	21
30	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
31	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
32	MALDI-TOF mass spectrometry for sub-typing of <i>Streptococcus pneumoniae</i> . <i>BMC Microbiology</i> , 2020, 20, 367.	3.3	9
33	Case-based surveillance of antimicrobial resistance in the ACORN (A Clinically Oriented Antimicrobial) Tj ETQq1 1 0,784314 rgBT /Overl	2.1	4
34	Detection of colonisation by extended-spectrum beta-lactamase or carbapenemase producing Enterobacterales from frozen stool specimens. <i>BMC Research Notes</i> , 2020, 13, 429.	1.4	1
35	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
36	Salmonella Typhi and Paratyphi A infections in Cambodian children, 2012–2016. <i>International Journal of Infectious Diseases</i> , 2020, 97, 334-336.	3.3	7

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37	Setting priorities for patient-centered surveillance of drug-resistant infections. International Journal of Infectious Diseases, 2020, 97, 60-65.	3.3	4
38	Antimicrobial Resistance Surveillance in Low- and Middle-Income Countries: Progress and Challenges in Eight South Asian and Southeast Asian Countries. Clinical Microbiology Reviews, 2020, 33, .	13.6	105
39	Evaluation of the Coverage of 3 Antibiotic Regimens for Neonatal Sepsis in the Hospital Setting Across Asian Countries. JAMA Network Open, 2020, 3, e1921124.	5.9	11
40	<i>Elizabethkingia anophelis</i> Infection in Infants, Cambodia, 2012–2018. Emerging Infectious Diseases, 2020, 26, 320-322.	4.3	17
41	Genomic surveillance for hypervirulence and multi-drug resistance in invasive <i>Klebsiella pneumoniae</i> from South and Southeast Asia. Genome Medicine, 2020, 12, 11.	8.2	178
42	The spread of chloramphenicol-resistant <i>Neisseria meningitidis</i> in Southeast Asia. International Journal of Infectious Diseases, 2020, 95, 198-203.	3.3	6
43	Causes of fever in primary care in Southeast Asia and the performance of C-reactive protein in discriminating bacterial from viral pathogens. International Journal of Infectious Diseases, 2020, 96, 334-342.	3.3	8
44	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
45	Leapfrogging laboratories: the promise and pitfalls of high-tech solutions for antimicrobial resistance surveillance in low-income settings. BMJ Global Health, 2020, 5, e003622.	4.7	30
46	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
47	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
48	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
49	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
50	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
51	Global patterns in monthly activity of influenza virus, respiratory syncytial virus, parainfluenza virus, and metapneumovirus: a systematic analysis. The Lancet Global Health, 2019, 7, e1031-e1045.	6.3	266
52	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
53	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. PLoS Neglected Tropical Diseases, 2019, 13, e0007421.	3.0	51
54	Dissecting the molecular evolution of fluoroquinolone-resistant <i>Shigella sonnei</i> . Nature Communications, 2019, 10, 4828.	12.8	41

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55	Standardising the reporting of microbiology and antimicrobial susceptibility data. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 1163-1164.	9.1	8
56	'Candidatus <i>Ornithobacterium hominis</i> ': insights gained from draft genomes obtained from nasopharyngeal swabs. <i>Microbial Genomics</i> , 2019, 5, .	2.0	16
57	Antimicrobial resistance in Cambodia: a review. <i>International Journal of Infectious Diseases</i> , 2019, 85, 98-107.	3.3	27
58	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
59	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
60	Time to switch from CLSI to EUCAST? A Southeast Asian perspective. <i>Clinical Microbiology and Infection</i> , 2019, 25, 782-785.	6.0	18
61	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
62	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
63	Seroprevalence of Dengue Virus and Rickettsial Infections in Cambodian Children. <i>American Journal of Tropical Medicine and Hygiene</i> , 2019, 100, 635-638.	1.4	8
64	Pantonâ€“Valentine leucocidin is the key determinant of <i>Staphylococcus aureus</i> pyomyositis in a bacterial GWAS. <i>ELife</i> , 2019, 8, .	6.0	56
65	Transmission dynamics and control of multidrug-resistant <i>Klebsiella pneumoniae</i> in neonates in a developing country. <i>ELife</i> , 2019, 8, .	6.0	17
66	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
67	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
68	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
69	Climatic drivers of melioidosis in Laos and Cambodia: a 16-year case series analysis. <i>Lancet Planetary Health</i> , The, 2018, 2, e334-e343.	11.4	23
70	Utilization of a clinical microbiology service at a Cambodian paediatric hospital and its impact on appropriate antimicrobial prescribing. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 509-516.	3.0	14
71	Grading antimicrobial susceptibility data quality: room for improvement. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 603-604.	9.1	12
72	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.	3.5	24

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73	Antimicrobial Resistance in Invasive Bacterial Infections in Hospitalized Children, Cambodia, 2007–2016. <i>Emerging Infectious Diseases</i> , 2018, 24, 841-851.	4.3	50
74	Serotype Distribution of Clinical <i>Streptococcus pneumoniae</i> Isolates before the Introduction of the 13-Valent Pneumococcal Conjugate Vaccine in Cambodia. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 98, 791-796.	1.4	7
75	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
76	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
77	Prospective surveillance of healthcare associated infections in a Cambodian pediatric hospital. <i>Antimicrobial Resistance and Infection Control</i> , 2017, 6, 16.	4.1	12
78	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
79	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
80	A current perspective on antimicrobial resistance in Southeast Asia. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 2963-2972.	3.0	139
81	The aetiologies of central nervous system infections in hospitalised Cambodian children. <i>BMC Infectious Diseases</i> , 2017, 17, 806.	2.9	26
82	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
83	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
84	Genome-wide identification of lineage and locus specific variation associated with pneumococcal carriage duration. <i>ELife</i> , 2017, 6, .	6.0	95
85	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
86	Molecular Surveillance Identifies Multiple Transmissions of Typhoid in West Africa. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004781.	3.0	46
87	The Molecular and Spatial Epidemiology of Typhoid Fever in Rural Cambodia. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004785.	3.0	40
88	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
89	Lack of Utility of Nasopharyngeal Swabs for Diagnosis of <i>Burkholderia pseudomallei</i> Pneumonia in Paediatric Patients. <i>Journal of Tropical Pediatrics</i> , 2016, 62, 328-330.	1.5	0
90	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

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91	Understanding pneumococcal serotype 1 biology through population genomic analysis. BMC Infectious Diseases, 2016, 16, 649.	2.9	22
92	A retrospective analysis of melioidosis in Cambodian children, 2009–2013. BMC Infectious Diseases, 2016, 16, 688.	2.9	29
93	Use of Blood Smears and Dried Blood Spots for Polymerase Chain Reaction–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
94	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
95	Evolutionary History of the Global Emergence of the Escherichia coli Epidemic Clone ST131. MBio, 2016, 7, e02162.	4.1	289
96	Capsular Typing Method for Streptococcus agalactiae Using Whole-Genome Sequence Data. Journal of Clinical Microbiology, 2016, 54, 1388-1390.	3.9	35
97	Horizontal DNA Transfer Mechanisms of Bacteria as Weapons of Intragenomic Conflict. PLoS Biology, 2016, 14, e1002394.	5.6	127
98	Characterisation of Invasive Streptococcus pneumoniae Isolated from Cambodian Children between 2007 – 2012. PLoS ONE, 2016, 11, e0159358.	2.5	15
99	Molecular Epidemiology of Group A Streptococcus Infections in Cambodian Children, 2007–2012. Pediatric Infectious Disease Journal, 2015, 34, 1414-1415.	2.0	3
100	Necrotizing fasciitis complicating snakebite in Cambodia. IDCases, 2015, 2, 86-87.	0.9	3
101	Performance of C-reactive protein and procalcitonin to distinguish viral from bacterial and malarial causes of fever in Southeast Asia. BMC Infectious Diseases, 2015, 15, 511.	2.9	103
102	Pneumococcal Infection among Children before Introduction of 13-Valent Pneumococcal Conjugate Vaccine, Cambodia. Emerging Infectious Diseases, 2015, 21, 2080-2083.	4.3	19
103	A Novel Technique for Detecting Antibiotic-Resistant Typhoid from Rapid Diagnostic Tests. Journal of Clinical Microbiology, 2015, 53, 1758-1760.	3.9	7
104	Clinically and Microbiologically Derived Azithromycin Susceptibility Breakpoints for Salmonella enterica Serovars Typhi and Paratyphi A. Antimicrobial Agents and Chemotherapy, 2015, 59, 2756-2764.	3.2	44
105	Climate induces seasonality in pneumococcal transmission. Scientific Reports, 2015, 5, 11344.	3.3	41
106	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
107	Phylogeographical analysis of the dominant multidrug-resistant H58 clade of Salmonella Typhi identifies inter- and intracontinental transmission events. Nature Genetics, 2015, 47, 632-639.	21.4	403
108	Diagnostic Accuracy Assessment of Immunochromatographic Tests for the Rapid Detection of Antibodies Against Orientia tsutsugamushi Using Paired Acute and Convalescent Specimens. American Journal of Tropical Medicine and Hygiene, 2015, 93, 1168-1171.	1.4	12

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109	Introduction and establishment of fluoroquinolone-resistant <i>Shigella sonnei</i> into Bhutan. <i>Microbial Genomics</i> , 2015, 1, e000042.	2.0	11
110	Ophthalmic infections in children presenting to Angkor Hospital for Children, Siem Reap, Cambodia. <i>BMC Research Notes</i> , 2014, 7, 784.	1.4	9
111	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
112	Trimethoprim/sulfamethoxazole resistance in <i>Burkholderia pseudomallei</i> . <i>International Journal of Antimicrobial Agents</i> , 2014, 44, 368-369.	2.5	24
113	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
114	Reagent and laboratory contamination can critically impact sequence-based microbiome analyses. <i>BMC Biology</i> , 2014, 12, 87.	3.8	2,677
115	Two-phase importance sampling for inference about transmission trees. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20141324.	2.6	22
116	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
117	Dense genomic sampling identifies highways of pneumococcal recombination. <i>Nature Genetics</i> , 2014, 46, 305-309.	21.4	371
118	Two fatal cases of melioidosis on the Thai-Myanmar border. <i>F1000Research</i> , 2014, 3, 4.	1.6	4
119	Two fatal cases of melioidosis on the Thai-Myanmar border. <i>F1000Research</i> , 2014, 3, 4.	1.6	7
120	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
121	Respiratory virus surveillance in hospitalised pneumonia patients on the Thailand-Myanmar border. <i>BMC Infectious Diseases</i> , 2013, 13, 434.	2.9	27
122	A three year descriptive study of early onset neonatal sepsis in a refugee population on the Thailand Myanmar border. <i>BMC Infectious Diseases</i> , 2013, 13, 601.	2.9	12
123	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
124	Rectal pH in Well and Unwell Infants. <i>Journal of Tropical Pediatrics</i> , 2013, 59, 162-162.	1.5	0
125	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
126	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

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127	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
128	Neonatal Intensive Care in a Karen Refugee Camp: A 4 Year Descriptive Study. PLoS ONE, 2013, 8, e72721.	2.5	43
129	Rectal pH in Well and Unwell Infants. Journal of Tropical Pediatrics, 2012, 58, 311-313.	1.5	8
130	Assessment of Streptococcus pneumoniae pilus islet-1 prevalence in carried and transmitted isolates from motherâ€“infant pairs on the Thailandâ€“Burma border. Clinical Microbiology and Infection, 2012, 18, 970-975.	6.0	21
131	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
132	Aeromonasspp. Bacteremia in Pregnant Women, Thailandâ€“Myanmar Border, 2011. Emerging Infectious Diseases, 2012, 18, 1522-3.	4.3	5
133	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. BMC Infectious Diseases, 2012, 12, 34.	2.9	36
134	A Longitudinal Study of Streptococcus pneumoniae Carriage in a Cohort of Infants and Their Mothers on the Thailand-Myanmar Border. PLoS ONE, 2012, 7, e38271.	2.5	129
135	Improved Detection of Nasopharyngeal Cocolonization by Multiple Pneumococcal Serotypes by Use of Latex Agglutination or Molecular Serotyping by Microarray. Journal of Clinical Microbiology, 2011, 49, 1784-1789.	3.9	134
136	Susceptibility of communityâ€“acquired pathogens to antibiotics in Africa and Asia in neonates â€“ an alarmingly short review. Tropical Medicine and International Health, 2011, 16, 145-151.	2.3	36
137	Antimicrobial susceptibility of bacterial isolates from community acquired infections in Subâ€“Saharan Africa and Asian low and middle income countries. Tropical Medicine and International Health, 2011, 16, 1167-1179.	2.3	67
138	Susceptibility of bacterial isolates from communityâ€“acquired infections in subâ€“Saharan Africa and Asia to macrolide antibiotics. Tropical Medicine and International Health, 2011, 16, 1192-1205.	2.3	15
139	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
140	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
141	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
142	Influenza in Refugees on the Thailandâ€“Myanmar Border, Mayâ€“October 2009. Emerging Infectious Diseases, 2010, 16, 1366-1372.	4.3	8
143	Duplex Real-Time PCR Assay for Detection of <i>Streptococcus pneumoniae</i> in Clinical Samples and Determination of Penicillin Susceptibility. Journal of Clinical Microbiology, 2008, 46, 2751-2758.	3.9	35
144	Dermal absorption of isopropyl alcohol from a commercial hand rub: implications for its use in hand decontamination. Journal of Hospital Infection, 2004, 56, 287-290.	2.9	52

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145	A case of human syngamosis. Travel Medicine and Infectious Disease, 2003, 1, 231-233.	3.0	2