

# Duy Thanh Pham

## List of Publications by Year in descending order

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75  
papers

3,727  
citations

117453

34  
h-index

143772

57  
g-index

85  
all docs

85  
docs citations

85  
times ranked

4090  
citing authors

#	ARTICLE	IF	CITATIONS
1	A fatal outbreak of neonatal sepsis caused by mcr-10-carrying <i>Enterobacter kobei</i> in a tertiary care hospital in Nepal. <i>Journal of Hospital Infection</i> , 2022, 125, 60-66.	1.4	6
2	Evolutionary histories and antimicrobial resistance in <i>Shigella flexneri</i> and <i>Shigella sonnei</i> in Southeast Asia. <i>Access Microbiology</i> , 2022, 4, .	0.2	0
3	Spontaneous Emergence of Azithromycin Resistance in Independent Lineages of <i>Salmonella</i> Typhi in Northern India. <i>Clinical Infectious Diseases</i> , 2021, 72, e120-e127.	2.9	39
4	Colonization with <i>Staphylococcus aureus</i> and <i>Klebsiella pneumoniae</i> causes infections in a Vietnamese intensive care unit. <i>Microbial Genomics</i> , 2021, 7, .	1.0	10
5	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.	2.0	17
6	Emerging carbapenem-resistant <i>Klebsiella pneumoniae</i> sequence type 16 causing multiple outbreaks in a tertiary hospital in southern Vietnam. <i>Microbial Genomics</i> , 2021, 7, .	1.0	14
7	Bactericidal activities and post-antibiotic effects of ofloxacin and ceftriaxone against drug-resistant <i>Salmonella enterica</i> serovar Typhi. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2606-2609.	1.3	1
8	The genomic epidemiology of multi-drug resistant invasive non-typhoidal <i>Salmonella</i> in selected sub-Saharan African countries. <i>BMJ Global Health</i> , 2021, 6, e005659.	2.0	16
9	A genomic snapshot of <i>Salmonella enterica</i> serovar Typhi in Colombia. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009755.	1.3	7
10	Azithromycin and cefixime combination versus azithromycin alone for the out-patient treatment of clinically suspected or confirmed uncomplicated typhoid fever in South Asia: a randomised controlled trial protocol. <i>Wellcome Open Research</i> , 2021, 6, 207.	0.9	0
11	Genomic insights into the circulation of pandemic fluoroquinolone-resistant extra-intestinal pathogenic <i>Escherichia coli</i> ST1193 in Vietnam. <i>Microbial Genomics</i> , 2021, 7, .	1.0	8
12	The evolutionary history of <i>Shigella flexneri</i> serotype 6 in Asia. <i>Microbial Genomics</i> , 2021, 7, .	1.0	3
13	Gallbladder carriage generates genetic variation and genome degradation in <i>Salmonella</i> Typhi. <i>PLoS Pathogens</i> , 2020, 16, e1008998.	2.1	20
14	Pathogenic <i>Escherichia coli</i> Possess Elevated Growth Rates under Exposure to Sub-Inhibitory Concentrations of Azithromycin. <i>Antibiotics</i> , 2020, 9, 735.	1.5	5
15	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008040.	1.3	6
16	Commensal <i>Escherichia coli</i> are a reservoir for the transfer of XDR plasmids into epidemic fluoroquinolone-resistant <i>Shigella sonnei</i> . <i>Nature Microbiology</i> , 2020, 5, 256-264.	5.9	43
17	The emergence of azithromycin-resistant <i>Salmonella</i> Typhi in Nepal. <i>JAC-Antimicrobial Resistance</i> , 2020, 2, dlaa109.	0.9	30
18	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0

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19	Surveillance of Salmonella enterica serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0
20	Surveillance of Salmonella enterica serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0
21	Surveillance of Salmonella enterica serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0
22	Dissecting the molecular evolution of fluoroquinolone-resistant Shigella sonnei. Nature Communications, 2019, 10, 4828.	5.8	41
23	Clinical and laboratory-induced colistin-resistance mechanisms in Acinetobacter baumannii. Microbial Genomics, 2019, 5, .	1.0	30
24	Endemic fluoroquinolone-resistant Salmonella enterica serovar Kentucky ST198 in northern India. Microbial Genomics, 2019, 5, .	1.0	21
25	Azithromycin Resistance in Shigella spp. in Southeast Asia. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	37
26	Assessing gut microbiota perturbations during the early phase of infectious diarrhea in Vietnamese children. Gut Microbes, 2018, 9, 38-54.	4.3	66
27	The phylogeography and incidence of multi-drug resistant typhoid fever in sub-Saharan Africa. Nature Communications, 2018, 9, 5094.	5.8	98
28	The Control of Typhoid Fever in Vietnam. American Journal of Tropical Medicine and Hygiene, 2018, 99, 72-78.	0.6	23
29	New Variant of Multidrug-Resistant <i>Salmonella enterica</i> Serovar Typhimurium Associated with Invasive Disease in Immunocompromised Patients in Vietnam. MBio, 2018, 9, .	1.8	53
30	Excess body weight and age associated with the carriage of fluoroquinolone and third-generation cephalosporin resistance genes in commensal Escherichia coli from a cohort of urban Vietnamese children. Journal of Medical Microbiology, 2018, 67, 1457-1466.	0.7	8
31	An evaluation of purified Salmonella Typhi protein antigens for the serological diagnosis of acute typhoid fever. Journal of Infection, 2017, 75, 104-114.	1.7	23
32	A universal genome sequencing method for rotavirus A from human fecal samples which identifies segment reassortment and multi-genotype mixed infection. BMC Genomics, 2017, 18, 324.	1.2	10
33	The emergence of G8P[8] rotavirus group A across Vietnam. Virus Evolution, 2017, 3, .	2.2	0
34	Whole Genome Sequence Analysis of Salmonella Typhi Isolated in Thailand before and after the Introduction of a National Immunization Program. PLoS Neglected Tropical Diseases, 2017, 11, e0005274.	1.3	26
35	Phenotypic and genotypic characteristics of ESBL and AmpC producing organisms associated with bacteraemia in Ho Chi Minh City, Vietnam. Antimicrobial Resistance and Infection Control, 2017, 6, 105.	1.5	15
36	Evaluation of the Clinical and Microbiological Response to Salmonella Paratyphi A Infection in the First Paratyphoid Human Challenge Model. Clinical Infectious Diseases, 2017, 64, 1066-1073.	2.9	60

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37	A novel ciprofloxacin-resistant subclade of H58 Salmonella Typhi is associated with fluoroquinolone treatment failure. <i>ELife</i> , 2016, 5, e14003.	2.8	111
38	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.	3.9	84
39	The Molecular and Spatial Epidemiology of Typhoid Fever in Rural Cambodia. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004785.	1.3	40
40	The induction and identification of novel Colistin resistance mutations in <i>Acinetobacter baumannii</i> and their implications. <i>Scientific Reports</i> , 2016, 6, 28291.	1.6	88
41	Inducible colistin resistance via a disrupted plasmid-borne <i>mcr-1</i> gene in a 2008 Vietnamese <i>Shigella sonnei</i> isolate. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 2314-2317.	1.3	82
42	Garifloxacin versus ceftriaxone for uncomplicated enteric fever in Nepal: an open-label, two-centre, randomised controlled trial. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 535-545.	4.6	54
43	The genomic signatures of <i>Shigella</i> evolution, adaptation and geographical spread. <i>Nature Reviews Microbiology</i> , 2016, 14, 235-250.	13.6	142
44	Clinical implications of reduced susceptibility to fluoroquinolones in paediatric <i>Shigella sonnei</i> and <i>Shigella flexneri</i> infections. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 807-815.	1.3	13
45	Repeated local emergence of carbapenem-resistant <i>Acinetobacter baumannii</i> in a single hospital ward. <i>Microbial Genomics</i> , 2016, 2, e000050.	1.0	65
46	The Ecological Dynamics of Fecal Contamination and Salmonella Typhi and Salmonella Paratyphi A in Municipal Kathmandu Drinking Water. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004346.	1.3	70
47	A high-resolution genomic analysis of multidrug-resistant hospital outbreaks of <i>Klebsiella pneumoniae</i> . <i>EMBO Molecular Medicine</i> , 2015, 7, 227-239.	3.3	104
48	The Rising Dominance of <i>Shigella sonnei</i> : An Intercontinental Shift in the Etiology of Bacillary Dysentery. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003708.	1.3	140
49	Species-wide whole genome sequencing reveals historical global spread and recent local persistence in <i>Shigella flexneri</i> . <i>ELife</i> , 2015, 4, e07335.	2.8	94
50	The diagnostic accuracy of three rapid diagnostic tests for typhoid fever at <i>C/hittagong</i> <i>M/edical College</i> <i>H/ospital</i> , <i>C/hittagong</i> , <i>B/angladesh</i> . <i>Tropical Medicine and International Health</i> , 2015, 20, 1376-1384.	1.0	22
51	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.	1.4	44
52	Intercontinental dissemination of azithromycin-resistant shigellosis through sexual transmission: a cross-sectional study. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 913-921.	4.6	204
53	Phylogeographical analysis of the dominant multidrug-resistant H58 clade of <i>Salmonella Typhi</i> identifies inter- and intracontinental transmission events. <i>Nature Genetics</i> , 2015, 47, 632-639.	9.4	403
54	A Prospective Multi-Center Observational Study of Children Hospitalized with Diarrhea in Ho Chi Minh City, Vietnam. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 92, 1045-1052.	0.6	56

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55	Introduction and establishment of fluoroquinolone-resistant <i>Shigella sonnei</i> into Bhutan. <i>Microbial Genomics</i> , 2015, 1, e000042.	1.0	11
56	Ophthalmic infections in children presenting to Angkor Hospital for Children, Siem Reap, Cambodia. <i>BMC Research Notes</i> , 2014, 7, 784.	0.6	9
57	Evaluation of the Diagnostic Accuracy of a Typhoid IgM Flow Assay for the Diagnosis of Typhoid Fever in Cambodian Children Using a Bayesian Latent Class Model Assuming an Imperfect Gold Standard. <i>American Journal of Tropical Medicine and Hygiene</i> , 2014, 90, 114-120.	0.6	34
58	The validation and utility of a quantitative one-step multiplex RT real-time PCR targeting Rotavirus A and Norovirus. <i>Journal of Virological Methods</i> , 2013, 187, 138-143.	1.0	42
59	Whole-genome sequences of <i>Chlamydia trachomatis</i> directly from clinical samples without culture. <i>Genome Research</i> , 2013, 23, 855-866.	2.4	115
60	Identification of <i>Salmonella enterica</i> Serovar Typhi Genotypes by Use of Rapid Multiplex Ligation-Dependent Probe Amplification. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2950-2958.	1.8	9
61	Tracking the establishment of local endemic populations of an emergent enteric pathogen. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 17522-17527.	3.3	124
62	Fitness benefits in fluoroquinolone-resistant <i>Salmonella</i> Typhi in the absence of antimicrobial pressure. <i>ELife</i> , 2013, 2, e01229.	2.8	103
63	The characterization of ESBL genes in <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> causing nosocomial infections in Vietnam. <i>Journal of Infection in Developing Countries</i> , 2013, 7, 922-928.	0.5	21
64	Highly Resistant <i>Salmonella enterica</i> Serovar Typhi with a Novel <i>gyrA</i> Mutation Raises Questions about the Long-Term Efficacy of Older Fluoroquinolones for Treating Typhoid Fever. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 2761-2762.	1.4	32
65	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
66	The Microbiological and Clinical Characteristics of Invasive <i>Salmonella</i> in Gallbladders from Cholecystectomy Patients in Kathmandu, Nepal. <i>PLoS ONE</i> , 2012, 7, e47342.	1.1	56
67	Combined high-resolution genotyping and geospatial analysis reveals modes of endemic urban typhoid fever transmission. <i>Open Biology</i> , 2011, 1, 110008.	1.5	112
68	Emergence of a Globally Dominant IncHI1 Plasmid Type Associated with Multiple Drug Resistant Typhoid. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1245.	1.3	114
69	Temporal Fluctuation of Multidrug Resistant <i>Salmonella</i> Typhi Haplotypes in the Mekong River Delta Region of Vietnam. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e929.	1.3	47
70	A Multi-Center Randomized Trial to Assess the Efficacy of Gatifloxacin versus Ciprofloxacin for the Treatment of Shigellosis in Vietnamese Children. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1264.	1.3	22
71	The Burden and Characteristics of Enteric Fever at a Healthcare Facility in a Densely Populated Area of Kathmandu. <i>PLoS ONE</i> , 2010, 5, e13988.	1.1	58
72	The Sudden Dominance of bla <sub>CTX-M</sub> Harboring Plasmids in <i>Shigella</i> spp. Circulating in Southern Vietnam. <i>PLoS Neglected Tropical Diseases</i> , 2010, 4, e702.	1.3	48

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73	Suitable Disk Antimicrobial Susceptibility Breakpoints Defining <i>Salmonella enterica</i> Serovar Typhi Isolates with Reduced Susceptibility to Fluoroquinolones. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 5201-5208.	1.4	45
74	A changing picture of shigellosis in southern Vietnam: shifting species dominance, antimicrobial susceptibility and clinical presentation. <i>BMC Infectious Diseases</i> , 2009, 9, 204.	1.3	111
75	Azithromycin and cefixime combination versus azithromycin alone for the out-patient treatment of clinically suspected or confirmed uncomplicated typhoid fever in South Asia: a randomised controlled trial protocol. <i>Wellcome Open Research</i> , 0, 6, 207.	0.9	6