

Duy Thanh Pham

List of Publications by Year in descending order

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Version: 2024-02-01

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	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.	9.4	403
2	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
3	The genomic signatures of <i>Shigella</i> evolution, adaptation and geographical spread. <i>Nature Reviews Microbiology</i> , 2016, 14, 235-250.	13.6	142
4	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
5	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
6	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
7	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
8	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
9	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
10	A novel ciprofloxacin-resistant subclade of H58 <i>Salmonella</i> Typhi is associated with fluoroquinolone treatment failure. <i>ELife</i> , 2016, 5, e14003.	2.8	111
11	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
12	Fitness benefits in fluoroquinolone-resistant <i>Salmonella</i> Typhi in the absence of antimicrobial pressure. <i>ELife</i> , 2013, 2, e01229.	2.8	103
13	The phylogeography and incidence of multi-drug resistant typhoid fever in sub-Saharan Africa. <i>Nature Communications</i> , 2018, 9, 5094.	5.8	98
14	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
15	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
16	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
17	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
18	The Ecological Dynamics of Fecal Contamination and <i>Salmonella</i> Typhi and <i>Salmonella</i> Paratyphi A in Municipal Kathmandu Drinking Water. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004346.	1.3	70

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19	Assessing gut microbiota perturbations during the early phase of infectious diarrhea in Vietnamese children. <i>Gut Microbes</i> , 2018, 9, 38-54.	4.3	66
20	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
21	Evaluation of the Clinical and Microbiological Response to <i>Salmonella Paratyphi A</i> Infection in the First Paratyphoid Human Challenge Model. <i>Clinical Infectious Diseases</i> , 2017, 64, 1066-1073.	2.9	60
22	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
23	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
24	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
25	Gatifloxacin 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
26	New Variant of Multidrug-Resistant <i>Salmonella enterica</i> Serovar Typhimurium Associated with Invasive Disease in Immunocompromised Patients in Vietnam. <i>MBio</i> , 2018, 9, .	1.8	53
27	The Sudden Dominance of bla _{CTX-M} Harboring Plasmids in <i>Shigella</i> spp. Circulating in Southern Vietnam. <i>PLoS Neglected Tropical Diseases</i> , 2010, 4, e702.	1.3	48
28	Temporal Fluctuation of Multidrug Resistant <i>Salmonella Typhi</i> Haplotypes in the Mekong River Delta Region of Vietnam. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e929.	1.3	47
29	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
30	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
31	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
32	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
33	Dissecting the molecular evolution of fluoroquinolone-resistant <i>Shigella sonnei</i> . <i>Nature Communications</i> , 2019, 10, 4828.	5.8	41
34	The Molecular and Spatial Epidemiology of Typhoid Fever in Rural Cambodia. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004785.	1.3	40
35	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
36	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

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37	Azithromycin Resistance in <i>Shigella</i> spp. in Southeast Asia. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	1.4	37
38	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
39	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
40	The emergence of azithromycin-resistant <i>Salmonella</i> Typhi in Nepal. <i>JAC-Antimicrobial Resistance</i> , 2020, 2, dlaa109.	0.9	30
41	Clinical and laboratory-induced colistin-resistance mechanisms in <i>Acinetobacter baumannii</i> . <i>Microbial Genomics</i> , 2019, 5, .	1.0	30
42	Whole Genome Sequence Analysis of <i>Salmonella</i> Typhi Isolated in Thailand before and after the Introduction of a National Immunization Program. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005274.	1.3	26
43	An evaluation of purified <i>Salmonella</i> Typhi protein antigens for the serological diagnosis of acute typhoid fever. <i>Journal of Infection</i> , 2017, 75, 104-114.	1.7	23
44	The Control of Typhoid Fever in Vietnam. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 99, 72-78.	0.6	23
45	The diagnostic accuracy of three rapid diagnostic tests for typhoid fever at <i>C</i> hittagong <i>M</i> edical <i>C</i> ollege <i>H</i> ospital, <i>C</i> hittagong, <i>B</i> angladesh. <i>Tropical Medicine and International Health</i> , 2015, 20, 1376-1384.	1.0	22
46	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
47	Endemic fluoroquinolone-resistant <i>Salmonella enterica</i> serovar Kentucky ST198 in northern India. <i>Microbial Genomics</i> , 2019, 5, .	1.0	21
48	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
49	Gallbladder carriage generates genetic variation and genome degradation in <i>Salmonella</i> Typhi. <i>PLoS Pathogens</i> , 2020, 16, e1008998.	2.1	20
50	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
51	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
52	Phenotypic and genotypic characteristics of ESBL and AmpC producing organisms associated with bacteraemia in Ho Chi Minh City, Vietnam. <i>Antimicrobial Resistance and Infection Control</i> , 2017, 6, 105.	1.5	15
53	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
54	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

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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	A universal genome sequencing method for rotavirus A from human fecal samples which identifies segment reassortment and multi-genotype mixed infection. <i>BMC Genomics</i> , 2017, 18, 324.	1.2	10
57	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
58	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
59	Ophthalmic infections in children presenting to Angkor Hospital for Children, Siem Reap, Cambodia. <i>BMC Research Notes</i> , 2014, 7, 784.	0.6	9
60	Excess body weight and age associated with the carriage of fluoroquinolone and third-generation cephalosporin resistance genes in commensal <i>Escherichia coli</i> from a cohort of urban Vietnamese children. <i>Journal of Medical Microbiology</i> , 2018, 67, 1457-1466.	0.7	8
61	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
62	A genomic snapshot of <i>Salmonella enterica</i> serovar Typhi in Colombia. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009755.	1.3	7
63	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008040.	1.3	6
64	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
65	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
66	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
67	The evolutionary history of <i>Shigella flexneri</i> serotype 6 in Asia. <i>Microbial Genomics</i> , 2021, 7, .	1.0	3
68	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
69	The emergence of G8P[8] rotavirus group A across Vietnam. <i>Virus Evolution</i> , 2017, 3, .	2.2	0
70	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
71	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0
72	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0

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73	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0
74	Surveillance of <i>Salmonella enterica</i> serovar Typhi in Colombia, 2012–2015. , 2020, 14, e0008040.		0
75	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