

# Hao Chung The

## List of Publications by Year in descending order

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Version: 2024-02-01

24  
papers

995  
citations

623188

14  
h-index

610482

24  
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26  
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26  
docs citations

26  
times ranked

1544  
citing authors

#	ARTICLE	IF	CITATIONS
1	The genomic signatures of <i>Shigella</i> evolution, adaptation and geographical spread. <i>Nature Reviews Microbiology</i> , 2016, 14, 235-250.	13.6	142
2	Recent insights into <i>Shigella</i> : a major contributor to the global diarrhoeal disease burden. <i>Current Opinion in Infectious Diseases</i> , 2018, 31, 449-454.	1.3	134
3	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
4	Rapid environmental effects on gut nematode susceptibility in rewilded mice. <i>PLoS Biology</i> , 2018, 16, e2004108.	2.6	97
5	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
6	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
7	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
8	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
9	Dissecting the molecular evolution of fluoroquinolone-resistant <i>Shigella sonnei</i> . <i>Nature Communications</i> , 2019, 10, 4828.	5.8	41
10	Azithromycin Resistance in <i>Shigella</i> spp. in Southeast Asia. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	1.4	37
11	A Double-blind, Randomized, Placebo-controlled Trial of <i>Lactobacillus acidophilus</i> for the Treatment of Acute Watery Diarrhea in Vietnamese Children. <i>Pediatric Infectious Disease Journal</i> , 2018, 37, 35-42.	1.1	24
12	Dynamic of the human gut microbiome under infectious diarrhea. <i>Current Opinion in Microbiology</i> , 2022, 66, 79-85.	2.3	23
13	Genomic Serotyping, Clinical Manifestations, and Antimicrobial Resistance of Nontyphoidal <i>Salmonella</i> Gastroenteritis in Hospitalized Children in Ho Chi Minh City, Vietnam. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	1.8	18
14	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
15	Out of Asia: the independent rise and global spread of fluoroquinolone-resistant <i>Shigella</i> . <i>Microbial Genomics</i> , 2018, 4, .	1.0	14
16	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
17	Introduction and establishment of fluoroquinolone-resistant <i>Shigella sonnei</i> into Bhutan. <i>Microbial Genomics</i> , 2015, 1, e000042.	1.0	11
18	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

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19	The identification of novel immunogenic antigens as potential Shigella vaccine components. <i>Genome Medicine</i> , 2021, 13, 8.	3.6	9
20	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
21	Exploring the Genomic Diversity and Antimicrobial Susceptibility of <i>Bifidobacterium pseudocatenulatum</i> in a Vietnamese Population. <i>Microbiology Spectrum</i> , 2021, 9, e0052621.	1.2	6
22	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
23	The Gut Microbiome of Healthy Vietnamese Adults and Children Is a Major Reservoir for Resistance Genes Against Critical Antimicrobials. <i>Journal of Infectious Diseases</i> , 2021, 224, S840-S847.	1.9	3
24	The evolutionary history of <i>Shigella flexneri</i> serotype 6 in Asia. <i>Microbial Genomics</i> , 2021, 7, .	1.0	3