

# Rujirat Hatrongjit

## List of Publications by Year in descending order

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Version: 2024-02-01

24  
papers

369  
citations

933447

10  
h-index

839539

18  
g-index

25  
all docs

25  
docs citations

25  
times ranked

305  
citing authors

#	ARTICLE	IF	CITATIONS
1	Streptococcus suis serotyping by a new multiplex PCR. Journal of Medical Microbiology, 2014, 63, 824-830.	1.8	75
2	Emergence of Streptococcus suis serotype 9 infection in humans. Journal of Microbiology, Immunology and Infection, 2017, 50, 545-546.	3.1	62
3	First human case report of sepsis due to infection with Streptococcus suis serotype 31 in Thailand. BMC Infectious Diseases, 2015, 15, 392.	2.9	43
4	Detection of plasmid-mediated colistin-resistant and carbapenem-resistant genes by multiplex PCR. MethodsX, 2018, 5, 532-536.	1.6	28
5	Tools for Molecular Epidemiology of Streptococcus suis. Pathogens, 2020, 9, 81.	2.8	21
6	Distribution and Molecular Characterization of <i>Escherichia coli</i> Harboring <i>mcr</i> Genes Isolated from Slaughtered Pigs in Thailand. Microbial Drug Resistance, 2021, 27, 971-979.	2.0	17
7	Klebsiella pneumoniae Complex Harboring <i>mcr</i> -1, <i>mcr</i> -7, and <i>mcr</i> -8 Isolates from Slaughtered Pigs in Thailand. Microorganisms, 2021, 9, 2436.	3.6	17
8	Genomic Analysis of Aeromonas veronii C198, a Novel <i>Mcr</i> -3.41-Harboring Isolate from a Patient with Septicemia in Thailand. Pathogens, 2020, 9, 1031.	2.8	14
9	Non-Penicillin-Susceptible Streptococcus suis Isolated from Humans. Pathogens, 2021, 10, 1178.	2.8	14
10	Genomic characterization of an emerging blaKPC-2 carrying Enterobacteriaceae clinical isolates in Thailand. Scientific Reports, 2019, 9, 18521.	3.3	12
11	Zoonotic infection and clonal dissemination of <i>Streptococcus equi</i> subspecies <i>zooepidemicus</i> sequence type 194 isolated from humans in Thailand. Transboundary and Emerging Diseases, 2022, 69, .	3.0	10
12	Genomic Characterization of Clinical Extensively Drug-Resistant Acinetobacter pittii Isolates. Microorganisms, 2021, 9, 242.	3.6	10
13	Multiplex PCR for identification of six clinically relevant streptococci. Journal of Medical Microbiology, 2017, 66, 1590-1595.	1.8	10
14	Genomic Characterization of Streptococcus suis Serotype 24 Clonal Complex 221/234 From Human Patients. Frontiers in Microbiology, 2021, 12, 812436.	3.5	7
15	Complete Genome Sequences of Four Extensively Drug-Resistant Acinetobacter baumannii Isolates from Thailand. Microbiology Resource Announcements, 2020, 9, .	0.6	6
16	Development of a multiplex PCR assay to detect the major clonal complexes of Streptococcus suis relevant to human infection. Journal of Medical Microbiology, 2016, 65, 392-396.	1.8	5
17	Development of a multiplex PCR for identification of $\beta$ -hemolytic streptococci relevant to human infections and serotype distribution of invasive Streptococcus agalactiae in Thailand. Molecular and Cellular Probes, 2017, 36, 10-14.	2.1	4
18	Draft Genome Sequence of Methicillin-Resistant Staphylococcus aureus Harboring Staphylococcal Cassette Chromosome <i>mec</i> Type IX, Isolated from a Fatal Bacteremic Pneumonia Case. Microbiology Resource Announcements, 2021, 10, e0061621.	0.6	3

#	ARTICLE	IF	CITATIONS
19	Fluoroquinolone resistance in non-typhoidal <i>Salmonella enterica</i> isolated from slaughtered pigs in Thailand. <i>Journal of Medical Microbiology</i> , 2021, 70, .	1.8	3
20	Whole Genome Analysis of Extensively Drug-Resistant <i>Acinetobacter baumannii</i> Clinical Isolates in Thailand. <i>Infectious Disorders - Drug Targets</i> , 2021, 21, e270421188042.	0.8	3
21	Application of random amplified polymorphism DNA and 16S-23S rDNA intergenic spacer polymerase chain reaction-restriction fragment length polymorphism to predict major <i>Streptococcus suis</i> clonal complexes isolated from humans and pigs. <i>Molecular and Cellular Probes</i> , 2019, 43, 34-39.	2.1	2
22	Distinguishing Clinical <i>Enterococcus faecium</i> Strains and Resistance to Vancomycin Using a Simple In-House Screening Test. <i>Antibiotics</i> , 2022, 11, 286.	3.7	2
23	Evaluation of in-house cefoxitin screening broth to determine methicillin-resistant staphylococci. <i>Heliyon</i> , 2022, 8, e08950.	3.2	1
24	Phenotypic and molecular characterization of $\beta$ -lactamase and plasmid-mediated quinolone resistance genes in <i>Klebsiella oxytoca</i> isolated from slaughtered pigs in Thailand. <i>Veterinary World</i> , 2022, 15, 309-315.	1.7	0