Rujirat Hatrongjit

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	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
2	Evaluation of in-house cefoxitin screening broth to determine methicillin-resistant staphylococci. Heliyon, 2022, 8, e08950.	3.2	1
3	Phenotypic and molecular characterization of β-lactamase and plasmid-mediated quinolone resistance genes in Klebsiella oxytoca isolated from slaughtered pigs in Thailand. Veterinary World, 2022, 15, 309-315.	1.7	0
4	Distinguishing Clinical Enterococcus faecium Strains and Resistance to Vancomycin Using a Simple In-House Screening Test. Antibiotics, 2022, 11, 286.	3.7	2
5	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
6	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
7	Fluoroquinolone resistance in non-typhoidal Salmonella enterica isolated from slaughtered pigs in Thailand. Journal of Medical Microbiology, 2021, 70, .	1.8	3
8	Whole Genome Analysis of Extensively Drug-Resistant Acinetobacter baumannii Clinical Isolates in Thailand. Infectious Disorders - Drug Targets, 2021, 21, e270421188042.	0.8	3
9	Non-Penicillin-Susceptible Streptococcus suis Isolated from Humans. Pathogens, 2021, 10, 1178.	2.8	14
10	Genomic Characterization of Clinical Extensively Drug-Resistant Acinetobacter pittii Isolates. Microorganisms, 2021, 9, 242.	3.6	10
11	Klebsiella pneumoniae Complex Harboring mcr-1, mcr-7, and mcr-8 Isolates from Slaughtered Pigs in Thailand. Microorganisms, 2021, 9, 2436.	3.6	17
12	Genomic Characterization of Streptococcus suis Serotype 24 Clonal Complex 221/234 From Human Patients. Frontiers in Microbiology, 2021, 12, 812436.	3.5	7
13	Complete Genome Sequences of Four Extensively Drug-Resistant Acinetobacter baumannii Isolates from Thailand. Microbiology Resource Announcements, 2020, 9, .	0.6	6
14	Genomic Analysis of Aeromonas veronii C198, a Novel Mcr-3.41-Harboring Isolate from a Patient with Septicemia in Thailand. Pathogens, 2020, 9, 1031.	2.8	14
15	Tools for Molecular Epidemiology of Streptococcus suis. Pathogens, 2020, 9, 81.	2.8	21
16	Genomic characterization of an emerging blaKPC-2 carrying Enterobacteriaceae clinical isolates in Thailand. Scientific Reports, 2019, 9, 18521.	3.3	12
17	Application of random amplified polymorphism DNA and 16S-23S rDNA intergenic spacer polymerase chain reaction-restriction fragment length polymorphism to predict major Streptococcus suis clonal complexes isolated from humans and pigs. Molecular and Cellular Probes, 2019, 43, 34-39.	2.1	2
18	Detection of plasmid-mediated colistin-resistant and carbapenem-resistant genes by multiplex PCR. MethodsX. 2018. 5. 532-536.	1.6	28

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19	Emergence of Streptococcus suis serotype 9 infection in humans. Journal of Microbiology, Immunology and Infection, 2017, 50, 545-546.	3.1	62
20	Development of a multiplex PCR for identification of β-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
21	Multiplex PCR for identification of six clinically relevant streptococci. Journal of Medical Microbiology, 2017, 66, 1590-1595.	1.8	10
22	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
23	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
24	Streptococcus suis serotyping by a new multiplex PCR. Journal of Medical Microbiology, 2014, 63, 824-830.	1.8	75