Korakrit Imwattana

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genetically related <i>Clostridium difficile</i> from water sources and human <scp>CDI</scp> cases revealed by wholeâ€genome sequencing. Environmental Microbiology, 2022, 24, 1221-1230.	3.8	7
2	Antimicrobial-resistant Bacteroides fragilis in Thailand and their inhibitory effect inÂvitro on the growth of Clostridioides difficile. Anaerobe, 2022, 73, 102505.	2.1	1
3	Whole-genome sequencing links Clostridium (Clostridioides) difficile in a single hospital to diverse environmental sources in the community. Journal of Applied Microbiology, 2022, 133, 1156-1168.	3.1	13
4	Esculin hydrolysis negative and TcdA-only producing strains of <i>Clostridium (Clostridioides) difficile</i> from the environment in Western Australia. Journal of Applied Microbiology, 2022, 133, 1183-1196.	3.1	5
5	Clobal evolutionary dynamics and resistome analysis of Clostridioides difficile ribotype 017. Microbial Genomics, 2022, 8, .	2.0	4
6	Can sequencing improve the diagnosis and management of Clostridioides difficile infection?. Expert Review of Molecular Diagnostics, 2021, 21, 429-431.	3.1	1
7	Molecular Characterization of, and Antimicrobial Resistance in, <i>Clostridioides difficile</i> from Thailand, 2017–2018. Microbial Drug Resistance, 2021, 27, 1505-1512.	2.0	13
8	Major genetic discontinuity and novel toxigenic species in Clostridioides difficile taxonomy. ELife, 2021, 10, .	6.0	50
9	A species-wide genetic atlas of antimicrobial resistance in Clostridioides difficile. Microbial Genomics, 2021, 7, .	2.0	8
10	Antimicrobial resistance in <i>Clostridium difficile</i> ribotype 017. Expert Review of Anti-Infective Therapy, 2020, 18, 17-25.	4.4	28
11	Genomic basis of antimicrobial resistance in non-toxigenic Clostridium difficile in Southeast Asia. Anaerobe, 2020, 66, 102290.	2.1	16
12	Mild or Malign: Clinical Characteristics and Outcomes of Clostridium difficile Infection in Thailand. Journal of Clinical Microbiology, 2020, 58, .	3.9	8
13	<i>Clostridium difficile</i> ribotype 017 – characterization, evolution and epidemiology of the dominant strain in Asia. Emerging Microbes and Infections, 2019, 8, 796-807.	6.5	61
14	High prevalence and diversity of tcdA-negative and tcdB-positive, and non-toxigenic, Clostridium difficile in Thailand. Anaerobe, 2019, 57, 4-10.	2.1	19