

Yuan Wu

List of Publications by Year in descending order

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22
papers

374
citations

759233

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23
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23
times ranked

510
citing authors

#	ARTICLE	IF	CITATIONS
1	A narrative review of <i>Clostridioides difficile</i> infection in China. <i>Anaerobe</i> , 2022, , 102540.	2.1	8
2	Evaluation of the antimicrobial activity of ridinilazole and six comparators against Chinese, Japanese and South Korean strains of <i>Clostridioides difficile</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 967-972.	3.0	4
3	New ribotype <i>Clostridioides difficile</i> from ST11 group revealed higher pathogenic ability than RT078. <i>Emerging Microbes and Infections</i> , 2021, 10, 687-699.	6.5	6
4	<i>Clostridioides difficile</i> infection in the Asia-Pacific region. <i>Emerging Microbes and Infections</i> , 2020, 9, 42-52.	6.5	47
5	Response of the gut microbiota during the <i>Clostridioides difficile</i> infection in tree shrews mimics those in humans. <i>BMC Microbiology</i> , 2020, 20, 260.	3.3	5
6	Antimicrobial Susceptibilities of <i>Clostridium difficile</i> Isolates from 12 Asia-Pacific Countries in 2014 and 2015. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	3.2	26
7	The molecular characters and antibiotic resistance of <i>Clostridioides difficile</i> from economic animals in China. <i>BMC Microbiology</i> , 2020, 20, 70.	3.3	12
8	Microevolution within ST11 group <i>Clostridioides difficile</i> isolates through mobile genetic elements based on complete genome sequencing. <i>BMC Genomics</i> , 2019, 20, 796.	2.8	5
9	Antibiotic resistance of clinical isolates of <i>Clostridioides difficile</i> in China and its association with geographical regions and patient age. <i>Anaerobe</i> , 2019, 60, 102094.	2.1	15
10	Independent Microevolution Mediated by Mobile Genetic Elements of Individual <i>Clostridium difficile</i> Isolates from Clade 4 Revealed by Whole-Genome Sequencing. <i>MSystems</i> , 2019, 4, .	3.8	16
11	Polymorphism analysis of virulence-related genes among <i>Candida tropicalis</i> isolates. <i>Chinese Medical Journal</i> , 2019, 132, 446-453.	2.3	5
12	A retrospective study of community-acquired <i>Clostridium difficile</i> infection in southwest China. <i>Scientific Reports</i> , 2018, 8, 3992.	3.3	29
13	Molecular Characterization of <i>Clostridium difficile</i> Isolates in China From 2010 to 2015. <i>Frontiers in Microbiology</i> , 2018, 9, 845.	3.5	40
14	The Activities of Adhesion and Biofilm Formation by <i>Candida tropicalis</i> Clinical Isolates Display Significant Correlation with Its Multilocus Sequence Typing. <i>Mycopathologia</i> , 2017, 182, 459-469.	3.1	14
15	<i>Clostridium difficile</i> RT 078/ST11: A Threat to Community Population and Pigs Identified in Elder Hospitalized Patients in Beijing, China. <i>Infection Control and Hospital Epidemiology</i> , 2017, 38, 1383-1385.	1.8	9
16	Distinct Expression Levels of ALS, LIP, and SAP Genes in <i>Candida tropicalis</i> with Diverse Virulent Activities. <i>Frontiers in Microbiology</i> , 2016, 7, 1175.	3.5	25
17	A Genome-Wide Transcriptional Analysis of Yeast-Hyphal Transition in <i>Candida tropicalis</i> by RNA-Seq. <i>PLoS ONE</i> , 2016, 11, e0166645.	2.5	12
18	Multilocus microsatellite markers for molecular typing of <i>Candida tropicalis</i> isolates. <i>BMC Microbiology</i> , 2014, 14, 245.	3.3	33

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19	<i>Pichia fabianii</i> blood infection in a premature infant in China: case report. BMC Research Notes, 2013, 6, 77.	1.4	23
20	Identification and Molecular Analysis of Pathogenic Yeasts in Droppings of Domestic Pigeons in Beijing, China. Mycopathologia, 2012, 174, 203-214.	3.1	18
21	Analysis of the Clonality of <i>Candida tropicalis</i> Strains from a General Hospital in Beijing Using Multilocus Sequence Typing. PLoS ONE, 2012, 7, e47767.	2.5	15
22	Confocal Raman microspectroscopy combined with chemometrics as a discrimination method of clostridia and serotypes of <i>Clostridium botulinum</i> strains. Journal of Raman Spectroscopy, 0, , .	2.5	2