Melissa Chun-Jiao Liu

List of Publications by Year in descending order

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1478505 1281871 14 151 11 6 citations h-index g-index papers 15 15 15 217 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	IncX3 Epidemic Plasmid Carrying bla NDM-5 in Escherichia coli from Swine in Multiple Geographic Areas in China. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	51
2	Prevalence and characterization of hybrid blaCTX-M among Escherichia coli isolates from livestock and other animals. Diagnostic Microbiology and Infectious Disease, 2015, 82, 148-153.	1.8	24
3	Antimicrobial susceptibility of Bacteroides fragilis group organisms in Hong Kong by the tentative EUCAST disc diffusion method. Anaerobe, 2017, 47, 51-56.	2.1	17
4	Re-sensitization of <i>mcr</i> carrying multidrug resistant bacteria to colistin by silver. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2119417119.	7.1	15
5	Evaluation of disc diffusion tests and agar screening for predicting mecA-mediated oxacillin resistance in Staphylococcus lugdunensis revealed a cefoxitin-susceptible, mecA-positive S. lugdunensis clonal complex 27 clone. Journal of Global Antimicrobial Resistance, 2020, 20, 260-265.	2.2	8
6	Large-Scale Genomic Epidemiology of Klebsiella pneumoniae Identified Clone Divergence with Hypervirulent Plus Antimicrobial-Resistant Characteristics Causing Within-Ward Strain Transmissions. Microbiology Spectrum, 2022, 10, e0269821.	3.0	7
7	Emergence of <i>ileS2</i> -Carrying, Multidrug-Resistant Plasmids in Staphylococcus lugdunensis. Antimicrobial Agents and Chemotherapy, 2016, 60, 6411-6414.	3.2	6
8	Genomic investigation of a sequence type 67 Clostridium difficile causing community-acquired fulminant colitis in Hong Kong. International Journal of Medical Microbiology, 2019, 309, 270-273.	3.6	6
9	Structures of SCCmec elements in methicillin-resistant Staphylococcus lugdunensis are closely related to those harboured by community-associated methicillin-resistant Staphylococcus aureus. Journal of Medical Microbiology, 2019, 68, 1367-1372.	1.8	6
10	OMMA enables population-scale analysis of complex genomic features and phylogenomic relationships from nanochannel-based optical maps. GigaScience, 2019, 8, .	6.4	5
11	Diversity of genomic clusters and CfiA/cfiA alleles in Bacteroides fragilis isolates from human and animals. Anaerobe, 2022, 75, 102567.	2.1	5
12	Within-Ward Transmissions of <i>Klebsiella pneumoniae</i> Low-Endemic-Risk Clones Resolved by a Genomic Analysis: Highlighting the Role of Whole-Genome Sequencing in Infection Surveillance. SSRN Electronic Journal, 0, , .	0.4	1
13	Improved Detection of mecA-Mediated \hat{l}^2 -Lactam Resistance in Staphylococcus lugdunensis Using a New Oxacillin Salt Agar Screen. Frontiers in Microbiology, 2021, 12, 704552.	3.5	O
14	Carbapenemase-producing Enterobacteriaceae (CPE) isolated from pigs in China. Access Microbiology, $2019, 1, \dots$	0.5	0