Hermine V Mkrtchyan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1119981/publications.pdf

Version: 2024-02-01

		1163117	1372567	
10	268	8	10	
papers	citations	h-index	g-index	
10	1.0	1.0	200	
10	10	10	326	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	The prevalence, antibiotic resistance and mecA characterization of coagulase negative staphylococci recovered from non-healthcare settings in London, UK. Antimicrobial Resistance and Infection Control, 2018, 7, 73.	4.1	66
2	Could Public Restrooms Be an Environment for Bacterial Resistomes?. PLoS ONE, 2013, 8, e54223.	2.5	46
3	Antibiotic resistance and mecA characterization of coagulase-negative staphylococci isolated from three hotels in London, UK. Frontiers in Microbiology, 2015, 6, 947.	3.5	44
4	Surveillance and prevalence of antimicrobial resistant bacteria from public settings within urban built environments: Challenges and opportunities for hygiene and infection control. Environment International, 2021, 157, 106836.	10.0	28
5	Whole genome sequencing revealed new molecular characteristics in multidrug resistant staphylococci recovered from high frequency touched surfaces in London. Scientific Reports, 2019, 9, 9637.	3.3	26
6	Whole Genome Sequence and Comparative Genomics Analysis of Multi-drug Resistant Environmental <i>Staphylococcus epidermidis</i> ST59. G3: Genes, Genomes, Genetics, 2018, 8, 2225-2230.	1.8	21
7	Antibiotic resistance and molecular characteristics of methicillin-resistant Staphylococcus epidermidis recovered from hospital personnel in China. Journal of Global Antimicrobial Resistance, 2020, 22, 195-201.	2.2	16
8	Diversity of SCCmec elements in Staphylococci isolated from public washrooms. BMC Microbiology, 2015, 15, 120.	3.3	11
9	Comparative Proteomic Profiling of Methicillinâ€Susceptible and Resistant <i>Staphylococcus aureus</i> . Proteomics, 2020, 20, e1900221.	2.2	6
10	Comparative Genomics Analysis Demonstrated a Link Between Staphylococci Isolated From Different Sources: A Possible Public Health Risk. Frontiers in Microbiology, 2021, 12, 576696.	3.5	4