

Multidrug-Resistant Bacteria in the Community

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Antibacterial activity of <i>Hibiscus sabdariffa</i> L. calyces against hospital isolates of multidrug resistant <i>Acinetobacter baumannii</i> . <i>Journal of Acute Disease</i> , 2016, 5, 512-516.	0.0	46
2	Risk Factors for Drug-Resistant Cap in Immunocompetent Patients. <i>Current Infectious Disease Reports</i> , 2017, 19, 11.	1.3	4
3	The endless increase of antibiotic resistance in Enterobacteriaceae and the activity of new compounds to face the challenge. <i>Enfermedades Infecciosas Y Microbiología Clínica (English Ed)</i> , 2017, 35, 475-477.	0.2	0
4	The endless increase of antibiotic resistance in Enterobacteriaceae and the activity of new compounds to face the challenge. <i>Enfermedades Infecciosas Y Microbiología Clínica</i> , 2017, 35, 477-479.	0.3	1
5	Immune stealth-driven O2 serotype prevalence and potential for therapeutic antibodies against multidrug resistant <i>Klebsiella pneumoniae</i> . <i>Nature Communications</i> , 2017, 8, 1991.	5.8	70
6	Emergence of Multidrug-Resistant Bacteria in Freshwater Ecosystems (River) and Screening of Natural Therapeutics Against the Probable Drug Targets of Drug-Resistant Pathogens by Computational Biology Approaches. , 2017, , 109-132.		0
7	Antibody-Based Immunotherapy To Treat and Prevent Infection with Hypervirulent <i>Klebsiella pneumoniae</i> . <i>Vaccine Journal</i> , 2017, 24, .	3.2	66
8	The global epidemiology of carbapenemase-producing Enterobacteriaceae. <i>Virulence</i> , 2017, 8, 460-469.	1.8	613
9	Metagenomics: The Next Culture-Independent Game Changer. <i>Frontiers in Microbiology</i> , 2017, 8, 1069.	1.5	230
10	Charged Metallopolymer-Grafted Silica Nanoparticles for Antimicrobial Applications. <i>Biomacromolecules</i> , 2018, 19, 417-425.	2.6	34
11	Complete genome sequence of <i>Enterobacter cloacae</i> R11 reveals multiple genes potentially associated with high-level polymyxin E resistance. <i>Canadian Journal of Microbiology</i> , 2018, 64, 87-90.	0.8	4
12	The Infinity War: How to Cope with Carbapenem-resistant <i>Enterobacteriaceae</i> . <i>Journal of Korean Medical Science</i> , 2018, 33, e255.	1.1	13
13	Investigation of antimicrobial susceptibility, class I and II integrons among <i>Pseudomonas aeruginosa</i> isolates from hospitalized patients in Isfahan, Iran. <i>BMC Research Notes</i> , 2018, 11, 806.	0.6	29
14	Discovery of Potential Plant-Derived Peptide Deformylase (PDF) Inhibitors for Multidrug-Resistant Bacteria Using Computational Studies. <i>Journal of Clinical Medicine</i> , 2018, 7, 563.	1.0	21
15	Infected Pancreatic Necrosis due to Multidrug-Resistant Organisms and Persistent Organ failure Predict Mortality in Acute Pancreatitis. <i>Clinical and Translational Gastroenterology</i> , 2018, 9, e190.	1.3	50
16	Incisional Negative Pressure Wound Therapy for Surgical Site Infection Prophylaxis in the Post-Antibiotic Era. <i>Surgical Infections</i> , 2018, 19, 821-830.	0.7	5
17	Purification, characterization, mode of action, and enhanced production of Salivaricin mmaye1, a novel bacteriocin from <i>Lactobacillus salivarius</i> SPW1 of human gut origin. <i>Electronic Journal of Biotechnology</i> , 2018, 35, 39-47.	1.2	16
18	Interplay between Peptidoglycan Biology and Virulence in Gram-Negative Pathogens. <i>Microbiology and Molecular Biology Reviews</i> , 2018, 82, .	2.9	36

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19	Community acquired multi drug resistant (MDR) <i>Acinetobacter baumannii</i> pneumonia in Malaysia – A case report. <i>Respiratory Medicine Case Reports</i> , 2018, 24, 147-149.	0.2	6
20	Multiscale Simulations of Clavulanate Inhibition Identify the Reactive Complex in Class A β -Lactamases and Predict the Efficiency of Inhibition. <i>Biochemistry</i> , 2018, 57, 3560-3563.	1.2	17
21	A Far-Red Fluorescent DNA Binder for Interaction Studies of Live Multidrug-Resistant Pathogens and Host Cells. <i>Angewandte Chemie</i> , 2018, 130, 11738-11742.	1.6	5
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25	Factors associated with community-onset multidrug-resistant organisms in inner Brazil. <i>American Journal of Infection Control</i> , 2018, 46, 1423-1424.	1.1	0
26	Culture-independent analysis of liver abscess using nanopore sequencing. <i>PLoS ONE</i> , 2018, 13, e0190853.	1.1	22
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