

Roshan D'Souza

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Cross-Genus "Boot-Up" of Synthetic Bacteriophage in <i>Staphylococcus aureus</i> by Using a New and Efficient DNA Transformation Method. <i>Applied and Environmental Microbiology</i> , 2022, 88, AEM0148621.	3.1	6
2	Imipenem/Relebactam Resistance in Clinical Isolates of Extensively Drug Resistant <i>Pseudomonas aeruginosa</i> : Inhibitor-Resistant β -Lactamases and Their Increasing Importance. <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, e0179021.	3.2	8
3	Proof of the triple prerequisite conditions which are essential for carbapenem resistance development in <i>Klebsiella pneumoniae</i> by using radiation-mediated mutagenesis. <i>FEMS Microbiology Letters</i> , 2021, 368, .	1.8	0
4	In Vitro Activity of a Novel Siderophore-Cephalosporin LCB10-0200 (GT-1), and LCB10-0200/Avibactam, against Carbapenem-Resistant <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> , <i>Acinetobacter baumannii</i> , and <i>Pseudomonas aeruginosa</i> Strains at a Tertiary Hospital in Korea. <i>Pharmaceuticals</i> , 2021, 14, 370.	3.8	5
5	Adjustment of Modified Carbapenem Inactivation Method Conditions for Rapid Detection of Carbapenemase-Producing <i>Acinetobacter baumannii</i> . <i>Annals of Laboratory Medicine</i> , 2020, 40, 21-26.	2.5	7
6	Complete Genome Sequence of Broad-Host-Range <i>Staphylococcus aureus</i> Myophage ESa1. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	1
7	Complete Genome Sequence of <i>Staphylococcus aureus</i> Phage SA75, Isolated from Goat Feces. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	1
8	In Vitro Activity of a Novel Siderophore-Cephalosporin, GT-1 and Serine-Type β -Lactamase Inhibitor, GT-055, against <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> and <i>Acinetobacter</i> spp. Panel Strains. <i>Antibiotics</i> , 2020, 9, 267.	3.7	17
9	Resistome Profiles, Plasmid Typing, and Whole-Genome Phylogenetic Tree Analyses of Bla _{NDM-9} and Mcr-1 Co-Harboring <i>Escherichia coli</i> ST617 from a Patient without a History of Farm Exposure in Korea. <i>Pathogens</i> , 2019, 8, 212.	2.8	7
10	Phenotypic and Genotypic Characterization of <i>Acinetobacter</i> spp. Panel Strains: A Cornerstone to Facilitate Antimicrobial Development. <i>Frontiers in Microbiology</i> , 2019, 10, 559.	3.5	15
11	First Report of the Carbapenemase Gene bla _{OXA-499} in <i>Acinetobacter pittii</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	5
12	Whole genome and transcriptome analysis reveal MALDI-TOF MS and SDS-PAGE have limited performance for the detection of the key outer membrane protein in carbapenem-resistant <i>Klebsiella pneumoniae</i> isolates. <i>Oncotarget</i> , 2017, 8, 84818-84826.	1.8	4
13	Panel strain of <i>Klebsiella pneumoniae</i> for beta-lactam antibiotic evaluation: their phenotypic and genotypic characterization. <i>PeerJ</i> , 2017, 5, e2896.	2.0	23
14	Molecular epidemiology and resistome analysis of multidrug-resistant ST11 <i>Klebsiella pneumoniae</i> strain containing multiple copies of extended-spectrum β -lactamase genes using whole-genome sequencing. <i>New Microbiologica</i> , 2017, 40, 38-44.	0.1	11
15	Prediction of Putative Resistance Islands in a Carbapenem-Resistant <i>Acinetobacter baumannii</i> Global Clone 2 Clinical Isolate. <i>Annals of Laboratory Medicine</i> , 2016, 36, 320-324.	2.5	12
16	Insufficient Discriminatory Power of Matrix-Assisted Laser Desorption Ionization Time-of-Flight Mass Spectrometry Dendrograms to Determine the Clonality of Multi-Drug-Resistant <i>Acinetobacter baumannii</i> Isolates from an Intensive Care Unit. <i>BioMed Research International</i> , 2015, 2015, 1-8.	1.9	18
17	Complete genome sequence of the siphoviral bacteriophage ϕ R3177, which lyses an OXA-66-producing carbapenem-resistant <i>Acinetobacter baumannii</i> isolate. <i>Archives of Virology</i> , 2015, 160, 3157-3160.	2.1	6