

Ilana L Camargo

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

38
papers

602
citations

13
h-index

23
g-index

39
ext. papers

719
ext. citations

3.8
avg, IF

3.55
L-index

#	Paper	IF	Citations
38	Polyphasic characterization and antimicrobial properties of <i>Induratia</i> species isolated from <i>Coffea arabica</i> in Brazil. <i>Mycological Progress</i> , 2021 , 20, 1457-1477	1.9	2
37	Polymyxin Resistance Among XDR ST1 Carbapenem-Resistant Clone Expanding in a Teaching Hospital. <i>Frontiers in Microbiology</i> , 2021 , 12, 622704	5.7	8
36	Liquid Fungal Cocultivation as a Strategy to Access Bioactive Metabolites. <i>Planta Medica</i> , 2021 , 87, 187-195	3.7	1
35	Discovery of a Novel Lineage ST 1870 Endophytically Isolated from Medicinal Which Shows Potent Antileishmanial and Antimicrobial Effects. <i>International Journal of Microbiology</i> , 2021 , 2021, 6618559	3.6	0
34	Study of the Antibacterial Property of Tea Tree Oil and Its Incorporation Into Poly(Lactic Acid)-montmorillonite Clay Bionanocomposites. <i>Macromolecular Symposia</i> , 2020 , 394, 2000073	0.8	1
33	PVA/anionic collagen membranes as drug carriers of ciprofloxacin hydrochloride with sustained antibacterial activity and potential use in the treatment of ulcerative keratitis. <i>Journal of Biomaterials Applications</i> , 2020 , 35, 301-312	2.9	8
32	A mutation in the glycosyltransferase gene <i>lafB</i> causes daptomycin hypersusceptibility in <i>Enterococcus faecium</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2020 , 75, 36-45	5.1	6
31	Oxidative functionalization of a halimane diterpenoid achieved by fungal transformation. <i>Bioorganic Chemistry</i> , 2019 , 86, 550-556	5.1	2
30	Dissemination of <i>bla</i> in an NTE by an <i>IncX5</i> plasmid. <i>Plasmid</i> , 2019 , 106, 102446	3.3	4
29	Molecular basis for the emergence of a new hospital endemic tigecycline-resistant <i>Enterococcus faecalis</i> ST103 lineage. <i>Infection, Genetics and Evolution</i> , 2019 , 67, 23-32	4.5	5
28	Resistance in In Vitro Selected Tigecycline-Resistant Methicillin-Resistant <i>Staphylococcus aureus</i> Sequence Type 5 Is Driven by Mutations in <i>mepR</i> and <i>mepA</i> Genes. <i>Microbial Drug Resistance</i> , 2018 , 24, 519-526	2.9	17
27	Daptomycin and vancomycin heteroresistance revealed among CC5-SCCmecII MRSA clone and in vitro evaluation of treatment alternatives. <i>Journal of Global Antimicrobial Resistance</i> , 2018 , 14, 209-216	3.4	7
26	Antibacterial Activity of the Non-Cytotoxic Peptide (p-BthTX-I) and Its Serum Degradation Product against Multidrug-Resistant Bacteria. <i>Molecules</i> , 2017 , 22,	4.8	11
25	High-Quality Draft Genome Sequence of the Multidrug-Resistant Clinical Isolate <i>Enterococcus faecium</i> VRE16. <i>Genome Announcements</i> , 2016 , 4,		2
24	C-terminal Lysine-Linked Magainin 2 with Increased Activity Against Multidrug-Resistant Bacteria. <i>Protein and Peptide Letters</i> , 2016 , 23, 738-47	1.9	12
23	The Amazing World of Peptide Engineering: the Example of Antimicrobial Peptides from Frogs and Their Analogues. <i>Protein and Peptide Letters</i> , 2016 , 23, 722-37	1.9	10
22	Molecular analysis of methicillin-resistant <i>Staphylococcus aureus</i> dissemination among healthcare professionals and/or HIV patients from a tertiary hospital. <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2016 , 49, 51-6	1.5	5

21	Different VanA Elements in <i>E. faecalis</i> and in <i>E. faecium</i> Suggest at Least Two Origins of Tn1546 Among VRE in a Brazilian Hospital. <i>Microbial Drug Resistance</i> , 2015 , 21, 320-8	2.9	9
20	Molecular characterization of methicillin-resistant <i>Staphylococcus aureus</i> resistant to tigecycline and daptomycin isolated in a hospital in Brazil. <i>Epidemiology and Infection</i> , 2014 , 142, 479-83	4.3	17
19	Clonal complexes of <i>Staphylococcus aureus</i> : all mixed and together. <i>FEMS Microbiology Letters</i> , 2014 , 351, 7-8	2.9	7
18	Draft Genome Sequence of Methicillin-Resistant <i>Staphylococcus aureus</i> Strain SA16, Representative of an Endemic Clone from a Brazilian Hospital. <i>Genome Announcements</i> , 2013 , 1,		5
17	The first report of the qnrB19, qnrS1 and aac(61)-Ib-cr genes in urinary isolates of ciprofloxacin-resistant <i>Escherichia coli</i> in Brazil. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2012 , 107, 687-9	2.6	11
16	An RpoB mutation confers dual heteroresistance to daptomycin and vancomycin in <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2010 , 54, 5222-33	5.9	132
15	Inactivation of formyltransferase (wbkC) gene generates a <i>Brucella abortus</i> rough strain that is attenuated in macrophages and in mice. <i>Vaccine</i> , 2010 , 28, 5627-34	4.1	23
14	Clonal relationships determined by multilocus sequence typing among enteropathogenic <i>Escherichia coli</i> isolated in Brazil. <i>Canadian Journal of Microbiology</i> , 2009 , 55, 672-9	3.2	10
13	<i>Staphylococcus aureus</i> --probing for host weakness?. <i>Journal of Bacteriology</i> , 2008 , 190, 2253-6	3.5	22
12	Serial daptomycin selection generates daptomycin-nonsusceptible <i>Staphylococcus aureus</i> strains with a heterogeneous vancomycin-intermediate phenotype. <i>Antimicrobial Agents and Chemotherapy</i> , 2008 , 52, 4289-99	5.9	93
11	Virulence factors in vancomycin-resistant and vancomycin - susceptible <i>Enterococcus faecalis</i> from Brazil. <i>Brazilian Journal of Microbiology</i> , 2008 , 39, 273-278	2.2	4
10	Multilocus sequence typing of uropathogenic ESBL-producing <i>Escherichia coli</i> isolated in a Brazilian community. <i>Current Microbiology</i> , 2007 , 55, 524-9	2.4	16
9	The role of the vacB gene in the pathogenesis of <i>Brucella abortus</i> . <i>Microbes and Infection</i> , 2007 , 9, 375-80	3.3	16
8	Molecular characterization and T and B cell epitopes prediction of <i>Mycoplasma synoviae</i> 53 strain VlhA hemagglutinin. <i>Genetics and Molecular Biology</i> , 2007 , 30, 264-269	2	1
7	Sequence analysis of <i>Enterococcus faecium</i> strain 10/96A (VanD4), the original vancomycin-resistant <i>E. faecium</i> strain in Brazil. <i>Journal of Clinical Microbiology</i> , 2006 , 44, 2635-7	9.7	6
6	Multilocus sequence typing and analysis of putative virulence factors in vancomycin-resistant and vancomycin-sensitive <i>Enterococcus faecium</i> isolates from Brazil. <i>Clinical Microbiology and Infection</i> , 2006 , 12, 1123-30	9.5	53
5	Evaluation of clonality in enterococci isolated in Brazil carrying Tn1546-like elements associated with vanA plasmids. <i>FEMS Microbiology Letters</i> , 2006 , 258, 29-36	2.9	16
4	Occurrence of insertion sequences within the genomes and Tn1546-like elements of glycopeptide-resistant enterococci isolated in Brazil, and identification of a novel element, ISEfa5. <i>International Journal of Medical Microbiology</i> , 2005 , 294, 513-9	3.7	18

3	Enterococcus gallinarum carrying the vanA gene cluster: first report in Brazil. <i>Brazilian Journal of Medical and Biological Research</i> , 2004 , 37, 1669-71	2.8	18
2	Identification of an unusual VanA element in glycopeptide-resistant Enterococcus faecium in Brazil following international transfer of a bone marrow transplant patient. <i>Canadian Journal of Microbiology</i> , 2004 , 50, 767-70	3.2	16
1	Effect of C-terminal and N-terminal dimerization and alanine scanning on antibacterial activity of the analogs of the peptide p-BthTX-I. <i>Peptide Science</i> , e24243	3	1