Rosineide Marques Ribas

List of Publications by Year
in descending order

53	789	471509 17	552781 26
papers	citations	h-index	g-index

54	54	54	1289
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	WHO Critical Priority Escherichia coli as One Health Challenge for a Post-Pandemic Scenario: Genomic Surveillance and Analysis of Current Trends in Brazil. Microbiology Spectrum, 2022, 10, e0125621.	3.0	31
2	Novel ST1465/CC216 Nosocomial Lineage of Carbapenem-Resistant Acinetobacter baumannii Harboring an Unusual Plasmid Carrying bla_{NDM-1} Gene. Microbial Drug Resistance, 2021, 27, 471-475.	2.0	6
3	Gram-negative bacilli bacteremia: a 7 year retrospective study in a referral Brazilian tertiary-care teaching hospital. Journal of Medical Microbiology, 2021, 70, .	1.8	2
4	Public health in Brazil: Before COVID-19, and after. Travel Medicine and Infectious Disease, 2021, 40, 101974.	3.0	0
5	Dengue, Influenza and COVID-19 in Brazil: The â€œPerfect Stormâ€• Coronaviruses, 2021, 2, 4-5.	0.3	0
6	Genetic Alterations Associated with Polymyxin B Resistance in Nosocomial KPC-2-Producing Klebsiella pneumoniae from Brazil. Microbial Drug Resistance, 2021, 27, 1677-1684.	2.0	3
7	Financial impact of healthcare-associated infections on intensive care units estimated for fifty Brazilian university hospitals affiliated to the unified health system. Journal of Hospital Infection, 2021, 117, 96-102.	2.9	3
8	Health care-associated infections: Significant challenge and it's to-be. American Journal of Infection Control, 2021, 49, 1212-1213.	2.3	0
9	Novel small IncX3 plasmid carrying the blaKPC-2 gene in high-risk Klebsiella pneumoniae ST11/CG258. Diagnostic Microbiology and Infectious Disease, 2020, 96, 114900.	1.8	8
10	Costs of healthcare-associated infections to the Brazilian public Unified Health System in a tertiary-care teaching hospital: a matched caseâ€“control study. Journal of Hospital Infection, 2020, 106, 303-310.	2.9	11
11	The rising problem of hospital antimicrobial resistance and the challenges of antibiotic prescription in Brazil. Journal of Chemotherapy, 2020, 33, 1-3.	1.5	1
12	Implications of social distancing in Brazil in the COVID-19 pandemic. Infection Control and Hospital Epidemiology, 2020, , 1-2.	1.8	12
13	Coronavirus Disease 2019 (COVID-19) and healthcare-associated infections: Emerging and future challenges for public health in Brazil. Travel Medicine and Infectious Disease, 2020, 37, 101675.	3.0	12
14	High mortality by nosocomial infections caused by carbapenem-resistant P. aeruginosa in a referral hospital in Brazil: facing the perfect storm. Journal of Medical Microbiology, 2020, 69, 1388-1397.	1.8	5
15	2021 olympic games Tokyo: Safety issues and protection against COVID-19 transmission. Journal of Global Infectious Diseases, 2020, 12, 114.	0.5	2
16	Infections and antimicrobial resistance in an adult intensive care unit in a Brazilian hospital and the influence of drug resistance on the thirty-day mortality among patients with bloodstream infections. Revista Da Sociedade Brasileira De Medicina Tropical, 2020, 53, e20190106.	0.9	4
17	Origin of Catheter-Related Bloodstream Infections Caused by Staphylococcus epidermidis in Critical Neonates. Journal of Child Science, 2020, 10, e196-e201.	0.2	0
18	Incidence of infections caused by carbapenem-resistant Acinetobacter baumannii. American Journal of Infection Control, 2019, 47, 1431-1435.	2.3	17

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19	Early Dissemination of IncQ1 Plasmids in KPC-2-Producing Klebsiella pneumoniae CG258. Microbial Drug Resistance, 2019, 25, 1257-1259.	2.0	5
20	Genomic features of a clinical ESBL-producing and colistin-resistant hypermucoviscous K. quasipneumoniae subsp. similipneumoniae from Brazil. Brazilian Journal of Infectious Diseases, 2019, 23, 207-209.	0.6	4
21	Molecular Detection of Class 1 Integron-Associated Gene Cassettes in KPC-2-Producing Klebsiella pneumoniae Clones by Whole-Genome Sequencing. Microbial Drug Resistance, 2019, 25, 1127-1131.	2.0	8
22	Using point prevalence survey to define burden of antimicrobial use among 35 adult intensive care units in Brazil. Infectious Diseases, 2019, 51, 459-462.	2.8	4
23	Small IncQ1 and Col-Like Plasmids Harboring bla _{KPC-2} and Non-Tn 4401 Elements (NTE _{KPC} -IId) in High-Risk Lineages of Klebsiella pneumoniae CG258. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	27
24	Detection of ISE cp1- associated bla CTX-M-15 â€“mediated resistance to colistin in KPC-producing Klebsiella pneumoniae isolates. International Journal of Antimicrobial Agents, 2018, 51, 810-811.	2.5	4
25	Insights into a novel Tn4401 deletion (Tn4401i) in a multidrug-resistant Klebsiella pneumoniae clinical strain belonging to the high-risk clonal group 258 producing KPC-2. International Journal of Antimicrobial Agents, 2018, 52, 525-527.	2.5	9
26	Association of Colistin-Resistant KPC Clonal Strains with Subsequent Infections and Colonization and Biofilm Production. Microbial Drug Resistance, 2018, 24, 1441-1449.	2.0	6
27	Molecular characterization and clonal dynamics of nosocomial blaOXA-23 producing XDR Acinetobacter baumannii. PLoS ONE, 2018, 13, e0198643.	2.5	23
28	Hypervirulence and biofilm production in KPC-2-producing Klebsiella pneumoniae CG258 isolated in Brazil. Journal of Medical Microbiology, 2018, 67, 523-528.	1.8	27
29	Pressure ulcer as a reservoir of multiresistant Gram-negative bacilli: risk factors for colonization and development of bacteremia. Brazilian Journal of Infectious Diseases, 2017, 21, 171-175.	0.6	31
30	Carbapenem-resistant Pseudomonas aeruginosa : association with virulence genes and biofilm formation. Brazilian Journal of Microbiology, 2017, 48, 211-217.	2.0	59
31	IncX3 plasmid harboring a non-Tn 4401 genetic element (NTE KPC) in a hospital-associated clone of KPC-2-producing Klebsiella pneumoniae ST340/CG258. Diagnostic Microbiology and Infectious Disease, 2017, 89, 164-167.	1.8	24
32	Molecular epidemiological survey of bacteremia by multidrug resistant Pseudomonas aeruginosa: the relevance of intrinsic resistance mechanisms. PLoS ONE, 2017, 12, e0176774.	2.5	24
33	High frequency of the combined presence of QRDR mutations and PMQR determinants in multidrug-resistant Klebsiella pneumoniae and Escherichia coli isolates from nosocomial and community-acquired infections. Journal of Medical Microbiology, 2017, 66, 1144-1150.	1.8	18
34	Clinical and Molecular Epidemiology of Multidrug-Resistant P. aeruginosa Carrying aac(6')-Ib-cr, qnrS1 and blaSPM Genes in Brazil. PLoS ONE, 2016, 11, e0155914.	2.5	30
35	Multidrug Resistance Related to Biofilm Formation in Acinetobacter baumannii and Klebsiella pneumoniae Clinical Strains from Different Pulsotypes. Current Microbiology, 2016, 72, 617-627.	2.2	43
36	Biofilm formation of Brazilian meticillin-resistant Staphylococcus aureus strains: prevalence of biofilm determinants and clonal profiles. Journal of Medical Microbiology, 2016, 65, 286-297.	1.8	18

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37	Spread of Multidrug-resistant microrganisms: a global threat and critical healthcare problem. Revista De Epidemiologia E Controle De InfecÃ§Ã£o, 2016, 6, .	0.0	0
38	The nares as a CA-MRSA reservoir in the healthy elderly. Revista Da Sociedade Brasileira De Medicina Tropical, 2015, 48, 614-616.	0.9	6
39	Spread of multidrug-resistant Acinetobacter baumannii and Pseudomonas aeruginosa clones in patients with ventilator-associated pneumonia in an adult intensive care unit at a university hospital. Brazilian Journal of Infectious Diseases, 2015, 19, 350-357.	0.6	35
40	Late onset sepsis in newborn babies: epidemiology and effect of a bundle to prevent central line associated bloodstream infections in the neonatal intensive care unit. Brazilian Journal of Infectious Diseases, 2015, 19, 52-57.	0.6	33
41	Molecular epidemiological survey of the quinolone- and carbapenem-resistant genotype and its association with the type III secretion system in Pseudomonas aeruginosa. Journal of Medical Microbiology, 2015, 64, 262-271.	1.8	8
42	Genotypic study documents divergence in the pathogenesis of bloodstream infection related central venous catheters in neonates. Brazilian Journal of Infectious Diseases, 2014, 18, 387-393.	0.6	6
43	A sustained endemic outbreak of vancomycin-resistant Enterococcus faecium: A 30-month surveillance study. Scandinavian Journal of Infectious Diseases, 2014, 46, 547-554.	1.5	8
44	Pseudomonas aeruginosa bacteraemia: independent risk factors for mortality and impact of resistance on outcome. Journal of Medical Microbiology, 2014, 63, 1679-1687.	1.8	78
45	Authors' reply: Emergence of antibiotic-resistant bacterial strains, methicillin-resistant Staphylococcus aureus and extended spectrum Î²-lactamases, and multi-drug resistance are problems similar to global warming. Revista Da Sociedade Brasileira De Medicina Tropical, 2014, 47, 817-818.	0.9	2
46	Relationship between nasal colonization and ventilator-associated pneumonia and the role of the environment in transmission of Staphylococcus aureus inÂintensive care units. American Journal of Infection Control, 2013, 41, 1236-1240.	2.3	25
47	Active surveillance to determine the impact of methicillin resistance on mortality in patients with bacteremia and influences of the use of antibiotics on the development of MRSA infection. Revista Da Sociedade Brasileira De Medicina Tropical, 2013, 46, 713-718.	0.9	17
48	Nosocomial infections in a pediatric intensive care unit of a developing country: NHSN surveillance. Revista Da Sociedade Brasileira De Medicina Tropical, 2012, 45, 475-479.	0.9	29
49	Risk factors for vancomycin-resistant enterococci colonisation in critically ill patients. Memorias Do Instituto Oswaldo Cruz, 2012, 107, 57-63.	1.6	36
50	Nosocomial bloodstream infections: organisms, risk factors and resistant phenotypes in the Brazilian University Hospital. Brazilian Journal of Infectious Diseases, 2007, 11, 351-354.	0.6	8
51	Vancomycin-resistant van a phenotype Enterococcus faecalis: first case in Minas Gerais state and epidemiological considerations. Brazilian Journal of Infectious Diseases, 2007, 11, 439-40.	0.6	5
52	Infection and colonization by Gram-negative bacilli in neonates hospitalized in High Risk Nursery at Uberlandia Federal University Hospital: etiology, resistant phenotypes and risk factors. Brazilian Journal of Microbiology, 2004, 35, 193-198.	2.0	3
53	Conventional versus molecular tests (Multiplex PCR and PCR mecA gene) for detection of methicillin resistant Staphylococcus aureus. Brazilian Journal of Microbiology, 0, 34, 35-37.	2.0	2