

Ana R Freitas

List of Publications by Year in descending order

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Version: 2024-02-01

54
papers

1,841
citations

293460

24
h-index

325983

40
g-index

59
all docs

59
docs citations

59
times ranked

2065
citing authors

#	ARTICLE	IF	CITATIONS
1	High-Resolution Genotyping Unveils Identical Ampicillin-Resistant <i>Enterococcus faecium</i> Strains in Different Sources and Countries: A One Health Approach. <i>Microorganisms</i> , 2022, 10, 632.	1.6	6
2	Evolution of Chlorhexidine Susceptibility and of the EfrEF Operon among <i>Enterococcus faecalis</i> from Diverse Environments, Clones, and Time Spans. <i>Microbiology Spectrum</i> , 2022, 10, .	1.2	0
3	Multidrug-resistant high-risk <i>Enterococcus faecium</i> clones: can we really define them?. <i>International Journal of Antimicrobial Agents</i> , 2021, 57, 106227.	1.1	24
4	MicroMundo@UPorto: an experimental microbiology project fostering student's antimicrobial resistance awareness and personal and social development. <i>FEMS Microbiology Letters</i> , 2021, 368, .	0.7	3
5	Apparent nosocomial adaptation of <i>Enterococcus faecalis</i> predates the modern hospital era. <i>Nature Communications</i> , 2021, 12, 1523.	5.8	69
6	Industrial dog food is a vehicle of multidrug-resistant enterococci carrying virulence genes often linked to human infections. <i>International Journal of Food Microbiology</i> , 2021, 358, 109284.	2.1	13
7	Fitness cost of vancomycin-resistant <i>Enterococcus faecium</i> plasmids associated with hospital infection outbreaks. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2757-2764.	1.3	6
8	Linezolid- and Multidrug-Resistant Enterococci in Raw Commercial Dog Food, Europe, 2019–2020. <i>Emerging Infectious Diseases</i> , 2021, 27, 2221-2224.	2.0	17
9	Diversity of metal and antibiotic resistance genes in <i>Enterococcus</i> spp. from the last century reflects multiple pollution and genetic exchange among phyla from overlapping ecosystems. <i>Science of the Total Environment</i> , 2021, 787, 147548.	3.9	13
10	<i>Enterococcus</i> spp. as a Producer and Target of Bacteriocins: A Double-Edged Sword in the Antimicrobial Resistance Crisis Context. <i>Antibiotics</i> , 2021, 10, 1215.	1.5	23
11	From farm to fork: identical clones and Tn6674-like elements in linezolid-resistant <i>Enterococcus faecalis</i> from food-producing animals and retail meat. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 30-35.	1.3	28
12	Comment on: Emergence of plasmid-mediated oxazolidinone resistance gene <i>poxTA</i> from CC17 <i>Enterococcus faecium</i> of pig origin. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 1358-1359.	1.3	1
13	Editorial: Antimicrobials and Anticancers of Bacterial Origins. <i>Frontiers in Microbiology</i> , 2020, 11, 842.	1.5	4
14	Linezolid-resistant (Tn6246:: <i>fexB</i> - <i>poxTA</i>) <i>Enterococcus faecium</i> strains colonizing humans and bovines on different continents: similarity without epidemiological link. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 2416-2423.	1.3	34
15	Transmission of Antibiotic Resistant Bacteria and Genes: Unveiling the Jigsaw Pieces of a One Health Problem. <i>Pathogens</i> , 2020, 9, 497.	1.2	7
16	Silent clonal spread of vancomycin-resistant <i>Enterococcus faecalis</i> ST6 and ST525 colonizing patients at hospital admission in Natal, Brazil. <i>Infection Control and Hospital Epidemiology</i> , 2020, 41, 485-487.	1.0	2
17	Comparative genomics of global <i>optrA</i> -carrying <i>Enterococcus faecalis</i> uncovers a common chromosomal hotspot for <i>optrA</i> acquisition within a diversity of core and accessory genomes. <i>Microbial Genomics</i> , 2020, 6, .	1.0	31
18	Isolation and Visualization of Plasmids from Gram-Positive Bacteria of Interest in Public Health. <i>Methods in Molecular Biology</i> , 2020, 2075, 21-38.	0.4	3

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19	Methods to Quantify DNA Transfer in Enterococcus. <i>Methods in Molecular Biology</i> , 2020, 2075, 111-122.	0.4	0
20	Dispersal of linezolid-resistant enterococci carrying <i>poxtA</i> or <i>optrA</i> in retail meat and food-producing animals from Tunisia. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 2865-2869.	1.3	65
21	2CS-CHX ^T Operon Signature of Chlorhexidine Tolerance among <i>Enterococcus faecium</i> Isolates. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	10
22	Phylogenomics of <i>Enterococcus faecalis</i> from wild birds: new insights into host-associated differences in core and accessory genomes of the species. <i>Environmental Microbiology</i> , 2019, 21, 3046-3062.	1.8	14
23	Fourier transform infrared spectroscopy: unlocking fundamentals and prospects for bacterial strain typing. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2019, 38, 427-448.	1.3	92
24	Dissemination of <i>Staphylococcus epidermidis</i> ST22 With Stable, High-Level Resistance to Linezolid and Tedizolid in the Greek-Turkish Region (2008–2016). <i>Infection Control and Hospital Epidemiology</i> , 2018, 39, 492-494.	1.0	8
25	Water supply and feed as sources of antimicrobial-resistant <i>Enterococcus</i> spp. in aquacultures of rainbow trout (<i>Oncorhynchus mykiss</i>), Portugal. <i>Science of the Total Environment</i> , 2018, 625, 1102-1112.	3.9	29
26	Distribution of putative virulence markers in <i>Enterococcus faecium</i> : towards a safety profile review. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 306-319.	1.3	40
27	High rates of colonisation by ampicillin-resistant enterococci in residents of long-term care facilities in Porto, Portugal. <i>International Journal of Antimicrobial Agents</i> , 2018, 51, 503-507.	1.1	11
28	Update on prevalence and mechanisms of resistance to linezolid, tigecycline and daptomycin in enterococci in Europe: Towards a common nomenclature. <i>Drug Resistance Updates</i> , 2018, 40, 25-39.	6.5	165
29	Wild corvid birds colonized with vancomycin-resistant <i>Enterococcus faecium</i> of human origin harbor epidemic <i>vanA</i> plasmids. <i>Environment International</i> , 2018, 118, 125-133.	4.8	13
30	Rapid detection of high-risk <i>Enterococcus faecium</i> clones by matrix-assisted laser desorption ionization time-of-flight mass spectrometry. <i>Diagnostic Microbiology and Infectious Disease</i> , 2017, 87, 299-307.	0.8	14
31	Detection of <i>optrA</i> in the African continent (Tunisia) within a mosaic <i>Enterococcus faecalis</i> plasmid from urban wastewaters. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 3245-3251.	1.3	61
32	Co-diversification of <i>Enterococcus faecium</i> Core Genomes and PBP5: Evidences of <i>pbp5</i> Horizontal Transfer. <i>Frontiers in Microbiology</i> , 2016, 7, 1581.	1.5	34
33	Co-infection with three linezolid-resistant <i>Enterococcus faecium</i> ST117 strain variants: what are we missing in diagnosis?. <i>International Journal of Antimicrobial Agents</i> , 2016, 47, 500-501.	1.1	5
34	Multilevel population genetic analysis of <i>vanA</i> and <i>vanB</i> <i>Enterococcus faecium</i> causing nosocomial outbreaks in 27 countries (1986–2012). <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 3351-3366.	1.3	129
35	Relevance of <i>trcYAZB</i> operon acquisition for <i>Enterococcus</i> survival at high copper concentrations under anaerobic conditions: Table 1.. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 560-563.	1.3	10
36	Filling the map for antimicrobial resistance in sub-Saharan Africa: ampicillin-resistant <i>Enterococcus</i> from non-clinical sources in Angola: Table 1.. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 2914-2916.	1.3	16

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37	A hospital sewage ST17 <i>Enterococcus faecium</i> with a transferable Inc18-like plasmid carrying genes coding for resistance to antibiotics and quaternary ammonium compounds (qacZ). <i>Journal of Global Antimicrobial Resistance</i> , 2015, 3, 49-51.	0.9	9
38	Co-transfer of resistance to high concentrations of copper and first-line antibiotics among <i>Enterococcus</i> from different origins (humans, animals, the environment and foods) and clonal lineages. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 899-906.	1.3	68
39	Spread of multidrug-resistant <i>Enterococcus</i> to animals and humans: an underestimated role for the pig farm environment. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 2746-2754.	1.3	74
40	Microevolutionary Events Involving Narrow Host Plasmids Influences Local Fixation of Vancomycin-Resistance in <i>Enterococcus</i> Populations. <i>PLoS ONE</i> , 2013, 8, e60589.	1.1	56
41	Clonal outbreak of ST17 multidrug-resistant <i>Enterococcus faecium</i> harbouring an Inc18-like::Tn1546 plasmid in a haemo-oncology ward of a Spanish hospital. <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 832-836.	1.3	32
42	Different Genetic Supports for the <i>tet(S)</i> Gene in <i>Enterococci</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 6014-6018.	1.4	15
43	A <i>tet(S/M)</i> hybrid from CTn6000 and CTn916 recombination. <i>Microbiology (United Kingdom)</i> , 2012, 158, 2710-2711.	0.7	8
44	Non-susceptibility to tigecycline in enterococci from hospitalised patients, food products and community sources. <i>International Journal of Antimicrobial Agents</i> , 2011, 38, 174-176.	1.1	23
45	A multiresistance megaplasmid pLG1 bearing a <i>hylEfm</i> genomic island in hospital <i>Enterococcus faecium</i> isolates. <i>International Journal of Medical Microbiology</i> , 2011, 301, 165-175.	1.5	66
46	Characterization of antibiotic resistant enterococci isolated from untreated waters for human consumption in Portugal. <i>International Journal of Food Microbiology</i> , 2011, 145, 315-319.	2.1	30
47	Human and Swine Hosts Share Vancomycin-Resistant <i>Enterococcus faecium</i> CC17 and CC5 and <i>Enterococcus faecalis</i> CC2 Clonal Clusters Harboring Tn 1546 on Indistinguishable Plasmids. <i>Journal of Clinical Microbiology</i> , 2011, 49, 925-931.	1.8	126
48	Host range of enterococcal <i>vanA</i> plasmids among Gram-positive intestinal bacteria. <i>Journal of Antimicrobial Chemotherapy</i> , 2011, 66, 273-282.	1.3	55
49	Global Spread of the <i>hylEfm</i> Colonization-Virulence Gene in Megaplasms of the <i>Enterococcus faecium</i> CC17 Polyclonal Subcluster. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 2660-2665.	1.4	67
50	Successful application of the DiversiLab repetitive-sequence-based PCR typing system for confirmation of the circulation of a multiresistant <i>Pseudomonas aeruginosa</i> clone in different hospital wards. <i>Diagnostic Microbiology and Infectious Disease</i> , 2010, 67, 202-206.	0.8	19
51	Dispersion of Multidrug-Resistant <i>Enterococcus faecium</i> Isolates Belonging to Major Clonal Complexes in Different Portuguese Settings. <i>Applied and Environmental Microbiology</i> , 2009, 75, 4904-4908.	1.4	52
52	Clonal expansion within clonal complex 2 and spread of vancomycin-resistant plasmids among different genetic lineages of <i>Enterococcus faecalis</i> from Portugal. <i>Journal of Antimicrobial Chemotherapy</i> , 2009, 63, 1104-1111.	1.3	76
53	Diversity of Tn 1546 and Its Role in the Dissemination of Vancomycin-Resistant <i>Enterococci</i> in Portugal. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 1001-1008.	1.4	64
54	Antibiotic susceptibility testing for therapy and antimicrobial resistance surveillance: Âgenotype beats phenotype?. <i>Future Microbiology</i> , 0, , .	1.0	0