PatrÃ-cia Antunes

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	High diversity of pathogenic <i>Escherichia coli</i> clones carrying <i>mcrâ€l</i> among gulls underlines the need for strategies at the environment–livestock–human interface. Environmental Microbiology, 2022, 24, 4702-4713.	1.8	4
2	Evolution of Chlorhexidine Susceptibility and of the EfrEF Operon among Enterococcus faecalis from Diverse Environments, Clones, and Time Spans. Microbiology Spectrum, 2022, 10, .	1.2	0
3	MicroMundo@UPorto: an experimental microbiology project fostering student's antimicrobial resistance awareness and personal and social development. FEMS Microbiology Letters, 2021, 368, .	0.7	3
4	From farm to fork: Colistin voluntary withdrawal in Portuguese farms reflected in decreasing occurrence of <i>mcrâ€lâ€</i> carrying <i>Enterobacteriaceae</i> from chicken meat. Environmental Microbiology, 2021, 23, 7563-7577.	1.8	15
5	Diversity of metal and antibiotic resistance genes in Enterococcus spp. from the last century reflects multiple pollution and genetic exchange among phyla from overlapping ecosystems. Science of the Total Environment, 2021, 787, 147548.	3.9	13
6	Atypical Non-H2S-Producing Monophasic Salmonella Typhimurium ST3478 Strains from Chicken Meat at Processing Stage Are Adapted to Diverse Stresses. Pathogens, 2020, 9, 701.	1.2	10
7	Food-to-Humans Bacterial Transmission. Microbiology Spectrum, 2020, 8, .	1.2	27
8	Tolerance to arsenic contaminant among multidrugâ€resistant and copperâ€tolerant <scp><i>Salmonella</i></scp> successful clones is associated with diverse <scp><i>ars</i></scp> operons and genetic contexts. Environmental Microbiology, 2020, 22, 2829-2842.	1.8	17
9	2CS-CHX ^T Operon Signature of Chlorhexidine Tolerance among Enterococcus faecium Isolates. Applied and Environmental Microbiology, 2019, 85, .	1.4	10
10	A hybrid modelling approach for eliciting health state preferences: the Portuguese EQ-5D-5L value set. Quality of Life Research, 2019, 28, 3163-3175.	1.5	52
11	Non-typhoidal Salmonella in the Pig Production Chain: A Comprehensive Analysis of Its Impact on Human Health. Pathogens, 2019, 8, 19.	1.2	92
12	Food-to-Humans Bacterial Transmission. , 2019, , 161-193.		3
13	Water supply and feed as sources of antimicrobial-resistant Enterococcus spp. in aquacultures of rainbow trout (Oncorhyncus mykiss), Portugal. Science of the Total Environment, 2018, 625, 1102-1112.	3.9	29
14	Imported poultry meat as a source of extended-spectrum cephalosporin-resistant CMY-2-producing Salmonella Heidelberg and Salmonella Minnesota in the European Union, 2014–2015. International Journal of Antimicrobial Agents, 2018, 51, 151-154.	1.1	47
15	Occurrence of mcr-1 in Escherichia coli from rabbits of intensive farming. Veterinary Microbiology, 2018, 227, 78-81.	0.8	13
16	Inflow water is a major source of trout farming contamination with Salmonella and multidrug resistant bacteria. Science of the Total Environment, 2018, 642, 1163-1171.	3.9	27
17	Discrimination of non-typhoid Salmonella serogroups and serotypes by Fourier Transform Infrared Spectroscopy: A comprehensive analysis. International Journal of Food Microbiology, 2018, 285, 34-41.	2.1	28
18	<i>mcr-1</i> in Carbapenemase-Producing <i>Klebsiella pneumoniae</i> with Hospitalized Patients, Portugal, 2016–2017. Emerging Infectious Diseases, 2018, 24, 762-766.	2.0	48

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19	High occurrence and unusual serotype diversity of non-typhoidal Salmonella in non-clinical niches, Angola. Epidemiology and Infection, 2017, 145, 883-886.	1.0	2
20	Tolerance to multiple metal stressors in emerging non-typhoidal MDR <i>Salmonella</i> serotypes: a relevant role for copper in anaerobic conditions. Journal of Antimicrobial Chemotherapy, 2016, 71, 2147-2157.	1.3	48
21	Stimulating Participation and Learning in Microbiology: Presence and Identification of Bacteria from Student's Hands. Journal of Food Science Education, 2016, 15, 51-55.	1.0	0
22	Clinical <i>Salmonella</i> Typhimurium ST34 with metal tolerance genes and an IncHI2 plasmid carrying <i>oqxAB-aac(6′)-lb-cr</i> from Europe. Journal of Antimicrobial Chemotherapy, 2016, 71, 843-845.	1.3	27
23	Salmonellosis: the role of poultry meat. Clinical Microbiology and Infection, 2016, 22, 110-121.	2.8	398
24	Relevance of <i>tcrYAZB</i> operon acquisition for <i>Enterococcus</i> survival at high copper concentrations under anaerobic conditions: TableÂ1 Journal of Antimicrobial Chemotherapy, 2016, 71, 560-563.	1.3	10
25	MCR-1 in multidrug-resistant and copper-tolerant clinically relevant Salmonella 1,4,[5],12:i:- and S. Rissen clones in Portugal, 2011 to 2015. Eurosurveillance, 2016, 21, .	3.9	103
26	Metal tolerance in emerging clinically relevant multidrug-resistant Salmonella enterica serotype 4,[5],12:i:â^' clones circulating in Europe. International Journal of Antimicrobial Agents, 2015, 45, 610-616.	1.1	85
27	Filling the map for antimicrobial resistance in sub-Saharan Africa: ampicillin-resistant <i>Enterococcus</i> from non-clinical sources in Angola: Table 1 Journal of Antimicrobial Chemotherapy, 2015, 70, 2914-2916.	1.3	16
28	A hospital sewage ST17 Enterococcus faecium with a transferable Inc18-like plasmid carrying genes coding for resistance to antibiotics and quaternary ammonium compounds (qacZ). Journal of Global Antimicrobial Resistance, 2015, 3, 49-51.	0.9	9
29	Ready-to-eat street-vended food as a potential vehicle of bacterial pathogens and antimicrobial resistance: An exploratory study in Porto region, Portugal. International Journal of Food Microbiology, 2015, 206, 1-6.	2.1	63
30	Characterization of the emerging clinically-relevant multidrug-resistant Salmonella enterica serotype 4,[5],12:i:- (monophasic variant of S. Typhimurium) clones. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 2249-2257.	1.3	39
31	Co-transfer of resistance to high concentrations of copper and first-line antibiotics among Enterococcus from different origins (humans, animals, the environment and foods) and clonal lineages. Journal of Antimicrobial Chemotherapy, 2014, 69, 899-906.	1.3	68
32	Clinically relevant multidrug resistant Salmonella enterica in swine and meat handlers at the abattoir. Veterinary Microbiology, 2014, 168, 229-233.	0.8	36
33	Microbiological quality of ready-to-eat salads: An underestimated vehicle of bacteria and clinically relevant antibiotic resistance genes. International Journal of Food Microbiology, 2013, 166, 464-470.	2.1	94
34	Salmonella enterica serotype Bovismorbificans, a new host for CTX-M-9. International Journal of Antimicrobial Agents, 2013, 41, 91-93.	1.1	5
35	Spread of multidrug-resistant Enterococcus to animals and humans: an underestimated role for the pig farm environment. Journal of Antimicrobial Chemotherapy, 2013, 68, 2746-2754.	1.3	74
36	Characterization of extended-spectrum beta-lactamases, antimicrobial resistance genes, and plasmid content in Escherichia coli isolates from different sources in Rio de Janeiro, Brazil. Diagnostic Microbiology and Infectious Disease, 2012, 74, 91-94.	0.8	12

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37	Spread of an OmpK36-modified ST15 Klebsiella pneumoniae variant during an outbreak involving multiple carbapenem-resistant Enterobacteriaceae species and clones. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 3057-3063.	1.3	54
38	Salmonella cross-contamination in swine abattoirs in Portugal: Carcasses, meat and meat handlers. International Journal of Food Microbiology, 2012, 157, 82-87.	2.1	53
39	First description of qnrS1-IncN plasmid in a ST11 Salmonella Enteritidis clinical isolate from Portugal. Diagnostic Microbiology and Infectious Disease, 2011, 69, 463-465.	0.8	14
40	Leakage of emerging clinically relevant multidrug-resistant Salmonella clones from pig farms. Journal of Antimicrobial Chemotherapy, 2011, 66, 2028-2032.	1.3	78
41	Emergence of an Incl plasmid encoding CMY-2 Â-lactamase associated with the international ST19 OXA-30-producing Â-lactamase Salmonella Typhimurium multidrug-resistant clone. Journal of Antimicrobial Chemotherapy, 2010, 65, 2097-2100.	1.3	22
42	Successful application of the DiversiLab repetitive-sequence-based PCR typing system for confirmation of the circulation of a multiresistant Pseudomonas aeruginosa clone in different hospital wards. Diagnostic Microbiology and Infectious Disease, 2010, 67, 202-206.	0.8	19
43	Dissemination of sul3 -Containing Elements Linked to Class 1 Integrons with an Unusual 3′ Conserved Sequence Region among Salmonella Isolates. Antimicrobial Agents and Chemotherapy, 2007, 51, 1545-1548.	1.4	113
44	Characterization of antimicrobial resistance and class 1 and 2 integrons in Salmonella enterica isolates from different sources in Portugal. Journal of Antimicrobial Chemotherapy, 2006, 58, 297-304.	1.3	100
45	Illegal use of nitrofurans in food animals: contribution to human salmonellosis?. Clinical Microbiology and Infection, 2006, 12, 1047-1049.	2.8	55
46	Dissemination of Sulfonamide Resistance Genes (sul1 , sul2 , and sul3) in Portuguese Salmonella enterica Strains and Relation with Integrons. Antimicrobial Agents and Chemotherapy, 2005, 49, 836-839.	1.4	235
47	Dissemination amongst humans and food products of animal origin of a Salmonella typhimurium clone expressing an integron-borne OXA-30 Â-lactamase. Journal of Antimicrobial Chemotherapy, 2004, 54, 429-434.	1.3	47
48	Incidence of Salmonella from poultry products and their susceptibility to antimicrobial agents. International Journal of Food Microbiology, 2003, 82, 97-103.	2.1	173
49	Incidence and Susceptibility to Antimicrobial Agents of Listeria spp. and Listeria monocytogenes Isolated from Poultry Carcasses in Porto, Portugal. Journal of Food Protection, 2002, 65, 1888-1893.	0.8	34