## Vera Manageiro

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1306926/publications.pdf

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75 papers 4,999 citations

218677 26 h-index 95266 68 g-index

76 all docs

76 docs citations

76 times ranked 7217 citing authors

#	Article	IF	Citations
1	SARS-CoV-2 introductions and early dynamics of the epidemic in Portugal. Communications Medicine, 2022, 2, .	4.2	5
2	Nocturnal Birds of Prey as Carriers of Staphylococcus aureus and Other Staphylococci: Diversity, Antimicrobial Resistance and Clonal Lineages. Antibiotics, 2022, 11, 240.	3.7	15
3	A One Health Approach Molecular Analysis of Staphylococcus aureus Reveals Distinct Lineages in Isolates from Miranda Donkeys (Equus asinus) and Their Handlers. Antibiotics, 2022, 11, 374.	3.7	7
4	Staphylococcus aureus and Methicillin-Resistant Coagulase-Negative Staphylococci in Nostrils and Buccal Mucosa of Healthy Camels Used for Recreational Purposes. Animals, 2022, 12, 1255.	2.3	3
5	Antimicrobial Resistance and Molecular Epidemiology of Staphylococcus aureus from Hunters and Hunting Dogs. Pathogens, 2022, 11, 548.	2.8	3
6	Clonal Diversity and Antimicrobial Resistance of Methicillin-Resistant Staphylococcus pseudintermedius Isolated from Canine Pyoderma. Microorganisms, 2021, 9, 482.	3.6	17
7	Antimicrobial Resistance and Genetic Lineages of Staphylococcus aureus from Wild Rodents: First Report of mecC-Positive Methicillin-Resistant S. aureus (MRSA) in Portugal. Animals, 2021, 11, 1537.	2.3	19
8	Prevalence and Characteristics of Multidrug-Resistant Livestock-Associated Methicillin-Resistant Staphylococcus aureus (LA-MRSA) CC398 Isolated from Quails (Coturnix Coturnix Japonica) Slaughtered for Human Consumption. Animals, 2021, 11, 2038.	2.3	22
9	Biofilm Formation of Multidrug-Resistant MRSA Strains Isolated from Different Types of Human Infections. Pathogens, 2021, 10, 970.	2.8	27
10	Assessing the Bacterial Community Composition of Bivalve Mollusks Collected in Aquaculture Farms and Respective Susceptibility to Antibiotics. Antibiotics, 2021, 10, 1135.	3.7	7
11	Distribution and Clonal Diversity of Staphylococcus aureus and Other Staphylococci in Surface Waters: Detection of ST425-t742 and ST130-t843 mecC-Positive MRSA Strains. Antibiotics, 2021, 10, 1416.	3.7	18
12	Emergence of community-acquired methicillin-resistant Staphylococcus aureus EMRSA-15 clone as the predominant cause of diabetic foot ulcer infections in Portugal. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 179-186.	2.9	34
13	Diversity of methicillin-resistant staphylococci among wild Lepus granatensis: first detection of mecA-MRSA in hares. FEMS Microbiology Ecology, 2020, 96, .	2.7	17
14	Genetic Relatedness and Diversity of Staphylococcus aureus from Different Reservoirs: Humans and Animals of Livestock, Poultry, Zoo, and Aquaculture. Microorganisms, 2020, 8, 1345.	3.6	15
15	Integrated chromosomal and plasmid sequence analyses reveal diverse modes of carbapenemase gene spread among <i>Klebsiella pneumoniae</i> . Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 25043-25054.	7.1	97
16	Bacterial Diversity and Antibiotic Susceptibility of Sparus aurata from Aquaculture. Microorganisms, 2020, 8, 1343.	3.6	20
17	Plasmid-Mediated Colistin Resistance (mcr-1) in Escherichia coli from Non-Imported Fresh Vegetables for Human Consumption in Portugal. Microorganisms, 2020, 8, 429.	3.6	14
18	Epidemiological situation, laboratory capacity and preparedness for carbapenem-resistant Acinetobacter baumannii in Europe, 2019. Eurosurveillance, 2020, 25, .	7.0	18

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19	Antibiotic resistance in foodborne bacteria. Trends in Food Science and Technology, 2019, 84, 41-44.	15.1	159
20	Epidemic of carbapenem-resistant Klebsiella pneumoniae in Europe is driven by nosocomial spread. Nature Microbiology, 2019, 4, 1919-1929.	13.3	476
21	First report of linezolid-resistant cfr-positive methicillin-resistant Staphylococcus aureus in humans in Portugal. Journal of Global Antimicrobial Resistance, 2019, 17, 323-325.	2.2	30
22	Revealing mcr-1-positive ESBL-producing Escherichia coli strains among Enterobacteriaceae from food-producing animals (bovine, swine and poultry) and meat (bovine and swine), Portugal, 2010–2015. International Journal of Food Microbiology, 2019, 296, 37-42.	4.7	41
23	IncX4 Plasmid Carrying the New mcr-1.9 Gene Variant in a CTX-M-8-Producing Escherichia coli Isolate Recovered From Swine. Frontiers in Microbiology, 2019, 10, 367.	3.5	28
24	Deciphering the role of cyanobacteria in water resistome: Hypothesis justifying the antibiotic resistance (phenotype and genotype) in Planktothrix genus. Science of the Total Environment, 2019, 652, 447-454.	8.0	24
25	Attributable deaths and disability-adjusted life-years caused by infections with antibiotic-resistant bacteria in the EU and the European Economic Area in 2015: a population-level modelling analysis. Lancet Infectious Diseases, The, 2019, 19, 56-66.	9.1	1,908
26	Molecular Epidemiology and Risk Factors of Carbapenemase-Producing Enterobacteriaceae Isolates in Portuguese Hospitals: Results From European Survey on Carbapenemase-Producing Enterobacteriaceae (EuSCAPE). Frontiers in Microbiology, 2018, 9, 2834.	<b>3.</b> 5	27
27	First report on MRSA CC398 recovered from wild boars in the north of Portugal. Are we facing a problem?. Science of the Total Environment, 2017, 596-597, 26-31.	8.0	28
28	Biochemical characterization of CTX-M-166, a new CTX-M $\hat{I}^2$ -lactamase produced by a commensal Escherichia coli isolate. Journal of Antibiotics, 2017, 70, 809-810.	2.0	2
29	Quantitative proteome analysis of an antibiotic resistant Escherichia coli exposed to tetracycline reveals multiple affected metabolic and peptidoglycan processes. Journal of Proteomics, 2017, 156, 20-28.	2.4	20
30	Bacterial Resistances., 2017,, 403-415.		0
31	New insights into resistance to colistin and third-generation cephalosporins of Escherichia coli in poultry, Portugal: Novel blaCTX-M-166 and blaESAC genes. International Journal of Food Microbiology, 2017, 263, 67-73.	4.7	23
32	Occurrence of carbapenemase-producing Klebsiella pneumoniae and Escherichia coli in the European survey of carbapenemase-producing Enterobacteriaceae (EuSCAPE): a prospective, multinational study. Lancet Infectious Diseases, The, 2017, 17, 153-163.	9.1	522
33	Genetic Background and Expression of the New qepA4 Gene Variant Recovered in Clinical TEM-1- and CMY-2-Producing Escherichia coli. Frontiers in Microbiology, 2017, 8, 1899.	3 <b>.</b> 5	8
34	Livestock-associated meticillin-resistant Staphylococcus aureus (MRSA) among human MRSA isolates, European Union/European Economic Area countries, 2013. Eurosurveillance, 2017, 22, .	7.0	66
35	QnrS1- and Aac(6′)-lb-cr-Producing Escherichia coli among Isolates from Animals of Different Sources: Susceptibility and Genomic Characterization. Frontiers in Microbiology, 2016, 7, 671.	3.5	22
36	Serotypes and Antibiotic Susceptibility of Streptococcus pneumoniae Isolates from Invasive Pneumococcal Disease and Asymptomatic Carriage in a Pre-vaccination Period, in Algeria. Frontiers in Microbiology, 2016, 7, 803.	3 <b>.</b> 5	14

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37	Salmonella Enteritidis Isolate Harboring Multiple Efflux Pumps and Pathogenicity Factors, Shows Absence of O Antigen Polymerase Gene. Frontiers in Microbiology, 2016, 7, 1130.	3.5	1
38	Architecture of Class 1, 2, and 3 Integrons from Gram Negative Bacteria Recovered among Fruits and Vegetables. Frontiers in Microbiology, 2016, 7, 1400.	3.5	61
39	Draft Genomic Analysis of an Avian Multidrug Resistant Morganella morganii Isolate Carrying qnrD1. Frontiers in Microbiology, 2016, 7, 1660.	3.5	18
40	Draft Genome Sequence of an Escherichia coli Strain Isolated from a Gallus gallus Broiler Producing the Novel CTX-M-166 Variant. Genome Announcements, 2016, 4, .	0.8	3
41	Influence of agricultural practice on mobile <scp><i>bla</i></scp> genes: <scp>Incl1</scp> â€bearing <scp>CTX</scp> â€vscp>M, <scp>SHV</scp> , <scp>CMY</scp> and <scp>TEM</scp> in <scp><i>E</i></scp> <i>Scp&gt;<i>EScp&gt;<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>E<i>EBEE<!--</td--><td>3.8</td><td>28</td></i></i></i></i></i></i></i></i></i></i></i></i></i></i></i></i></i></i></i>	3.8	28
42	Genetic Diversity and Antibiotic Resistance Among Coagulase-Negative Staphylococci Recovered from Birds of Prey in Portugal. Microbial Drug Resistance, 2016, 22, 727-730.	2.0	14
43	New Class 2 Integron In <i>2-4</i> Among Incl1-Positive <i>Escherichia coli</i> Isolates Carrying ESBL and PMAβ Genes from Food Animals in Portugal. Foodborne Pathogens and Disease, 2016, 13, 36-39.	1.8	15
44	Complete Sequence of a bla OXA-48 -Harboring IncL Plasmid from an Enterobacter cloacae Clinical Isolate. Genome Announcements, 2015, 3, .	0.8	6
45	Draft Genome Sequence of a Pathogenic O86:H25 Sequence Type 57 Escherichia coli Strain Isolated from Poultry and Carrying 12 Acquired Antibiotic Resistance Genes. Genome Announcements, 2015, 3, .	0.8	2
46	Assessing the antibiotic susceptibility of freshwater Cyanobacteria spp Frontiers in Microbiology, 2015, 6, 799.	3.5	46
47	CTX-M-15–ProducingEscherichia coliin Dolphin, Portugal. Emerging Infectious Diseases, 2015, 21, 2249-2251.	4.3	7
48	Antimicrobial susceptibility and oxymino- $\hat{l}^2$ -lactam resistance mechanisms in Salmonella enterica and Escherichia coli isolates from different animal sources. Research in Microbiology, 2015, 166, 574-583.	2.1	30
49	Draft Genome Sequence of the First NDM-1-Producing Providencia stuartii Strain Isolated in Portugal. Genome Announcements, 2015, 3, .	0.8	22
50	Capsular typing of Streptococcus pneumoniae isolated in an Algerian hospital using a new multiplex PCR-based scheme. Journal of Microbiological Methods, 2015, 119, 243-246.	1.6	8
51	Current perspectives on the dynamics of antibiotic resistance in different reservoirs. Research in Microbiology, 2015, 166, 594-600.	2.1	26
52	First description of food-borne Salmonella enterica resistance regions R1 and R3 associated with IS26 elements. Research in Microbiology, 2015, 166, 570-573.	2.1	3
53	Two novel CMY-2-type $\hat{l}^2$ -lactamases encountered in clinical Escherichia coli isolates. Annals of Clinical Microbiology and Antimicrobials, 2015, 14, 12.	3.8	9
54	Predominance of KPC-3 in a Survey for Carbapenemase-Producing Enterobacteriaceae in Portugal. Antimicrobial Agents and Chemotherapy, 2015, 59, 3588-3592.	3.2	41

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55	Molecular evidence of the close relatedness of clinical, gull and wastewater isolates of quinolone-resistant Escherichia coli. Journal of Global Antimicrobial Resistance, 2015, 3, 286-289.	2.2	35
56	GES-5 among the $\langle b \rangle \hat{l}^2 \langle b \rangle$ -lactamases detected in ubiquitous bacteria isolated from aquatic environment samples. FEMS Microbiology Letters, 2014, 351, 64-69.	1.8	34
57	Human, food and animal Campylobacter spp. isolated in Portugal: High genetic diversity and antibiotic resistance rates. International Journal of Antimicrobial Agents, 2014, 44, 306-313.	2.5	52
58	First Description of OXA-48 Carbapenemase Harbored by Escherichia coli and Enterobacter cloacae from a Single Patient in Portugal. Antimicrobial Agents and Chemotherapy, 2014, 58, 7613-7614.	3.2	18
59	Diversity of extended-spectrum and plasmid-mediated AmpC $\hat{l}^2$ -lactamases in Enterobacteriaceae isolates from portuguese health care facilities. Journal of Microbiology, 2014, 52, 496-503.	2.8	19
60	Clinically relevant multidrug resistant Salmonella enterica in swine and meat handlers at the abattoir. Veterinary Microbiology, 2014, 168, 229-233.	1.9	36
61	Occurrence of extended-spectrum $\hat{l}^2$ -lactamases among isolates of Salmonella enterica subsp. enterica from food-producing animals and food products, in Portugal. International Journal of Food Microbiology, 2013, 167, 221-228.	4.7	66
62	Assessing the molecular basis of transferable quinolone resistance in Escherichia coli and Salmonella spp. from food-producing animals and food products. Veterinary Microbiology, 2013, 167, 523-531.	1.9	42
63	Emergence and risk factors of β-lactamase–mediated resistance to oxyimino-β-lactams in Enterobacteriaceae isolates. Diagnostic Microbiology and Infectious Disease, 2012, 72, 272-277.	1.8	8
64	Genetic diversity and clonal evolution of carbapenem-resistant Acinetobacter baumannii isolates from Portugal and the dissemination of ST118. International Journal of Antimicrobial Agents, 2012, 40, 398-403.	2.5	23
65	Characterization of the Inhibitor-Resistant SHV $\hat{I}^2$ -Lactamase SHV-107 in a Clinical Klebsiella pneumoniae Strain Coproducing GES-7 Enzyme. Antimicrobial Agents and Chemotherapy, 2012, 56, 1042-1046.	3.2	11
66	Role of SHV $\hat{I}^2$ -lactamase variants in resistance of clinical Klebsiella pneumoniae strains to $\hat{I}^2$ -lactams in an Algerian hospital. Journal of Medical Microbiology, 2011, 60, 983-987.	1.8	18
67	Escherichia coli and Staphylococcus aureus: bad news and good news from the European Antimicrobial Resistance Surveillance Network (EARS-Net, formerly EARSS), 2002 to 2009. Eurosurveillance, 2011, 16, .	7.0	142
68	Biochemical Study of a New Inhibitor-Resistant β-Lactamase, SHV-84, Produced by a Clinical <i>Escherichia coli</i> Strain. Antimicrobial Agents and Chemotherapy, 2010, 54, 2271-2272.	3.2	11
69	Carbapenem-non-susceptible Enterobacteriaceae in Europe: conclusions from a meeting of national experts. Eurosurveillance, 2010, 15, .	7.0	212
70	The Lys234Arg Substitution in the Enzyme SHV-72 Is a Determinant for Resistance to Clavulanic Acid Inhibition. Antimicrobial Agents and Chemotherapy, 2008, 52, 1806-1811.	3.2	22
71	Biochemical Characterization of SHV-55, an Extended-Spectrum Class A $\hat{I}^2$ -Lactamase from <i>Klebsiella pneumoniae</i> . Antimicrobial Agents and Chemotherapy, 2008, 52, 1897-1898.	3.2	2
72	Spread of Extended-Spectrum β-Lactamase CTX-M-Producing Escherichia coli Clinical Isolates in Community and Nosocomial Environments in Portugal. Antimicrobial Agents and Chemotherapy, 2007, 51, 1946-1955.	3.2	137

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73	P563 Beta-lactam resistance mechanisms in clinical isolates of Proteus spp. in Portugal:plasmid-mediated inhibitor resistant TEM and extended-spectrum Î <sup>2</sup> -lactamases. International Journal of Antimicrobial Agents, 2007, 29, S127.	2.5	0
74	Binding of NorR to three DNA sites is essential for promoter activation of the flavorubredoxin gene, the nitric oxide reductase of Escherichia coli. Biochemical and Biophysical Research Communications, 2005, 328, 540-544.	2.1	15
75	Clonal diversity and antimicrobial resistance of & Department of the Clonal diversity and antimicrobial resistance of the Clonal diversity and the Clonal diversity an		0