Ã,ngela Novais

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	NDM-1 Introduction in Portugal through a ST11 KL105 Klebsiella pneumoniae Widespread in Europe. Antibiotics, 2022, 11, 92.	3.7	16
2	The Darkest Place Is under the Candlestick-Healthy Urogenital Tract as a Source of Worldwide Disseminated Extraintestinal Pathogenic Escherichia coli Lineages. Microorganisms, 2022, 10, 27.	3.6	0
3	High occurrence of colistin- and multidrug-resistant strains carrying mcr-1 or an underestimated mcr-1.26 allelic variant along a large Brazilian river. Journal of Global Antimicrobial Resistance, 2022, 30, 127-129.	2.2	4
4	MicroMundo@UPorto: an experimental microbiology project fostering student's antimicrobial resistance awareness and personal and social development. FEMS Microbiology Letters, 2021, 368, .	1.8	3
5	Phylogenomic analysis of a highly virulent Escherichia coli ST83 lineage with potential animal-human transmission. Microbial Pathogenesis, 2021, 155, 104920.	2.9	4
6	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.	3.8	15
7	Fourier Transform Infrared Spectroscopy (FT-IR) for Food and Water Microbiology. , 2021, , 191-217.		2
8	Allogenous Selection of Mutational Collateral Resistance: Old Drugs Select for New Resistance Within Antibiotic Families. Frontiers in Microbiology, 2021, 12, 757833.	3.5	15
9	Multiplicity of Carbapenemase-Producers Three Years after a KPC-3-Producing K. pneumoniae ST147-K64 Hospital Outbreak. Antibiotics, 2020, 9, 806.	3.7	16
10	Fourier transform infrared (FT-IR) spectroscopy typing: a real-time analysis of an outbreak by carbapenem-resistant Klebsiella pneumoniae. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 2471-2475.	2.9	15
11	Escherichia coli resistant to fosfomycin from urinary tract infections: Detection of the fosA3 gene in Spain. Journal of Global Antimicrobial Resistance, 2020, 21, 414-416.	2.2	16
12	A Front Line on Klebsiella pneumoniae Capsular Polysaccharide Knowledge: Fourier Transform Infrared Spectroscopy as an Accurate and Fast Typing Tool. MSystems, 2020, 5, .	3.8	32
13	Acquired AmpC β-Lactamases among Enterobacteriaceae from Healthy Humans and Animals, Food, Aquatic and Trout Aquaculture Environments in Portugal. Pathogens, 2020, 9, 273.	2.8	8
14	Commonality of Multidrug-Resistant Klebsiella pneumoniae ST348 Isolates in Horses and Humans in Portugal. Frontiers in Microbiology, 2019, 10, 1657.	3.5	14
15	Fourier transform infrared spectroscopy: unlocking fundamentals and prospects for bacterial strain typing. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 427-448.	2.9	92
16	Virulence genes, capsular and plasmid types of multidrug-resistant CTX-M(-2, -8, -15) and KPC-2-producing Klebsiella pneumoniae isolates from four major hospitals in Brazil. Diagnostic Microbiology and Infectious Disease, 2018, 91, 164-168.	1.8	22
17	New fluorescent rosamine chelator showing promising antibacterial activity against Gram-positive bacteria. Bioorganic Chemistry, 2018, 79, 341-349.	4.1	8
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.	4.3	48

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19	Different Escherichia coli B2-ST131 clades (B and C) producing extended-spectrum β-lactamases (ESBL) colonizing residents of Portuguese nursing homes. Epidemiology and Infection, 2017, 145, 3303-3306.	2.1	11
20	Long-Term Care Facility (LTCF) Residents Colonized With Multidrug-Resistant (MDR) Klebsiella pneumoniae Lineages Frequently Causing Infections in Portuguese Clinical Institutions. Infection Control and Hospital Epidemiology, 2017, 38, 1127-1130.	1.8	7
21	Elucidating constraints for differentiation of major human Klebsiella pneumoniae clones using MALDI-TOF MS. European Journal of Clinical Microbiology and Infectious Diseases, 2017, 36, 379-386.	2.9	18
22	Citrobacter portucalensis sp. nov., isolated from an aquatic sample. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3513-3517.	1.7	40
23	KPC-3-Producing Klebsiella pneumoniae in Portugal Linked to Previously Circulating Non-CG258 Lineages and Uncommon Genetic Platforms (Tn4401d-IncFIA and Tn4401d-IncN). Frontiers in Microbiology, 2016, 7, 1000.	3.5	54
24	Importation of Fosfomycin Resistance <i>fosA3</i> Gene to Europe. Emerging Infectious Diseases, 2016, 22, 346-348.	4.3	25
25	Efficient transmission of IncFIIY and IncL plasmids and Klebsiella pneumoniae ST101 clone producing OXA-48, NDM-1 or OXA-181 in Bucharest hospitals. International Journal of Antimicrobial Agents, 2016, 48, 223-224.	2.5	14
26	An update on faecal carriage of ESBL-producing Enterobacteriaceae by Portuguese healthy humans: detection of theH30 subclone of B2-ST131Escherichia coliproducing CTX-M-27: TableÂ1 Journal of Antimicrobial Chemotherapy, 2016, 71, 1120-1122.	3.0	35
27	Atypical epidemiology of CTX-M-15 among Enterobacteriaceae from a high diversity of non-clinical niches in Angola: TableÂ1 Journal of Antimicrobial Chemotherapy, 2016, 71, 1169-1173.	3.0	28
28	Antibiotic-Resistant Klebsiella pneumoniae and Escherichia coli High-Risk Clones and an IncFII _k Mosaic Plasmid Hosting Tn <i>1</i> (<i>bla</i> _{TEM-4}) in Isolates from 1990 to 2004. Antimicrobial Agents and Chemotherapy, 2015, 59, 2904-2908.	3.2	9
29	Snapshot on carbapenemase-producing <i>Pseudomonas aeruginosa</i> and <i>Acinetobacter baumannii</i> in Bucharest hospitals reveals unusual clones and novel genetic surroundings for <i>bla</i> OXA-23. Journal of Antimicrobial Chemotherapy, 2015, 70, 1016-1020.	3.0	27
30	Increase of widespread A, B1 and D Escherichia coli clones producing a high diversity of CTX-M-types in a Portuguese hospital. Future Microbiology, 2015, 10, 1125-1131.	2.0	18
31	Phylogeny and Comparative Genomics Unveil Independent Diversification Trajectories of <i>qnrB</i> and Genetic Platforms within Particular Citrobacter Species. Antimicrobial Agents and Chemotherapy, 2015, 59, 5951-5958.	3.2	55
32	Evaluation of the Recently Launched Rapid Carb Blue Kit for Detection of Carbapenemase-Producing Gram-Negative Bacteria. Journal of Clinical Microbiology, 2015, 53, 3105-3107.	3.9	22
33	Detection of VIM-34, a novel VIM-1 variant identified in the intercontinental ST15 Klebsiella pneumoniae clone. Journal of Antimicrobial Chemotherapy, 2014, 69, 274-275.	3.0	10
34	MALDI-TOF mass spectrometry as a tool for the discrimination of high-risk Escherichia coli clones from phylogenetic groups B2 (ST131) and D (ST69, ST405, ST393). European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 1391-1399.	2.9	48
35	Expansion of ESBL-producing Klebsiella pneumoniae in hospitalized patients: A successful story of international clones (ST15, ST147, ST336) and epidemic plasmids (IncR, IncFIIK). International Journal of Medical Microbiology, 2014, 304, 1100-1108.	3.6	120
36	Long-term dissemination of acquired AmpC β-lactamases among Klebsiella spp. and Escherichia coli in Portuguese clinical settings. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 551-558.	2.9	17

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37	Diversity and biofilm-production ability among isolates of Escherichia coli phylogroup D belonging to ST69, ST393 and ST405 clonal groups. BMC Microbiology, 2013, 13, 144.	3.3	35
38	Incl1/ST3 and IncN/ST1 plasmids drive the spread of blaTEM-52 and blaCTX-M-1/-32 in diverse Escherichia coli clones from different piggeries. Journal of Antimicrobial Chemotherapy, 2013, 68, 2245-8.	3.0	30
39	Salmonella enterica serotype Bovismorbificans, a new host for CTX-M-9. International Journal of Antimicrobial Agents, 2013, 41, 91-93.	2.5	5
40	Blue-Carba, an Easy Biochemical Test for Detection of Diverse Carbapenemase Producers Directly from Bacterial Cultures. Journal of Clinical Microbiology, 2013, 51, 4281-4283.	3.9	172
41	Diverse high-risk B2 and D Escherichia coli clones depicted by Fourier Transform Infrared Spectroscopy. Scientific Reports, 2013, 3, 3278.	3.3	39
42	Contribution of IncFII and Broad-Host IncA/C and IncN Plasmids to the Local Expansion and Diversification of Phylogroup B2 Escherichia coli ST131 Clones Carrying <i>bla</i> _{CTX-M-15} and <i>qnrS1</i> Genes. Antimicrobial Agents and Chemotherapy, 2012, 56, 2763-2766.	3.2	27
43	Role of Common bla OXA-24/OXA-40 -Carrying Platforms and Plasmids in the Spread of OXA-24/OXA-40 among Acinetobacter Species Clinical Isolates. Antimicrobial Agents and Chemotherapy, 2012, 56, 3969-3972.	3.2	59
44	Characterization of Globally Spread Escherichia coli ST131 Isolates (1991 to 2010). Antimicrobial Agents and Chemotherapy, 2012, 56, 3973-3976.	3.2	49
45	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.	2.9	54
46	<i>In Vitro</i> Selection of Variants Resistant to β-Lactams plus β-Lactamase Inhibitors in CTX-M β-Lactamases: Predicting the <i>In Vivo</i> Scenario?. Antimicrobial Agents and Chemotherapy, 2011, 55, 4530-4536.	3.2	39
47	International Spread and Persistence of TEM-24 Is Caused by the Confluence of Highly Penetrating <i>Enterobacteriaceae</i> Clones and an IncA/C ₂ Plasmid Containing Tn <i>1696</i> ::Tn <i>1</i> and IS <i>5075</i> -Tn <i>21</i> . Antimicrobial Agents and Chemotherapy, 2010, 54, 825-834.	3.2	41
48	Evolutionary Trajectories of Beta-Lactamase CTX-M-1 Cluster Enzymes: Predicting Antibiotic Resistance. PLoS Pathogens, 2010, 6, e1000735.	4.7	100
49	Spread of <i>bla</i> _{CTX-M-14} Is Driven Mainly by IncK Plasmids Disseminated among <i>Escherichia coli</i> Phylogroups A, B1, and D in Spain. Antimicrobial Agents and Chemotherapy, 2009, 53, 5204-5212.	3.2	112
50	Extended-spectrum β-lactamase-producing Escherichia coli in Spain belong to a large variety of multilocus sequence typing types, including ST10 complex/A, ST23 complex/A and ST131/B2. International Journal of Antimicrobial Agents, 2009, 34, 173-176.	2.5	164
51	Prevalence and spread of extended-spectrum \hat{l}^2 -lactamase-producing Enterobacteriaceae in Europe. Clinical Microbiology and Infection, 2008, 14, 144-153.	6.0	495
52	Mutational Events in Cefotaximase Extended-Spectrum β-Lactamases of the CTX-M-1 Cluster Involved in Ceftazidime Resistance. Antimicrobial Agents and Chemotherapy, 2008, 52, 2377-2382.	3.2	40
53	Dissemination of Clonally Related <i>Escherichia coli</i> Strains Expressing Extended-Spectrum β-Lactamase CTX-M-15. Emerging Infectious Diseases, 2008, 14, 195-200.	4.3	672
54	Emergence and Dissemination of Enterobacteriaceae Isolates Producing CTX-M-1-Like Enzymes in Spain Are Associated with IncFII (CTX-M-15) and Broad-Host-Range (CTX-M-1, -3, and -32) Plasmids. Antimicrobial Agents and Chemotherapy, 2007, 51, 796-799.	3.2	110

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55	Preservation of Integron Types among <i>Enterobacteriaceae</i> Producing Extended-Spectrum β-Lactamases in a Spanish Hospital over a 15-Year Period (1988 to 2003). Antimicrobial Agents and Chemotherapy, 2007, 51, 2201-2204.	3.2	42
56	High diversity of extended-spectrum Â-lactamases among clinical isolates of Enterobacteriaceae from Portugal. Journal of Antimicrobial Chemotherapy, 2007, 60, 1370-1374.	3.0	53
57	Dissemination and Persistence of bla CTX-M-9 Are Linked to Class 1 Integrons Containing CR1 Associated with Defective Transposon Derivatives from Tn 402 Located in Early Antibiotic Resistance Plasmids of IncHI2, IncP1-î±, and IncFI Groups. Antimicrobial Agents and Chemotherapy, 2006, 50, 2741-2750.	3.2	108
58	Antibiotic Coresistance in Extended-Spectrum-β-Lactamase-Producing <i>Enterobacteriaceae</i> and In Vitro Activity of Tigecycline. Antimicrobial Agents and Chemotherapy, 2006, 50, 2695-2699.	3.2	145