Marta Fernãndez-Martãnez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3232668/publications.pdf

Version: 2024-02-01

20 papers 750 citations

759233 12 h-index 677142 22 g-index

22 all docs 22 docs citations

times ranked

22

1317 citing authors

#	Article	IF	CITATIONS
1	Prospective Multicenter Study of Carbapenemase-Producing Enterobacteriaceae from 83 Hospitals in Spain Reveals High <i>In Vitro</i> Susceptibility to Colistin and Meropenem. Antimicrobial Agents and Chemotherapy, 2015, 59, 3406-3412.	3.2	130
2	Genomics and Susceptibility Profiles of Extensively Drug-Resistant Pseudomonas aeruginosa Isolates from Spain. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	108
3	Comprehensive clinical and epidemiological assessment of colonisation and infection due to carbapenemase-producing Enterobacteriaceae in Spain. Journal of Infection, 2016, 72, 152-160.	3.3	73
4	Occurrence of Corynebacterium striatum as an emerging antibiotic-resistant nosocomial pathogen in a Tunisian hospital. Scientific Reports, 2017, 7, 9704.	3.3	69
5	Carbapenem-resistant Klebsiella pneumoniae isolates from Egypt containing bla NDM-1 on IncR plasmids and its association with rmtF. International Journal of Infectious Diseases, 2016, 43, 17-20.	3.3	60
6	Biofilm formation by multidrug resistant Enterobacteriaceae strains isolated from solid organ transplant recipients. Scientific Reports, 2019, 9, 8928.	3.3	59
7	First identification of NDM-5 associated with OXA-181 in <i>Escherichia coli</i> from Egypt. Emerging Microbes and Infections, 2016, 5, 1-12.	6.5	52
8	Molecular identification of aminoglycoside-modifying enzymes in clinical isolates of Escherichia coli resistant to amoxicillin/clavulanic acid isolated in Spain. International Journal of Antimicrobial Agents, 2015, 46, 157-163.	2.5	36
9	Isolation of VIM-2-Producing Pseudomonas monteilii Clinical Strains Disseminated in a Tertiary Hospital in Northern Spain. Antimicrobial Agents and Chemotherapy, 2015, 59, 1334-1336.	3.2	29
10	Prevalence of Aminoglycoside-Modifying Enzymes in <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> Producing Extended Spectrum β-Lactamases Collected in Two Multicenter Studies in Spain. Microbial Drug Resistance, 2018, 24, 367-376.	2.0	26
11	Susceptibility to Aminoglycosides and Distribution of aph and aac(3)-XI Genes among Corynebacterium striatum Clinical Isolates. PLoS ONE, 2016, 11, e0167856.	2.5	18
12	Activity of ceftazidime–avibactam against multidrug-resistance Enterobacteriaceae expressing combined mechanisms of resistance. Enfermedades Infecciosas Y MicrobiologÃa ClÃnica, 2017, 35, 499-504.	0.5	13
13	Oral decontamination with colistin plus neomycin in solid organ transplant recipients colonized by multidrug-resistant Enterobacterales: a multicentre, randomized, controlled, open-label, parallel-group clinical trial. Clinical Microbiology and Infection, 2021, 27, 856-863.	6.0	13
14	Rates of faecal colonization by carbapenemase-producing Enterobacteriaceae among patients admitted to ICUs in Spain: Table 1 Journal of Antimicrobial Chemotherapy, 2015, 70, 2916-2918.	3.0	11
15	Molecular characterization of multidrug resistant Enterobacterales strains isolated from liver and kidney transplant recipients in Spain. Scientific Reports, 2021, 11, 11875.	3.3	10
16	Aminoglycoside resistance determinants in multiresistant Escherichia coli and Klebsiella pneumoniae clinical isolates from Turkish and Syrian patients. Acta Microbiologica Et Immunologica Hungarica, 2019, 66, 327-335.	0.8	8
17	Antimicrobial Susceptibility and Characterization of Resistance Mechanisms of Corynebacterium urealyticum Clinical Isolates. Antibiotics, 2020, 9, 404.	3.7	8
18	High-risk clones and novel sequence type ST4497 of Klebsiella pneumoniae clinical isolates producing different alleles of NDM-type and other carbapenemases from a single tertiary-care centre in Egypt. International Journal of Antimicrobial Agents, 2020, 56, 106164.	2.5	7

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19	Draft Genome Sequence of the Quorum-Sensing and Biofilm-Producing Pseudomonas aeruginosa Strain Pae221, Belonging to the Epidemic High-Risk Clone Sequence Type 274. Genome Announcements, 2015, 3, .	0.8	5
20	Whole-genome sequencing reveals misidentification of a multidrug-resistant urine clinical isolate as Corynebacterium urealyticum. Journal of Global Antimicrobial Resistance, 2020, 23, 16-19.	2.2	5