

Paula GÃ“mez

List of Publications by Year in descending order

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Version: 2024-02-01

30
papers

914
citations

430874

18
h-index

477307

29
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30
all docs

30
docs citations

30
times ranked

997
citing authors

#	ARTICLE	IF	CITATIONS
1	Prevalence, antibiotic resistance, virulence traits and genetic lineages of <i>Staphylococcus aureus</i> in healthy sheep in Tunisia. <i>Veterinary Microbiology</i> , 2012, 156, 367-373.	1.9	77
2	Detection of methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) carrying the <i>mecC</i> gene in wild small mammals in Spain. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 2061-2064.	3.0	74
3	High prevalence of <i>spa</i> types associated with the clonal lineage CC398 among tetracycline-resistant methicillin-resistant <i>Staphylococcus aureus</i> strains in a Spanish hospital. <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 330-334.	3.0	69
4	Detection of MRSA ST3061-t843- <i>mecC</i> and ST398-t011- <i>mecA</i> in white stork nestlings exposed to human residues: Table 1.. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 53-57.	3.0	69
5	Identification of novel <i>vga(A)</i> -carrying plasmids and a Tn5406-like transposon in methicillin-resistant <i>Staphylococcus aureus</i> and <i>Staphylococcus epidermidis</i> of human and animal origin. <i>International Journal of Antimicrobial Agents</i> , 2012, 40, 306-312.	2.5	48
6	Antimicrobial resistance determinants in <i>Staphylococcus</i> spp. recovered from birds of prey in Portugal. <i>Veterinary Microbiology</i> , 2014, 171, 436-440.	1.9	46
7	Methicillin-resistant <i>Staphylococcus aureus</i> of lineage ST398 as cause of mastitis in cows. <i>Letters in Applied Microbiology</i> , 2014, 59, 665-669.	2.2	45
8	Characterization of staphylococci in urban wastewater treatment plants in Spain, with detection of methicillin resistant <i>Staphylococcus aureus</i> ST398. <i>Environmental Pollution</i> , 2016, 212, 71-76.	7.5	41
9	High prevalence of methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) carrying the <i>mecC</i> gene in a semi-extensive red deer (<i>Cervus elaphus hispanicus</i>) farm in Southern Spain. <i>Veterinary Microbiology</i> , 2015, 177, 326-331.	1.9	40
10	Characterization of <i>Staphylococcus aureus</i> from Raw Meat Samples in Tunisia: Detection of Clonal Lineage ST398 from the African Continent. <i>Foodborne Pathogens and Disease</i> , 2015, 12, 686-692.	1.8	39
11	Human <i>mecC</i> -Carrying MRSA: Clinical Implications and Risk Factors. <i>Microorganisms</i> , 2020, 8, 1615.	3.6	35
12	Detection of MRSA of Lineages CC130- <i>mecC</i> and CC398- <i>mecA</i> and <i>Staphylococcus delphini</i> - <i>lnu(A)</i> in Magpies and Cinereous Vultures in Spain. <i>Microbial Ecology</i> , 2019, 78, 409-415.	2.8	33
13	Characterization of methicillin-resistant coagulase-negative staphylococci in milk from cows with mastitis in Brazil. <i>Antonie Van Leeuwenhoek</i> , 2014, 106, 227-233.	1.7	31
14	Molecular Characterization and Clonal Diversity of Methicillin-Resistant and -Susceptible <i>Staphylococcus aureus</i> Isolates of Milk of Cows with Clinical Mastitis in Tunisia. <i>Microbial Drug Resistance</i> , 2018, 24, 1210-1216.	2.0	27
15	Frequency and Characterization of Antimicrobial Resistance and Virulence Genes of Coagulase-Negative Staphylococci from Wild Birds in Spain. Detection of <i>tst</i> -Carrying <i>S. sciuri</i> Isolates. <i>Microorganisms</i> , 2020, 8, 1317.	3.6	24
16	Genetic Lineages, Antimicrobial Resistance, and Virulence in <i>Staphylococcus aureus</i> of Meat Samples in Spain: Analysis of Immune Evasion Cluster (IEC) Genes. <i>Foodborne Pathogens and Disease</i> , 2014, 11, 354-356.	1.8	23
17	Diversity of species and antimicrobial resistance determinants of staphylococci in superficial waters in Spain. <i>FEMS Microbiology Ecology</i> , 2017, 93, fiw208.	2.7	22
18	Molecular characterization of <i>Staphylococcus aureus</i> isolated from humans related to a livestock farm in Spain, with detection of MRSA-CC130 carrying <i>mecC</i> gene: A zoonotic case?. <i>Enfermedades Infecciosas Y Microbiologa Clnica</i> , 2016, 34, 280-285.	0.5	21

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19	Staphylococcus aureus isolated from wastewater treatment plants in Tunisia: occurrence of human and animal associated lineages. <i>Journal of Water and Health</i> , 2017, 15, 638-643.	2.6	19
20	Diversity of staphylococcal species in food producing animals in Spain, with detection of PVL-positive MRSA ST8 (USA300). <i>Veterinary Microbiology</i> , 2019, 233, 5-10.	1.9	16
21	Clonal lineages detected amongst tetracycline-resistant methicillin-resistant <i>Staphylococcus aureus</i> isolates of a Tunisian hospital, with detection of lineage ST398. <i>Journal of Medical Microbiology</i> , 2015, 64, 623-629.	1.8	15
22	Antimicrobial Resistance, Virulence, and Genetic Lineages of Staphylococci from Horses Destined for Human Consumption: High Detection of <i>S. aureus</i> Isolates of Lineage ST1640 and Those Carrying the lukPQ Gene. <i>Animals</i> , 2019, 9, 900.	2.3	15
23	Genetic Diversity and Antibiotic Resistance Among Coagulase-Negative Staphylococci Recovered from Birds of Prey in Portugal. <i>Microbial Drug Resistance</i> , 2016, 22, 727-730.	2.0	14
24	Genetic lineages and antimicrobial resistance genotypes in <i>Staphylococcus aureus</i> from children with atopic dermatitis: detection of clonal complexes CC1, CC97 and CC398. <i>Journal of Chemotherapy</i> , 2016, 28, 359-366.	1.5	14
25	Livestock-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) in Purulent Subcutaneous Lesions of Farm Rabbits. <i>Foods</i> , 2020, 9, 439.	4.3	14
26	Genetic characterization of <i>Staphylococcus aureus</i> isolated from nasal samples of healthy ewes in Tunisia. High prevalence of CC130 and CC522 lineages. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2017, 51, 37-40.	1.6	12
27	Genomic Analysis of <i>Staphylococcus aureus</i> of the Lineage CC130, Including mecC-Carrying MRSA and MSSA Isolates Recovered of Animal, Human, and Environmental Origins. <i>Frontiers in Microbiology</i> , 2021, 12, 655994.	3.5	12
28	Molecular Epidemiology of <i>Staphylococcus aureus</i> Lineages in the Animal-Human Interface. , 2018, , 189-214.		9
29	Identification of Enterococci, Staphylococci, and Enterobacteriaceae from Slurries and Air in and around Two Pork Farms. <i>Journal of Food Protection</i> , 2018, 81, 1776-1782.	1.7	6
30	Simultaneous Nasal Carriage by Methicillin-Resistant and Methicillin Susceptible <i>Staphylococcus aureus</i> of Lineage ST398 in a Live Pig Transporter. <i>Pathogens</i> , 2020, 9, 401.	2.8	4