

# Marcia Giambiagi-Demarval

## List of Publications by Year in descending order

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Version: 2024-02-01

25  
papers

476  
citations

759233

12  
h-index

713466

21  
g-index

25  
all docs

25  
docs citations

25  
times ranked

539  
citing authors

#	ARTICLE	IF	CITATIONS
1	Detection of ileS-2 gene encoding mupirocin resistance in methicillin-resistant <i>Staphylococcus aureus</i> by multiplex PCR. <i>Diagnostic Microbiology and Infectious Disease</i> , 1999, 34, 77-81.	1.8	73
2	The gene <i>bap</i> , involved in biofilm production, is present in <i>Staphylococcus</i> spp. strains from nosocomial infections. <i>Journal of Microbiology</i> , 2009, 47, 319-326.	2.8	57
3	Underrated <i>Staphylococcus</i> species and their role in antimicrobial resistance spreading. <i>Genetics and Molecular Biology</i> , 2020, 43, e20190065.	1.3	48
4	CRISPR-Cas Systems Features and the Gene-Reservoir Role of Coagulase-Negative Staphylococci. <i>Frontiers in Microbiology</i> , 2017, 8, 1545.	3.5	40
5	Identification of coagulase-negative staphylococci from bovine mastitis using RFLP-PCR of the <i>groEL</i> gene. <i>Veterinary Microbiology</i> , 2008, 130, 134-140.	1.9	31
6	Heat-Resistance and Heat-Shock Response in the Nosocomial Pathogen <i>Enterococcus faecium</i> . <i>Current Microbiology</i> , 2003, 46, 313-317.	2.2	26
7	Transfer of mupirocin resistance from <i>Staphylococcus haemolyticus</i> clinical strains to <i>Staphylococcus aureus</i> through conjugative and mobilizable plasmids. <i>FEMS Microbiology Letters</i> , 2016, 363, fnw121.	1.8	25
8	Phenotypic and Genotypic Characterization of Biofilm Formation in <i>Staphylococcus haemolyticus</i> . <i>Current Microbiology</i> , 2015, 70, 829-834.	2.2	23
9	<i>Staphylococcus chromogenes</i> , a Coagulase-Negative <i>Staphylococcus</i> Species That Can Clot Plasma. <i>Journal of Clinical Microbiology</i> , 2016, 54, 1372-1375.	3.9	21
10	Occurrence of virulence-associated genes among <i>Staphylococcus saprophyticus</i> isolated from different sources. <i>Microbial Pathogenesis</i> , 2018, 119, 9-11.	2.9	20
11	The oral microbiota of domestic cats harbors a wide variety of <i>Staphylococcus</i> species with zoonotic potential. <i>Veterinary Microbiology</i> , 2017, 201, 136-140.	1.9	19
12	CRISPR tracking reveals global spreading of antimicrobial resistance genes by <i>Staphylococcus</i> of canine origin. <i>Veterinary Microbiology</i> , 2019, 232, 65-69.	1.9	16
13	The CtsR regulator controls the expression of <i>clpC</i> , <i>clpE</i> and <i>clpP</i> and is required for the virulence of <i>Enterococcus faecalis</i> in an invertebrate model. <i>Antonie Van Leeuwenhoek</i> , 2016, 109, 1253-1259.	1.7	13
14	Short communication: Diversity of species and transmission of antimicrobial resistance among <i>Staphylococcus</i> spp. isolated from goat milk. <i>Journal of Dairy Science</i> , 2019, 102, 5518-5524.	3.4	10
15	Identification of <i>Staphylococcus epidermidis</i> with transferrable mupirocin resistance from canine skin. <i>Veterinary Journal</i> , 2018, 235, 70-72.	1.7	9
16	The influence of pH on <i>Staphylococcus saprophyticus</i> iron metabolism and the production of siderophores. <i>Microbes and Infection</i> , 2019, 21, 456-463.	1.9	8
17	Interspecies transfer of plasmid-borne gentamicin resistance between <i>Staphylococcus</i> isolated from domestic dogs to <i>Staphylococcus aureus</i> . <i>Infection, Genetics and Evolution</i> , 2022, 98, 105230.	2.3	8
18	Expression of the stress-response regulators CtsR and HrcA in the uropathogen <i>Staphylococcus saprophyticus</i> during heat shock. <i>Antonie Van Leeuwenhoek</i> , 2017, 110, 1105-1111.	1.7	7

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19	The exoproteome profiles of three <i>Staphylococcus saprophyticus</i> strains reveal diversity in protein secretion contents. <i>Microbiological Research</i> , 2018, 216, 85-96.	5.3	6
20	<i>Staphylococcus saprophyticus</i> Proteomic Analyses Elucidate Differences in the Protein Repertoires among Clinical Strains Related to Virulence and Persistence. <i>Pathogens</i> , 2020, 9, 69.	2.8	6
21	Identification of coagulase-negative <i>Staphylococcus saprophyticus</i> by polymerase chain reaction based on the heat-shock repressor encoding <i>hrcA</i> gene. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 86, 253-256.	1.8	4
22	Characterization of biofilms and antimicrobial resistance of coagulase-negative <i>Staphylococcus</i> species involved with subclinical mastitis. <i>Journal of Dairy Research</i> , 2021, 88, 179-184.	1.4	3
23	<i>Staphylococcus nepalensis</i> , a commensal of the oral microbiota of domestic cats, is a reservoir of transferrable antimicrobial resistance. <i>Microbiology (United Kingdom)</i> , 2020, 166, 727-734.	1.8	3
24	A proteomic dataset of secreted proteins by three <i>Staphylococcus saprophyticus</i> strains. <i>Data in Brief</i> , 2018, 21, 1472-1476.	1.0	0
25	Accurate identification of atypical <i>Staphylococcus chromogenes</i> plasma-clotting strains causing bovine mastitis. <i>Ciencia Rural</i> , 2019, 49, .	0.5	0