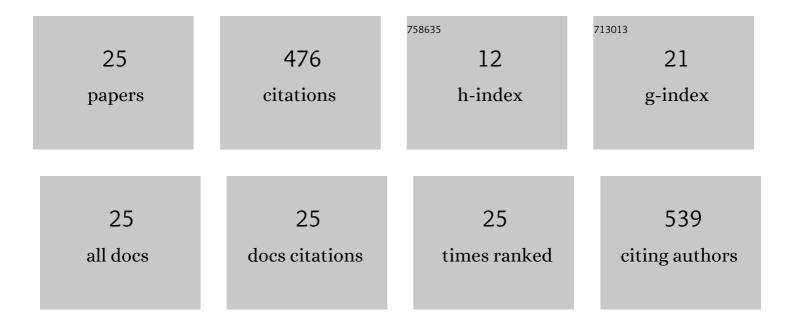
Marcia Giambiagi-Demarval

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Detection of ileS-2 gene encoding mupirocin resistance in methicillin-resistant Staphylococcus aureus by multiplex PCR. Diagnostic Microbiology and Infectious Disease, 1999, 34, 77-81.	0.8	73
2	The gene bap, involved in biofilm production, is present in Staphylococcus spp. strains from nosocomial infections. Journal of Microbiology, 2009, 47, 319-326.	1.3	57
3	Underrated Staphylococcus species and their role in antimicrobial resistance spreading. Genetics and Molecular Biology, 2020, 43, e20190065.	0.6	48
4	CRISPR-Cas Systems Features and the Gene-Reservoir Role of Coagulase-Negative Staphylococci. Frontiers in Microbiology, 2017, 8, 1545.	1.5	40
5	Identification of coagulase-negative staphylococci from bovine mastitis using RFLP-PCR of the groEL gene. Veterinary Microbiology, 2008, 130, 134-140.	0.8	31
6	Heat-Resistance and Heat-Shock Response in the Nosocomial Pathogen Enterococcus faecium. Current Microbiology, 2003, 46, 313-317.	1.0	26
7	Transfer of mupirocin resistance from <i>Staphylococcus haemolyticus</i> clinical strains to <i>Staphylococcus aureus</i> through conjugative and mobilizable plasmids. FEMS Microbiology Letters, 2016, 363, fnw121.	0.7	25
8	Phenotypic and Genotypic Characterization of Biofilm Formation in Staphylococcus haemolyticus. Current Microbiology, 2015, 70, 829-834.	1.0	23
9	Staphylococcus chromogenes, a Coagulase-Negative Staphylococcus Species That Can Clot Plasma. Journal of Clinical Microbiology, 2016, 54, 1372-1375.	1.8	21
10	Occurrence of virulence-associated genes among Staphylococcus saprophyticus isolated from different sources. Microbial Pathogenesis, 2018, 119, 9-11.	1.3	20
11	The oral microbiota of domestic cats harbors a wide variety of Staphylococcus species with zoonotic potential. Veterinary Microbiology, 2017, 201, 136-140.	0.8	19
12	CRISPR tracking reveals global spreading of antimicrobial resistance genes by Staphylococcus of canine origin. Veterinary Microbiology, 2019, 232, 65-69.	0.8	16
13	The CtsR regulator controls the expression of clpC, clpE and clpP and is required for the virulence of Enterococcus faecalis in an invertebrate model. Antonie Van Leeuwenhoek, 2016, 109, 1253-1259.	0.7	13
14	Short communication: Diversity of species and transmission of antimicrobial resistance among Staphylococcus spp. isolated from goat milk. Journal of Dairy Science, 2019, 102, 5518-5524.	1.4	10
15	Identification of Staphylococcus epidermidis with transferrable mupirocin resistance from canine skin. Veterinary Journal, 2018, 235, 70-72.	0.6	9
16	The influence of pH on Staphylococcus saprophyticus iron metabolism and the production of siderophores. Microbes and Infection, 2019, 21, 456-463.	1.0	8
17	Interspecies transfer of plasmid-borne gentamicin resistance between Staphylococcus isolated from domestic dogs to Staphylococcus aureus. Infection, Genetics and Evolution, 2022, 98, 105230.	1.0	8
18	Expression of the stress-response regulators CtsR and HrcA in the uropathogen Staphylococcus saprophyticus during heat shock. Antonie Van Leeuwenhoek, 2017, 110, 1105-1111.	0.7	7

#	Article	IF	CITATIONS
19	The exoproteome profiles of three Staphylococcus saprophyticus strains reveal diversity in protein secretion contents. Microbiological Research, 2018, 216, 85-96.	2.5	6
20	Staphylococcus saprophyticus Proteomic Analyses Elucidate Differences in the Protein Repertories among Clinical Strains Related to Virulence and Persistence. Pathogens, 2020, 9, 69.	1.2	6
21	Identification of coagulase-negative Staphylococcus saprophyticus by polymerase chain reaction based on the heat-shock repressor encoding hrcA gene. Diagnostic Microbiology and Infectious Disease, 2016, 86, 253-256.	0.8	4
22	Characterization of biofilms and antimicrobial resistance of coagulase-negative <i>Staphylococcus</i> species involved with subclinical mastitis. Journal of Dairy Research, 2021, 88, 179-184.	0.7	3
23	Staphylococcus nepalensis, a commensal of the oral microbiota of domestic cats, is a reservoir of transferrable antimicrobial resistance. Microbiology (United Kingdom), 2020, 166, 727-734.	0.7	3
24	A proteomic dataset of secreted proteins by three Staphylococcus saprophyticus strains. Data in Brief, 2018, 21, 1472-1476.	0.5	0
25	Accurate identification of atypical Staphylococcus chromogenes plasma-clotting strains causing bovine mastitis. Ciencia Rural. 2019. 49	0.3	0