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List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

751
citations

840776

11
h-index

713466

21
g-index

22
all docs

22
docs citations

22
times ranked

1250
citing authors

#	ARTICLE	IF	CITATIONS
1	Lactoferricin-inspired peptide AMC-109 augments the effect of ciprofloxacin against <i>Pseudomonas aeruginosa</i> biofilm in chronic murine wounds. <i>Journal of Global Antimicrobial Resistance</i> , 2022, 29, 185-193.	2.2	3
2	In vitro and in vivo antibacterial properties of peptide AMC-109 impregnated wound dressings and gels. <i>Journal of Antibiotics</i> , 2021, 74, 337-345.	2.0	13
3	Chronic lung disease in children and adolescents with HIV: a case-control study. <i>Tropical Medicine and International Health</i> , 2020, 25, 590-599.	2.3	12
4	Ribosome-Mediated Attenuation of <i>vga(A)</i> Expression Is Shaped by the Antibiotic Resistance Specificity of Vga(A) Protein Variants. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	3.2	17
5	Identification of surface proteins in a clinical <i>Staphylococcus haemolyticus</i> isolate by bacterial surface shaving. <i>BMC Microbiology</i> , 2020, 20, 80.	3.3	13
6	Effects of fecal microbiota transplantation in subjects with irritable bowel syndrome are mirrored by changes in gut microbiome. <i>Gut Microbes</i> , 2020, 12, 1794263.	9.8	31
7	<i>Staphylococcus borealis</i> sp. nov., isolated from human skin and blood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 6067-6078.	1.7	23
8	Comparative Genomic Analysis of <i>Staphylococcus haemolyticus</i> Reveals Key to Hospital Adaptation and Pathogenicity. <i>Frontiers in Microbiology</i> , 2019, 10, 2096.	3.5	33
9	Proteome profiling of secreted and membrane vesicle associated proteins of an invasive and a commensal <i>Staphylococcus haemolyticus</i> isolate. <i>Data in Brief</i> , 2019, 22, 914-919.	1.0	2
10	The synthetic antimicrobial peptide LTX 21 induces inflammatory responses in a human whole blood model and a murine peritoneum model. <i>Apmis</i> , 2019, 127, 475-483.	2.0	2
11	Comparative exoproteome profiling of an invasive and a commensal <i>Staphylococcus haemolyticus</i> isolate. <i>Journal of Proteomics</i> , 2019, 197, 106-114.	2.4	16
12	Fluorescence assay to predict activity of the glycopeptide antibiotics. <i>Journal of Antibiotics</i> , 2019, 72, 114-117.	2.0	13
13	Faecal microbiota transplantation versus placebo for moderate-to-severe irritable bowel syndrome: a double-blind, randomised, placebo-controlled, parallel-group, single-centre trial. <i>The Lancet Gastroenterology and Hepatology</i> , 2018, 3, 17-24.	8.1	325
14	Teicoplanin resistance in <i>Staphylococcus haemolyticus</i> is associated with mutations in histidine kinases VraS and Walk. <i>Diagnostic Microbiology and Infectious Disease</i> , 2018, 90, 233-240.	1.8	9
15	Effects of Probiotic Supplementation on the Gut Microbiota and Antibiotic Resistome Development in Preterm Infants. <i>Frontiers in Pediatrics</i> , 2018, 6, 347.	1.9	57
16	<i>Bifidobacterium</i> Bacteremia: Clinical Characteristics and a Genomic Approach To Assess Pathogenicity. <i>Journal of Clinical Microbiology</i> , 2017, 55, 2234-2248.	3.9	72
17	Coagulase-negative staphylococci in Southern Brazil: looking toward its high diversity. <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2016, 49, 292-299.	0.9	6
18	Antimicrobial susceptibility and body site distribution of community isolates of coagulase-negative staphylococci. <i>Apmis</i> , 2016, 124, 973-978.	2.0	18

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19	Whole-genome sequencing reveals clonal expansion of multiresistant <i>Staphylococcus haemolyticus</i> in European hospitals. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 2920-2927.	3.0	48
20	Efficacy of a synthetic antimicrobial peptidomimetic versus vancomycin in a <i>Staphylococcus epidermidis</i> device-related murine peritonitis model. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 2106-2110.	3.0	6
21	Core genome conservation of <i>Staphylococcus haemolyticus</i> limits sequence based population structure analysis. <i>Journal of Microbiological Methods</i> , 2012, 89, 159-166.	1.6	32