

Karim BEN SLAMA

List of Publications by Year in descending order

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Version: 2024-02-01

51
papers

1,926
citations

236925

25
h-index

265206

42
g-index

51
all docs

51
docs citations

51
times ranked

2392
citing authors

#	ARTICLE	IF	CITATIONS
1	Distribution, Diversity and Antibiotic Resistance of <i>Pseudomonas</i> spp. Isolated from the Water Dams in the North of Tunisia. <i>Current Microbiology</i> , 2022, 79, 188.	2.2	4
2	First Report of KPC-2 and KPC-3-Producing Enterobacteriaceae in Wild Birds in Africa. <i>Microbial Ecology</i> , 2020, 79, 30-37.	2.8	21
3	High prevalence of imipenem-resistant and metallo-beta-lactamase-producing <i>Pseudomonas aeruginosa</i> in the Burns Hospital in Tunisia: detection of a novel class 1 integron. <i>Journal of Chemotherapy</i> , 2019, 31, 120-126.	1.5	20
4	Extended-Spectrum β -Lactamases among Enterobacteriaceae Isolated from Urinary Tract Infections in Gaza Strip, Palestine. <i>BioMed Research International</i> , 2019, 2019, 1-11.	1.9	12
5	High diversity of genetic lineages and virulence genes of <i>Staphylococcus aureus</i> isolated from dairy products in Tunisia. <i>Annals of Microbiology</i> , 2019, 69, 73-78.	2.6	6
6	Detection of CTX-M-15 harboring <i>Escherichia coli</i> isolated from wild birds in Tunisia. <i>BMC Microbiology</i> , 2018, 18, 26.	3.3	20
7	Antimicrobial resistance and genetic lineages of faecal enterococci of wild birds: Emergence of vanA and vanB2 harbouring <i>Enterococcus faecalis</i> . <i>International Journal of Antimicrobial Agents</i> , 2018, 52, 936-941.	2.5	24
8	First Report of Extended-Spectrum β -Lactamases Among Clinical Isolates of <i>Klebsiella pneumoniae</i> in Gaza Strip, Palestine. <i>Microbial Drug Resistance</i> , 2017, 23, 169-176.	2.0	7
9	Characterization of two polyvalent phages infecting Enterobacteriaceae. <i>Scientific Reports</i> , 2017, 7, 40349.	3.3	115
10	Identification of LukPQ, a novel, equid-adapted leukocidin of <i>Staphylococcus aureus</i> . <i>Scientific Reports</i> , 2017, 7, 40660.	3.3	47
11	Antibiotic resistance in <i>Escherichia coli</i> in husbandry animals: the African perspective. <i>Letters in Applied Microbiology</i> , 2017, 64, 318-334.	2.2	119
12	Diversity of species and antibiotic resistance in enterococci isolated from seafood in Tunisia. <i>Annals of Microbiology</i> , 2017, 67, 135-141.	2.6	9
13	Species distribution, antibiotic resistance and virulence traits in canine and feline enterococci in Tunisia. <i>Acta Veterinaria Hungarica</i> , 2017, 65, 173-184.	0.5	19
14	Characterization of Five Podoviridae Phages Infecting <i>Citrobacter freundii</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 1023.	3.5	32
15	Prevalence, antimicrobial resistance and genetic lineages of <i>Enterococcus</i> spp. from vegetable food, soil and irrigation water in farm environments in Tunisia. <i>Journal of the Science of Food and Agriculture</i> , 2016, 96, 1627-1633.	3.5	48
16	Environmental <i>Staphylococcus aureus</i> contamination in a Tunisian hospital. <i>Journal of Chemotherapy</i> , 2016, 28, 506-509.	1.5	8
17	Characterization of extended-spectrum β -lactamase (ESBL)-producing <i>Klebsiella</i> , <i>Enterobacter</i> , and <i>Citrobacter</i> obtained in environmental samples of a Tunisian hospital. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 86, 190-193.	1.8	16
18	Antibiotic resistance and virulence of enterococci isolates from healthy humans in Tunisia. <i>Annals of Microbiology</i> , 2016, 66, 717-725.	2.6	7

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19	High prevalence of <i>Staphylococcus haemolyticus</i> and <i>Staphylococcus saprophyticus</i> in environmental samples of a Tunisian hospital. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 85, 136-140.	1.8	26
20	First report of extended-spectrum β -lactamases among clinical isolates of <i>Escherichia coli</i> in Gaza Strip, Palestine. <i>Journal of Global Antimicrobial Resistance</i> , 2016, 6, 17-21.	2.2	7
21	Characteristics of extended-spectrum β -lactamase (ESBL)- and pAmpC beta-lactamase-producing Enterobacteriaceae of water samples in Tunisia. <i>Science of the Total Environment</i> , 2016, 550, 1103-1109.	8.0	69
22	Detection of CTX-M-15-Producing <i>Escherichia coli</i> Isolates of Lineages ST131-B2 and ST167-A in Environmental Samples of a Tunisian Hospital. <i>Microbial Drug Resistance</i> , 2016, 22, 399-403.	2.0	17
23	Carbapenemases and extended-spectrum β -lactamases producing Enterobacteriaceae isolated from Tunisian and Libyan hospitals. <i>Journal of Infection in Developing Countries</i> , 2016, 10, 718-727.	1.2	36
24	Multidrug-resistant enterococci in the hospital environment: detection of novel vancomycin-resistant <i>E. faecium</i> clone ST910. <i>Journal of Infection in Developing Countries</i> , 2016, 10, 799-806.	1.2	19
25	Emergence of Carbapenem-Resistant <i>Pseudomonas aeruginosa</i> and <i>Acinetobacter baumannii</i> Clinical Isolates Collected from Some Libyan Hospitals. <i>Microbial Drug Resistance</i> , 2015, 21, 335-341.	2.0	39
26	Molecular Characterization of <i>Staphylococcus aureus</i> from Nasal Samples of Healthy Farm Animals and Pets in Tunisia. <i>Vector-Borne and Zoonotic Diseases</i> , 2015, 15, 109-115.	1.5	37
27	Diversity of enterococcal species and characterization of high-level aminoglycoside resistant enterococci of samples of wastewater and surface water in Tunisia. <i>Science of the Total Environment</i> , 2015, 530-531, 11-17.	8.0	28
28	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
29	Detection of extended-spectrum beta-lactamase (ESBL)-producing Enterobacteriaceae in vegetables, soil and water of the farm environment in Tunisia. <i>International Journal of Food Microbiology</i> , 2015, 203, 86-92.	4.7	111
30	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
31	Detection of CTX-M-15-producing <i>Escherichia coli</i> isolates of lineages ST410-A, ST617-A and ST354-D in faecal samples of hospitalized patients in a Mauritanian hospital. <i>Journal of Chemotherapy</i> , 2015, 27, 114-116.	1.5	20
32	Diversity of species and antibiotic resistance among fecal enterococci from wild birds in Tunisia. Detection of vanA-containing <i>Enterococcus faecium</i> isolates. <i>European Journal of Wildlife Research</i> , 2015, 61, 319-323.	1.4	23
33	Characterisation of nasal <i>Staphylococcus delphini</i> and <i>Staphylococcus pseudintermedius</i> isolates from healthy donkeys in Tunisia. <i>Equine Veterinary Journal</i> , 2015, 47, 463-466.	1.7	20
34	Antibiotic resistance and virulence of faecal enterococci isolated from food-producing animals in Tunisia. <i>Annals of Microbiology</i> , 2015, 65, 695-702.	2.6	21
35	vanA-containing <i>E. faecium</i> isolates of clonal complex CC17 in clinical and environmental samples in a Tunisian hospital. <i>Diagnostic Microbiology and Infectious Disease</i> , 2014, 79, 60-63.	1.8	19
36	Incl1 Plasmids Carrying <i>bla</i> _{CTX-M-1} or <i>bla</i> _{CMY-2} Genes in <i>Escherichia coli</i> from Healthy Humans and Animals in Tunisia. <i>Microbial Drug Resistance</i> , 2014, 20, 495-500.	2.0	66

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37	Antimicrobial Resistance, Virulence Genes, and Genetic Lineages of <i>Staphylococcus pseudintermedius</i> in Healthy Dogs in Tunisia. <i>Microbial Ecology</i> , 2013, 66, 363-368.	2.8	34
38	Species distribution, antibiotic resistance and virulence traits in enterococci from meat in Tunisia. <i>Meat Science</i> , 2013, 93, 675-680.	5.5	53
39	First Detection of CTX-M-1, CMY-2, and QnrB19 Resistance Mechanisms in Fecal <i>Escherichia coli</i> Isolates from Healthy Pets in Tunisia. <i>Vector-Borne and Zoonotic Diseases</i> , 2013, 13, 98-102.	1.5	36
40	Faecal enterococci from camels in Tunisia: species, antibiotic resistance and virulent genes. <i>Veterinary Record</i> , 2013, 172, 213-213.	0.3	8
41	Lineages and Virulence Gene Content among Extended-Spectrum β -Lactamase-Producing <i>Escherichia coli</i> Strains of Food Origin in Tunisia. <i>Journal of Food Protection</i> , 2013, 76, 323-327.	1.7	21
42	Genotypic Diversity, Antibiotic Resistance and Bacteriocin Production of Enterococci Isolated from Rhizospheres. <i>Microbes and Environments</i> , 2012, 27, 533-537.	1.6	10
43	High diversity of genetic lineages and virulence genes in nasal <i>Staphylococcus aureus</i> isolates from donkeys destined to food consumption in Tunisia with predominance of the ruminant associated CC133 lineage. <i>BMC Veterinary Research</i> , 2012, 8, 203.	1.9	42
44	Prevalence and Characterization of Extended-Spectrum Beta-Lactamase (ESBL) and CMY-2 Producing <i>Escherichia coli</i> Isolates from Healthy Food-Producing Animals in Tunisia. <i>Foodborne Pathogens and Disease</i> , 2012, 9, 1137-1142.	1.8	65
45	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
46	Nasal carriage of <i>Staphylococcus aureus</i> in healthy humans with different levels of contact with animals in Tunisia: genetic lineages, methicillin resistance, and virulence factors. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2011, 30, 499-508.	2.9	38
47	Diversity of Genetic Lineages Among CTX-M-15 and CTX-M-14 Producing <i>Escherichia coli</i> Strains in a Tunisian Hospital. <i>Current Microbiology</i> , 2011, 62, 1794-1801.	2.2	44
48	Prevalence of broad-spectrum cephalosporin-resistant <i>Escherichia coli</i> isolates in food samples in Tunisia, and characterization of integrons and antimicrobial resistance mechanisms implicated. <i>International Journal of Food Microbiology</i> , 2010, 137, 281-286.	4.7	62
49	Characterization of CTX-M and SHV extended-spectrum β -lactamases and associated resistance genes in <i>Escherichia coli</i> strains of food samples in Tunisia. <i>Journal of Antimicrobial Chemotherapy</i> , 2007, 60, 1137-1141.	3.0	170
50	Heterogeneity among infecting strains of <i>Pseudomonas aeruginosa</i> in diverse departments of a large Tunisian hospital. <i>Journal of Hospital Infection</i> , 2001, 47, 325-327.	2.9	7
51	Thuricin 7: a novel bacteriocin produced by <i>Bacillus thuringiensis</i> BMG1.7, a new strain isolated from soil. <i>Letters in Applied Microbiology</i> , 2001, 32, 243-247.	2.2	113