Karim BEN SLAMA

List of Publications by Year
in descending order

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	Characterization of CTX-M and SHV extended-spectrum Â-lactamases and associated resistance genes in Escherichia coli strains of food samples in Tunisia. Journal of Antimicrobial Chemotherapy, 2007, 60, 1137-1141.	3.0	170
2	Antibiotic resistance in Escherichia coli in husbandry animals: the African perspective. Letters in Applied Microbiology, 2017, 64, 318-334.	2.2	119
3	Characterization of two polyvalent phages infecting Enterobacteriaceae. Scientific Reports, 2017, 7, 40349.	3.3	115
4	Thuricin 7: a novel bacteriocin produced by Bacillus thuringiensis BMG1.7, a new strain isolated from soil. Letters in Applied Microbiology, 2001, 32, 243-247.	2.2	113
5	Detection of extended-spectrum beta-lactamase (ESBL)-producing Enterobacteriaceae in vegetables, soil and water of the farm environment in Tunisia. International Journal of Food Microbiology, 2015, 203, 86-92.	4.7	111
6	Prevalence, antibiotic resistance, virulence traits and genetic lineages of Staphylococcus aureus in healthy sheep in Tunisia. Veterinary Microbiology, 2012, 156, 367-373.	1.9	77
7	Characteristics of extended-spectrum Î²-lactamase (ESBL)- and pAmpC beta-lactamase-producing Enterobacteriaceae of water samples in Tunisia. Science of the Total Environment, 2016, 550, 1103-1109.	8.0	69
8	IncI1 Plasmids Carryingbla_{CTX-M-1}orbla_{CMY-2}Genes inEscherichia colifrom Healthy Humans and Animals in Tunisia. Microbial Drug Resistance, 2014, 20, 495-500.	2.0	66
9	Prevalence and Characterization of Extended-Spectrum Beta-Lactamase (ESBL)â€“ and CMY-2â€“ProducingEscherichia coliIsolates from Healthy Food-Producing Animals in Tunisia. Foodborne Pathogens and Disease, 2012, 9, 1137-1142.	1.8	65
10	Prevalence of broad-spectrum cephalosporin-resistant Escherichia coli isolates in food samples in Tunisia, and characterization of integrons and antimicrobial resistance mechanisms implicated. International Journal of Food Microbiology, 2010, 137, 281-286.	4.7	62
11	Species distribution, antibiotic resistance and virulence traits in enterococci from meat in Tunisia. Meat Science, 2013, 93, 675-680.	5.5	53
12	Prevalence, antimicrobial resistance and genetic lineages of Enterococcus spp. from vegetable food, soil and irrigation water in farm environments in Tunisia. Journal of the Science of Food and Agriculture, 2016, 96, 1627-1633.	3.5	48
13	Identification of LukPQ, a novel, equid-adapted leukocidin of Staphylococcus aureus. Scientific Reports, 2017, 7, 40660.	3.3	47
14	Diversity of Genetic Lineages Among CTX-M-15 and CTX-M-14 Producing Escherichia coli Strains in a Tunisian Hospital. Current Microbiology, 2011, 62, 1794-1801.	2.2	44
15	High diversity of genetic lineages and virulence genes in nasal Staphylococcus aureusisolates from donkeys destined to food consumption in Tunisia with predominance of the ruminant associated CC133 lineage. BMC Veterinary Research, 2012, 8, 203.	1.9	42
16	Emergence of Carbapenem-Resistant Pseudomonas aeruginosa and Acinetobacter baumannii Clinical Isolates Collected from Some Libyan Hospitals. Microbial Drug Resistance, 2015, 21, 335-341.	2.0	39
17	Characterization of Staphylococcus aureus from Raw Meat Samples in Tunisia: Detection of Clonal Lineage ST398 from the African Continent. Foodborne Pathogens and Disease, 2015, 12, 686-692.	1.8	39
18	Nasal carriage of Staphylococcus aureus in healthy humans with different levels of contact with animals in Tunisia: genetic lineages, methicillin resistance, and virulence factors. European Journal of Clinical Microbiology and Infectious Diseases, 2011, 30, 499-508.	2.9	38

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19	Molecular Characterization ofStaphylococcus aureusfrom Nasal Samples of Healthy Farm Animals and Pets in Tunisia. Vector-Borne and Zoonotic Diseases, 2015, 15, 109-115.	1.5	37
20	First Detection of CTX-M-1, CMY-2, and QnrB19 Resistance Mechanisms in FecalEscherichia coliIsolates from Healthy Pets in Tunisia. Vector-Borne and Zoonotic Diseases, 2013, 13, 98-102.	1.5	36
21	Carbapenemases and extended-spectrum Î²-lactamases producing Enterobacteriaceae isolated from Tunisian and Libyan hospitals. Journal of Infection in Developing Countries, 2016, 10, 718-727.	1.2	36
22	Antimicrobial Resistance, Virulence Genes, and Genetic Lineages of Staphylococcus pseudintermedius in Healthy Dogs in Tunisia. Microbial Ecology, 2013, 66, 363-368.	2.8	34
23	Characterization of Five Podoviridae Phages Infecting Citrobacter freundii. Frontiers in Microbiology, 2016, 7, 1023.	3.5	32
24	Diversity of enterococcal species and characterization of high-level aminoglycoside resistant enterococci of samples of wastewater and surface water in Tunisia. Science of the Total Environment, 2015, 530-531, 11-17.	8.0	28
25	High prevalence of Staphylococcus haemolyticus and Staphylococcus saprophyticus in environmental samples of a Tunisian hospital. Diagnostic Microbiology and Infectious Disease, 2016, 85, 136-140.	1.8	26
26	Antimicrobial resistance and genetic lineages of faecal enterococci of wild birds: Emergence of vanA and vanB2 harbouring Enterococcus faecalis. International Journal of Antimicrobial Agents, 2018, 52, 936-941.	2.5	24
27	Diversity of species and antibiotic resistance among fecal enterococci from wild birds in Tunisia. Detection of vanA-containing Enterococcus faecium isolates. European Journal of Wildlife Research, 2015, 61, 319-323.	1.4	23
28	Lineages and Virulence Gene Content among Extended-Spectrum Î²-Lactamase-Producing Escherichia coli Strains of Food Origin in Tunisia. Journal of Food Protection, 2013, 76, 323-327.	1.7	21
29	Antibiotic resistance and virulence of faecal enterococci isolated from food-producing animals in Tunisia. Annals of Microbiology, 2015, 65, 695-702.	2.6	21
30	First Report of KPC-2 and KPC-3-Producing Enterobacteriaceae in Wild Birds in Africa. Microbial Ecology, 2020, 79, 30-37.	2.8	21
31	Detection of CTX-M-15-producingEscherichia coliisolates of lineages ST410-A, ST617-A and ST354-D in faecal samples of hospitalized patients in a Mauritanian hospital. Journal of Chemotherapy, 2015, 27, 114-116.	1.5	20
32	Characterisation of nasal <scp>S</scp>taphylococcus delphini and <scp>S</scp>taphylococcus pseudintermedius isolates from healthy donkeys in <scp>T</scp>unisia. Equine Veterinary Journal, 2015, 47, 463-466.	1.7	20
33	Detection of CTX-M-15 harboring Escherichia coli isolated from wild birds in Tunisia. BMC Microbiology, 2018, 18, 26.	3.3	20
34	High prevalence of imipenem-resistant and metallo-beta-lactamase-producing Pseudomonas aeruginosa in the Burns Hospital in Tunisia: detection of a novel class 1 integron. Journal of Chemotherapy, 2019, 31, 120-126.	1.5	20
35	vanA-containing E. faecium isolates of clonal complex CC17 in clinical and environmental samples in a Tunisian hospital. Diagnostic Microbiology and Infectious Disease, 2014, 79, 60-63.	1.8	19
36	Species distribution, antibiotic resistance and virulence traits in canine and feline enterococci in Tunisia. Acta Veterinaria Hungarica, 2017, 65, 173-184.	0.5	19

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37	Multidrug-resistant enterococci in the hospital environment: detection of novel vancomycin-resistant E. faecium clone ST910. Journal of Infection in Developing Countries, 2016, 10, 799-806.	1.2	19
38	Detection of CTX-M-15-ProducingEscherichia coliIsolates of Lineages ST131-B2 and ST167-A in Environmental Samples of a Tunisian Hospital. Microbial Drug Resistance, 2016, 22, 399-403.	2.0	17
39	Characterization of extended-spectrum Î²-lactamase (ESBL)-producing Klebsiella, Enterobacter , and Citrobacter obtained in environmental samples of a Tunisian hospital. Diagnostic Microbiology and Infectious Disease, 2016, 86, 190-193.	1.8	16
40	Clonal lineages detected amongst tetracycline-resistant meticillin-resistant Staphylococcus aureus isolates of a Tunisian hospital, with detection of lineage ST398. Journal of Medical Microbiology, 2015, 64, 623-629.	1.8	15
41	Extended-SpectrumÎ²-Lactamases among Enterobacteriaceae Isolated from Urinary Tract Infections in Gaza Strip, Palestine. BioMed Research International, 2019, 2019, 1-11.	1.9	12
42	Genotypic Diversity, Antibiotic Resistance and Bacteriocin Production of Enterococci Isolated from Rhizospheres. Microbes and Environments, 2012, 27, 533-537.	1.6	10
43	Diversity of species and antibiotic resistance in enterococci isolated from seafood in Tunisia. Annals of Microbiology, 2017, 67, 135-141.	2.6	9
44	Faecal enterococci from camels in Tunisia: species, antibiotic resistance and virulent genes. Veterinary Record, 2013, 172, 213-213.	0.3	8
45	Environmental Staphylococcus aureus contamination in a Tunisian hospital. Journal of Chemotherapy, 2016, 28, 506-509.	1.5	8
46	Heterogeneity among infecting strains of Pseudomonas aeruginosa in diverse departments of a large Tunisian hospital. Journal of Hospital Infection, 2001, 47, 325-327.	2.9	7
47	Antibiotic resistance and virulence of enterococci isolates from healthy humans in Tunisia. Annals of Microbiology, 2016, 66, 717-725.	2.6	7
48	First report of extended-spectrum Î²-lactamases among clinical isolates of Escherichia coli in Gaza Strip, Palestine. Journal of Global Antimicrobial Resistance, 2016, 6, 17-21.	2.2	7
49	First Report of Extended-Spectrum Î²-Lactamases Among Clinical Isolates ofKlebsiella pneumoniaein Gaza Strip, Palestine. Microbial Drug Resistance, 2017, 23, 169-176.	2.0	7
50	High diversity of genetic lineages and virulence genes of Staphylococcus aureus isolated from dairy products in Tunisia. Annals of Microbiology, 2019, 69, 73-78.	2.6	6
51	Distribution, Diversity and Antibiotic Resistance of Pseudomonas spp. Isolated from the Water Dams in the North of Tunisia. Current Microbiology, 2022, 79, 188.	2.2	4