

Davood Kalantar-Neyestanaki

List of Publications by Year in descending order

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51
papers

529
citations

759233

12
h-index

752698

20
g-index

56
all docs

56
docs citations

56
times ranked

765
citing authors

#	ARTICLE	IF	CITATIONS
1	Determination of extended spectrum beta-lactamases, metallo-beta-lactamases and AmpC-beta-lactamases among carbapenem resistant <i>Pseudomonas aeruginosa</i> isolated from burn patients. <i>Burns</i> , 2014, 40, 1556-1561.	1.9	67
2	Characterization of AmpC, CTX-M and MBLs types of \hat{I}^2 -lactamases in clinical isolates of <i>Klebsiella pneumoniae</i> and <i>Escherichia coli</i> producing Extended Spectrum \hat{I}^2 -lactamases in Kerman, Iran. <i>Jundishapur Journal of Microbiology</i> , 2014, 7, e8756.	0.5	39
3	Endemic dissemination of different sequence types of carbapenem-resistant <i>Klebsiella pneumoniae</i> strains harboring <i>bla</i>_{<i>NDM</i>} and <i>16S</i> <i>rRNA methylase</i> genes in Kerman hospitals, Iran, from 2015 to 2017. <i>Infection and Drug Resistance</i> . 2019, Volume 12, 45-54.	2.7	32
4	Detection of AmpC- \hat{I}^2 -lactamases producing isolates among carbapenem resistant <i>P. aeruginosa</i> isolated from burn patient. <i>Iranian Journal of Microbiology</i> , 2014, 6, 306-10.	0.8	27
5	Association between virulence profile, biofilm formation and phylogenetic groups of <i>Escherichia coli</i> causing urinary tract infection and the commensal gut microbiota: A comparative analysis. <i>Microbial Pathogenesis</i> , 2017, 110, 540-545.	2.9	25
6	Virulence Factors, Capsular Serotypes and Antimicrobial Resistance of Hypervirulent <i>Klebsiella pneumoniae</i> and Classical <i>Klebsiella pneumoniae</i> in Southeast Iran. <i>Infection and Chemotherapy</i> , 2019, 51, .	2.3	24
7	Nanomaterials in the Management of Gram-Negative Bacterial Infections. <i>Nanomaterials</i> , 2021, 11, 2535.	4.1	23
8	Characterization of SCCmec, spa types and Multi Drug Resistant of methicillin-resistant <i>Staphylococcus aureus</i> isolates among inpatients and outpatients in a referral hospital in Shiraz, Iran. <i>BMC Research Notes</i> , 2019, 12, 614.	1.4	21
9	Characterization of SCCmec and spa types of methicillin-resistant <i>Staphylococcus aureus</i> isolates from health-care and community-acquired infections in Kerman, Iran. <i>Journal of Epidemiology and Global Health</i> , 2017, 7, 263.	2.9	20
10	Determination of carbapenem resistance mechanism in clinical isolates of <i>Pseudomonas aeruginosa</i> isolated from burn patients, in Tehran, Iran. <i>Journal of Epidemiology and Global Health</i> , 2017, 7, 155.	2.9	19
11	Emergence of co-existence of blaNDM with rmtC and qnrB genes in clinical carbapenem-resistant <i>Klebsiella pneumoniae</i> isolates in burning center from southeast of Iran. <i>Folia Microbiologica</i> , 2019, 64, 55-62.	2.3	19
12	Clonal relationships, antimicrobial susceptibilities, and molecular characterization of extended-spectrum beta-lactamase-producing <i>Escherichia coli</i> isolates from urinary tract infections and fecal samples in Southeast Iran. <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2018, 51, 44-51.	0.9	17
13	Molecular analysis of immune evasion cluster (IEC) genes and intercellular adhesion gene cluster (ICA) among methicillin-resistant and methicillin-sensitive isolates of. <i>Journal of Preventive Medicine and Hygiene</i> , 2017, 58, E308-E314.	0.9	13
14	Comparison five primer sets from different genome region of COVID-19 for detection of virus infection by conventional RT-PCR. <i>Iranian Journal of Microbiology</i> , 2020, 12, 185-193.	0.8	13
15	Distribution of Aminoglycoside-Modifying Enzymes and Molecular Analysis of the Coagulase Gene in Clinical Isolates of Methicillin-Resistant and Methicillin-Susceptible <i>Staphylococcus aureus</i>. <i>Microbial Drug Resistance</i> , 2019, 25, 47-53.	2.0	12
16	Clonal relation and antimicrobial resistance pattern of extended-spectrum \hat{I}^2 -lactamase- and AmpC \hat{I}^2 -lactamase-producing <i>Enterobacter</i> spp. isolated from different clinical samples in Tehran, Iran. <i>Revista Da Sociedade Brasileira De Medicina Tropical</i> , 2018, 51, 88-93.	0.9	11
17	The spa typing of methicillin-resistant <i>Staphylococcus aureus</i> isolates by High Resolution Melting (HRM) analysis. <i>Journal of Medical Microbiology</i> , 2017, 66, 1335-1337.	1.8	11
18	Identification of Extended-Spectrum \hat{I}^2 -Lactamase Genes and AmpC- \hat{I}^2 -Lactamase in Clinical Isolates of <i>Escherichia coli</i> Recovered from Patients with Urinary Tract Infections in Kerman, Iran. <i>Archives of Pediatric Infectious Diseases</i> , 2016, 5, .	0.3	9

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19	Investigation of the 5' flanking region and exon 3 polymorphisms of IGF-1 gene showed moderate association with semen quality in Sanjabi breed rams. <i>Theriogenology</i> , 2017, 104, 186-191.	2.1	8
20	Molecular Detection of Macrolide and Lincosamide-Resistance Genes in Clinical Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates from Kerman, Iran. <i>Archives of Pediatric Infectious Diseases</i> , 2016, 5, .	0.3	8
21	Inhibition of quorum sensing-controlled virulence factor production in by gall extracts. <i>Iranian Journal of Microbiology</i> , 2017, 9, 26-32.	0.8	8
22	Identification of g.170G>A and g.332G>A mutations in exon 3 of leptin gene (B c n l and C a i l) and their association with semen quality and testicular dimensions in Sanjabi rams. <i>Animal Reproduction Science</i> , 2017, 179, 49-56.	1.5	7
23	<p>Genetic characterization of two vancomycin-resistant &em> <i>Staphylococcus aureus</i> isolates in Kerman, Iran</p>. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 1869-1875.	2.7	7
24	Molecular identification of <i>Candida</i> species, assessment of the antifungal susceptibility and the genetic relationship of <i>Candida albicans</i> isolated from immunocompromised patients in Kerman, Iran. <i>Gene Reports</i> , 2019, 17, 100484.	0.8	7
25	The emergence of vancomycin-resistant <i>Staphylococcus aureus</i> in an intensive care unit in Kerman, Iran. <i>Wiener Medizinische Wochenschrift</i> , 2018, 168, 85-88.	1.1	6
26	Determining spa-type of methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) via high-resolution melting (HRM) analysis, Shiraz, Iran. <i>BMC Research Notes</i> , 2020, 13, 97.	1.4	6
27	Identification of <i>Candida</i> spp. isolated from oral mucosa in patients with leukemias and lymphomas in Iran. <i>Iranian Journal of Microbiology</i> , 0, , .	0.8	6
28	Molecular analysis and antimicrobial resistance pattern of distinct strains of <i>Pseudomonas aeruginosa</i> isolated from cystic fibrosis patients in Iran. <i>Iranian Journal of Microbiology</i> , 0, , .	0.8	6
29	Detection of methicillin-resistant (MRSA) in clinical samples of patients with external ocular infection. <i>Iranian Journal of Microbiology</i> , 2018, 10, 215-219.	0.8	6
30	Biofilm formation and molecular analysis of intercellular adhesion gene cluster (icaABCD) among <i>Staphylococcus aureus</i> strains isolated from children with adenoiditis. <i>Iranian Journal of Microbiology</i> , 2021, 13, 458-463.	0.8	5
31	Comparison of virulence genes and phylogenetic groups of <i>Escherichia coli</i> isolates from urinary tract infections and normal fecal flora. <i>Gene Reports</i> , 2020, 20, 100709.	0.8	5
32	ISPPu22, a novel insertion sequence in the oprD porin gene of a carbapenem-resistant <i>Pseudomonas aeruginosa</i> isolate from a burn patient in Tehran, Iran. <i>Iranian Journal of Microbiology</i> , 2015, 7, 247-50.	0.8	5
33	Molecular analysis and antimicrobial resistance pattern of distinct strains of isolated from cystic fibrosis patients in Iran. <i>Iranian Journal of Microbiology</i> , 2019, 11, 98-107.	0.8	5
34	Prevalence of plasmid-mediated quinolone resistance and ESBLs genes in <i>Escherichia coli</i> isolated from urinary tract infections and fecal samples in Southeast Iran. <i>Gene Reports</i> , 2019, 17, 100487.	0.8	4
35	Correlation Between hlyA and cnf1 Virulent Genes with Antibiotic Resistance and non-ESBLs <i>Escherichia coli</i> Isolates Collected from Patient with Urinary Tract Infections in Kerman, Iran. <i>Archives of Pediatric Infectious Diseases</i> , 2017, 5, .	0.3	4
36	Identification of spp. isolated from oral mucosa in patients with leukemias and lymphomas in Iran. <i>Iranian Journal of Microbiology</i> , 2019, 11, 114-119.	0.8	4

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37	Evaluation of chromosomally and acquired mechanisms of resistance to carbapenem antibiotics among clinical isolates of <i>Pseudomonas aeruginosa</i> in Kerman, Iran. <i>Gene Reports</i> , 2020, 21, 100918.	0.8	3
38	The Modified Hodge Test: Is it an appropriate method for detection of KPC enzyme or not?. <i>Iranian Journal of Microbiology</i> , 2015, 7, 123-4.	0.8	3
39	High prevalence of multi-drug resistant and different SCCmec types among coagulase-negative <i>Staphylococci</i> spp. collected from clinical samples and skin of healthcare workers in Kerman, Southeast Iran. <i>Gene Reports</i> , 2022, 26, 101428.	0.8	3
40	Determination of extended-spectrum $\hat{2}$ -lactamase producing and hybrid pathotypes of <i>Escherichia coli</i> isolates from diarrheic samples. <i>Gene Reports</i> , 2022, 27, 101583.	0.8	3
41	Determination of antibiotic resistance genes, immune evasion cluster and agr types among <i>Staphylococcus aureus</i> strains isolated from children with adenoiditis. <i>Gene Reports</i> , 2020, 21, 100875.	0.8	2
42	Determination of incompatibility group plasmids and copy number of the bla NDM-1 gene in carbapenem-resistant <i>Klebsiella pneumoniae</i> strains recovered from different hospitals in Kerman, Iran. <i>Journal of Medical Microbiology</i> , 2021, 70, .	1.8	2
43	Comparative Prevalence of blaCTX-M-15 Gene with Virulence Genes and Serotypes in <i>Klebsiella pneumoniae</i> . <i>Jundishapur Journal of Microbiology</i> , 2018, 11, .	0.5	2
44	First detection of insertion sequences ISpa1635 and IS1411 among non-carbapenemase producing strains of <i>Pseudomonas aeruginosa</i> in Kerman, Iran. <i>Gene Reports</i> , 2019, 15, 100373.	0.8	1
45	Dissemination of different sequence types lineages harboring among uropathogenic in Kerman, Iran. <i>Iranian Journal of Basic Medical Sciences</i> , 2020, 23, 1551-1557.	1.0	1
46	The Modified Hodge Test for identification of <i>Klebsiella pneumoniae</i> carbapenemase producing isolates. <i>Burns</i> , 2013, 39, 370-371.	1.9	0
47	Reply to: Molecular methods require for confirmation blaAIM (Adelaide imipenemase) producing <i>Pseudomonas aeruginosa</i> . <i>Burns</i> , 2014, 40, 1419-1420.	1.9	0
48	Reply to: Differentiation between KPC and IMP carbapenemase need phenotypic and genotypic methods. <i>Burns</i> , 2014, 40, 1242-1243.	1.9	0
49	The $\hat{2}$ -Lactamase Disk Test: A Modified Method to Detect Extended-Spectrum- $\hat{2}$ -Lactamases in Multidrug-Resistant <i>Escherichia coli</i> Isolates. <i>Archives of Clinical Infectious Diseases</i> , 2016, 12, .	0.2	0
50	Molecular Identification of <i>MefE</i> and <i>AmpC</i> Resistance Genes in ATCC Bacteria. <i>Avicenna Journal of Clinical Microbiology and Infection</i> , 2019, 6, 142-143.	0.4	0
51	Reducing Effect of Cloxacillin on Minimum Inhibitory Concentrations to Imipenem, Meropenem, Ceftazidime, and Cefepime in Carbapenem-resistant Isolates. <i>Yale Journal of Biology and Medicine</i> , 2020, 93, 29-34.	0.2	0