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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	High prevalence of multi-drug resistant and different SCCmec types among coagulase-negative Staphylococci spp. collected from clinical samples and skin of healthcare workers in Kerman, Southeast Iran. Gene Reports, 2022, 26, 101428.	0.8	3
2	Determination of extended-spectrum β-lactamase producing and hybrid pathotypes of Escherichia coli isolates from diarrheic samples. Gene Reports, 2022, 27, 101583.	0.8	3
3	Determination of incompatibility group plasmids and copy number of the bla NDM-1 gene in carbapenem-resistant Klebsiella pneumoniae strains recovered from different hospitals in Kerman, Iran. Journal of Medical Microbiology, 2021, 70, .	1.8	2
4	Biofilm formation and molecular analysis of intercellular adhesion gene cluster (icaABCD) among Staphylococcus aureus strains isolated from children with adenoiditis. Iranian Journal of Microbiology, 2021, 13, 458-463.	0.8	5
5	Nanomaterials in the Management of Gram-Negative Bacterial Infections. Nanomaterials, 2021, 11, 2535.	4.1	23
6	Determination of antibiotic resistance genes, immune evasion cluster and agr types among Staphylococcus aureus strains isolated from children with adenoiditis. Gene Reports, 2020, 21, 100875.	0.8	2
7	Evaluation of chromosomally and acquired mechanisms of resistance to carbapenem antibiotics among clinical isolates of Pseudomonas aeruginosa in Kerman, Iran. Gene Reports, 2020, 21, 100918.	0.8	3
8	Determining spa-type of methicillin-resistant Staphylococcus aureus (MRSA) via high-resolution melting (HRM) analysis, Shiraz, Iran. BMC Research Notes, 2020, 13, 97.	1.4	6
9	Comparison of virulence genes and phylogenetic groups of Escherichia coli isolates from urinary tract infections and normal fecal flora. Gene Reports, 2020, 20, 100709.	0.8	5
10	Reducing Effect of Cloxacillin on Minimum Inhibitory Concentrations to Imipenem, Meropenem, Ceftazidime, and Cefepime in Carbapenem-resistant Isolates. Yale Journal of Biology and Medicine, 2020, 93, 29-34.	0.2	0
11	Comparison five primer sets from different genome region of COVID-19 for detection of virus infection by conventional RT-PCR. Iranian Journal of Microbiology, 2020, 12, 185-193.	0.8	13
12	Dissemination of different sequence types lineages harboring among uropathogenic in Kerman, Iran. Iranian Journal of Basic Medical Sciences, 2020, 23, 1551-1557.	1.0	1
13	Endemic dissemination of different sequence types of carbapenem-resistant Klebsiella pneumoniae strains harboring bla _{NDM} and 16S rRNA methylase genes in Kerman hospitals, Iran, from 2015 to 2017. Infection and Drug Resistance, 2019. Volume 12, 45:54	2.7	32
14	Emergence of co-existence of blaNDM with rmtC and qnrB genes in clinical carbapenem-resistant Klebsiella pneumoniae isolates in burning center from southeast of Iran. Folia Microbiologica, 2019, 64, 55-62.	2.3	19
15	<p>Genetic characterization of two vancomycin-resistant Staphylococcus aureus isolates in Kerman, Iran</p> . Infection and Drug Resistance, 2019, Volume 12, 1869-1875.	2.7	7
16	Characterization of SCCmec, spa types and Multi Drug Resistant of methicillin-resistant Staphylococcus aureus isolates among inpatients and outpatients in a referral hospital in Shiraz, Iran. BMC Research Notes, 2019, 12, 614.	1.4	21
17	Molecular identification of Candida species, assessment of the antifungal susceptibility and the genetic relationship of Candida albicans isolated from immunocompromised patients in Kerman, Iran. Gene Reports, 2019, 17, 100484.	0.8	7
18	Prevalence of plasmid-mediated quinolone resistance and ESBLs genes in Escherichia coli isolated from urinary tract infections and fecal samples in Southeast Iran. Gene Reports, 2019, 17, 100487.	0.8	4

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19	First detection of insertion sequences ISpa1635 and IS1411 among non-carbapenemase producing strains of Pseudomonas aeruginosa in Kerman, Iran. Gene Reports, 2019, 15, 100373.	0.8	1
20	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> . Microbial Drug Resistance, 2019, 25, 47-53.	2.0	12
21	Virulence Factors, Capsular Serotypes and Antimicrobial Resistance of Hypervirulent <i>Klebsiella pneumoniae</i> and Classical <i>Klebsiella pneumoniae</i> in Southeast Iran. Infection and Chemotherapy, 2019, 51, .	2.3	24
22	Molecular Identification of <i>MefE </i> and <i>AmpC </i> Resistance Genes in ATCC Bacteria. Avicenna Journal of Clinical Microbiology and Infection, 2019, 6, 142-143.	0.4	0
23	Identification of spp. isolated from oral mucosa in patients with leukemias and lymphomas in Iran. Iranian Journal of Microbiology, 2019, 11, 114-119.	0.8	4
24	Molecular analysis and antimicrobial resistance pattern of distinct strains of isolated from cystic fibrosis patients in Iran. Iranian Journal of Microbiology, 2019, 11, 98-107.	0.8	5
25	The emergence of vancomycin-resistant Staphylococcus aureus in an intensive care unit in Kerman, Iran. Wiener Medizinische Wochenschrift, 2018, 168, 85-88.	1.1	6
26	Clonal relation and antimicrobial resistance pattern of extended-spectrum β-lactamase- and AmpC β-lactamase-producing Enterobacter spp. isolated from different clinical samples in Tehran, Iran. Revista Da Sociedade Brasileira De Medicina Tropical, 2018, 51, 88-93.	0.9	11
27	Clonal relationships, antimicrobial susceptibilities, and molecular characterization of extended-spectrum beta-lactamase-producing Escherichia coli isolates from urinary tract infections and fecal samples in Southeast Iran. Revista Da Sociedade Brasileira De Medicina Tropical, 2018, 51, 44-51.	0.9	17
28	Comparative Prevalence of blaCTX-M-15 Gene with Virulence Genes and Serotypes in Klebsiella pneumoniae. Jundishapur Journal of Microbiology, 2018, 11, .	0.5	2
29	Detection of methicillin-resistant (MRSA) in clinical samples of patients with external ocular infection. Iranian Journal of Microbiology, 2018, 10, 215-219.	0.8	6
30	ldentification of g.170G>A and g.332G>A mutations in exon 3 of leptin gene (B cn l and C ai l) and their association with semen quality and testicular dimensions in Sanjabi rams. Animal Reproduction Science, 2017, 179, 49-56.	1.5	7
31	Determination of carbapenem resistance mechanism in clinical isolates of Pseudomonas aeruginosa isolated from burn patients, in Tehran, Iran. Journal of Epidemiology and Global Health, 2017, 7, 155.	2.9	19
32	Characterization of SCCmec and spa types of methicillin-resistant Staphylococcus aureus isolates from health-care and community-acquired infections in Kerman, Iran. Journal of Epidemiology and Global Health, 2017, 7, 263.	2.9	20
33	Investigation of the 5′ flanking region and exon 3 polymorphisms of IGF-1 gene showed moderate association with semen quality in Sanjabi breed rams. Theriogenology, 2017, 104, 186-191.	2.1	8
34	Association between virulence profile, biofilm formation and phylogenetic groups of Escherichia coli causing urinary tract infection and the commensal gut microbiota: A comparative analysis. Microbial Pathogenesis, 2017, 110, 540-545.	2.9	25
35	The spa typing of methicillin-resistant Staphylococcus aureus isolates by High Resolution Melting (HRM) analysis. Journal of Medical Microbiology, 2017, 66, 1335-1337.	1.8	11
36	Molecular analysis of immune evasion cluster (IEC) genes and intercellular adhesion gene cluster (ICA) among methicillin-resistant and methicillin-sensitive isolates of. Journal of Preventive Medicine and Hygiene, 2017, 58, E308-E314.	0.9	13

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IF # ARTICLE CITATIONS Correlation Between hlyA and cnf1 Virulent Genes with Antibiotic Resistance and non-ESBLs Escherichia coli Isolates Collected from Patient with Urinary Tract Infections in Kerman, Iran. Archives of Pediatric Infectious Diseases, 2017, 5, . Inhibition of quorum sensing-controlled virulence factor production in by gall extracts. Iranian 38 0.8 8 Journal of Microbiology, 2017, 9, 26-32. Molecular Detection of Macrolide and Lincosamide-Resistance Genes in Clinical Methicillin-Resistant 0.3 Staphylococcus aureus Isolates from Kerman, Iran. Archives of Pediatric Infectious Diseases, 2016, 5, . Identification of Extended-Spectrum Î²-Lactamase Genes and AmpC-Î²-Lactamase in Clinical Isolates of Escherichia coli Recovered from Patients with Urinary Tract Infections in Kerman, Iran. Archives of 40 0.3 9 Pediatric Infectious Diseases, 2016, 5, . The Î2-Lactamase Disk Test: A Modified Method to Detect Extended-Spectrum-Î2-Lactamases in 0.2 Multidrug-Resistant Escherichia coli Isolates. Archives of Clinical Infectious Diseases, 2016, 12, . The Modified Hodge Test: Is it an appropriate method for detection of KPC enzyme or not?. Iranian 42 0.8 3 Journal of Microbiology, 2015, 7, 123-4. ISPpu22, a novel insertion sequence in the oprD porin gene of a carbapenem-resistant Pseudomonas 0.8 aeruginosa isolate from a burn patient in Tehran, Iran. Iranian Journal of Microbiology, 2015, 7, 247-50. Characterization of AmpC, CTX-M and MBLs types of Î²-lactamases in clinical isolates of Klebsiella pneumoniae and Escherichia coli producing Extended Spectrum Î²-lactamases in Kerman, Iran. 44 0.5 39 Jundishapur Journal of Microbiology, 2014, 7, e8756. Determination of extended spectrum beta-lactamases, metallo-beta-lactamases and AmpC-beta-lactamases among carbapenem resistant Pseudomonas aeruginosa isolated from burn patients. Burns, 2014, 40, 1556-1561. 1.9 Reply to: Molecular methods require for confirmation blaAIM (Adelaide imipenemase) producing 46 1.9 0 Pseudomonas aeruginosa. Burns, 2014, 40, 1419-1420. Reply to: Differentiation between KPC and IMP carbapenemase need phenotypic and genotypic methods. Burns, 2014, 40, 1242-1243. Detection of AmpC-Î²-lactamases producing isolates among carbapenem resistant P. aeruginosa isolated 48 0.8 27 from burn patient. Iranian Journal of Microbiology, 2014, 6, 306-10. The Modified Hodge Test for identification of Klebsiella pneumoniae carbapenemase producing 1.9 isolates. Burns, 2013, 39, 370-371. Identification of Candida spp. isolated from oral mucosa in patients with leukemias and lymphomas in 50 0.8 6 Iran. Iranian Journal of Microbiology, 0, , . Molecular analysis and antimicrobial resistance pattern of distinct strains of Pseudomonas aeruginosa isolated from cystic fibrosis patients in Iran. Iranian Journal of Microbiology, 0, , .