Fereshteh Jabalameli

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Multiple-locus variable-number tandem repeat analysis for genotyping of erythromycin-resistant group B streptococci in Iran. New Microbes and New Infections, 2022, 45, 100957.	0.8	0
2	Multiplex detection of five common respiratory pathogens from bronchoalveolar lavages using high resolution melting curve analysis. BMC Microbiology, 2022, 22, 141.	1.3	2
3	Phenotypic and Genotypic Prevalence of Extended-Spectrum β-Lactamase-Producing <i>Escherichia coli</i> : A Systematic Review and Meta-Analysis in Iran. Microbial Drug Resistance, 2021, 27, 73-86.	0.9	9
4	Efficacy of 16S rRNA variable regions high-resolution melt analysis for bacterial pathogens identification in periprosthetic joint infections. BMC Microbiology, 2021, 21, 112.	1.3	4
5	A More Positive Culture by Resin-containing Media Usage after Suspicious Arthroscopic Infections in Patients Receiving Antimicrobial Therapy. Archives of Bone and Joint Surgery, 2021, 9, 496-502.	0.1	0
6	Prevalence of Genes Encoding Aminoglycoside-Modifying Enzymes in Clinical Isolates of Gram-Positive Cocci in Iran: A Systematic Review and Meta-Analysis. Microbial Drug Resistance, 2020, 26, 126-135.	0.9	2
7	Molecular characterization, antibiotic resistance pattern and capsular types of invasive Streptococcus pneumoniae isolated from clinical samples in Tehran, Iran. BMC Microbiology, 2020, 20, 167.	1.3	11
8	Combinatorial effects of antibiotics and enzymes against dual-species Staphylococcus aureus and Pseudomonas aeruginosa biofilms in the wound-like medium. PLoS ONE, 2020, 15, e0235093.	1.1	35
9	Antimicrobial resistance pattern, virulence determinants and molecular analysis of Enterococcus faecium isolated from children infections in Iran. BMC Microbiology, 2019, 19, 156.	1.3	14
10	Status of carbapenem-resistant Acinetobacter baumannii harboring carbapenemase: First systematic review and meta-analysis from Iran. Infection, Genetics and Evolution, 2019, 73, 433-443.	1.0	18
11	Prevalence of extended-spectrum β-lactamase-producing Klebsiella pneumoniae: First systematic review and meta-analysis from Iran. Journal of Global Antimicrobial Resistance, 2019, 18, 12-21.	0.9	16
12	High prevalence of direct repeat unit types of 10di, 8 h and 8i among methicillin resistant Staphylococcus aureus strains with staphylococcal cassette chromosome mec type IIIA isolated in Tehran, Iran. Antimicrobial Resistance and Infection Control, 2019, 8, 50.	1.5	8
13	The efficacy of lyticase and β-glucosidase enzymes on biofilm degradation of Pseudomonas aeruginosa strains with different gene profiles. BMC Microbiology, 2019, 19, 291.	1.3	20
14	Cytotoxicity Evaluation of Minimum Antibacterial Values of Different Medicaments Used in Endodontic Regenerative Procedures. European Journal of Dentistry, 2019, 13, 514-520.	0.8	13
15	Molecular Epidemiology and Drug Resistance Pattern of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> Isolates from Iran. Microbial Drug Resistance, 2019, 25, 336-343.	0.9	36
16	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
17	Prevalence of methicillin-resistant Staphylococcus aureus isolated from burn patients in Iran: A systematic review and meta-analysis. Journal of Global Antimicrobial Resistance, 2018, 12, 202-206.	0.9	21
18	Prevalence of Group B Streptococcus in Pregnant Women in Iran. Pediatric Infectious Disease Journal, 2018, 37, 186-190.	1.1	8

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19	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.	0.4	11
20	Prevalence of metallo-β-lactamase-encoding genes among carbapenem-resistant Pseudomonas aeruginosa strains isolated from burn patients in Iran. Revista Da Sociedade Brasileira De Medicina Tropical, 2018, 51, 270-276.	0.4	21
21	High diversity in SCCmec elements among multidrug-resistant Staphylococcus haemolyticus strains originating from paediatric patients; characterization of a new composite island. Journal of Medical Microbiology, 2018, 67, 915-921.	0.7	13
22	Assessment of disinfectant and antibiotic susceptibility patterns and multi-locus variable number tandem repeat analysis of isolated from blood cultures. Iranian Journal of Microbiology, 2018, 10, 90-97.	0.8	2
23	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.	1.1	19
24	Characterization of biofilm formation, antimicrobial resistance, and staphylococcal cassette chromosome mec analysis of methicillin resistant Staphylococcus hominis from blood cultures of children. Revista Da Sociedade Brasileira De Medicina Tropical, 2017, 50, 329-333.	0.4	7
25	Nasal carriage rate of methicillin resistant Staphylococcus aureus among Iranian healthcare workers: a systematic review and meta-analysis. Revista Da Sociedade Brasileira De Medicina Tropical, 2017, 50, 590-597.	0.4	22
26	The Effects of Berberine and Palmatine on Efflux Pumps Inhibition with Different Gene Patterns in Isolated from Burn Infections. Avicenna Journal of Medical Biotechnology, 2017, 9, 2-7.	0.2	26
27	Monoterpene isolated from the essential oil of Trachyspermum ammi is cytotoxic to multidrug-resistant Pseudomonas aeruginosa and Staphylococcus aureus strains. Revista Da Sociedade Brasileira De Medicina Tropical, 2016, 49, 172-176.	0.4	15
28	Evaluation of Mannosidase and Trypsin Enzymes Effects on Biofilm Production of Pseudomonas aeruginosa Isolated from Burn Wound Infections. PLoS ONE, 2016, 11, e0164622.	1.1	60
29	Spreading of genes encoding enterotoxins, haemolysins, adhesin and biofilm among methicillin resistant Staphylococcus aureus strains with staphylococcal cassette chromosome mec type IIIA isolated from burn patients. Microbial Pathogenesis, 2016, 97, 34-37.	1.3	25
30	Characterization of virulence factors, antimicrobial resistance pattern and clonal complexes of group B streptococci isolated from neonates. Microbial Pathogenesis, 2016, 99, 119-122.	1.3	17
31	Prevalence of vancomycin-resistant Enterococcus in Iran: a systematic review and meta-analysis. European Journal of Clinical Microbiology and Infectious Diseases, 2016, 35, 1387-1392.	1.3	46
32	Investigation of biofilm formation ability, antimicrobial resistance and the staphylococcal cassette chromosome mec patterns of methicillin resistant Staphylococcus epidermidis with different sequence types isolated from children. Microbial Pathogenesis, 2016, 93, 126-130.	1.3	28
33	Comparison of virulence factors and capsular types of Streptococcus agalactiae isolated from human and bovine infections. Microbial Pathogenesis, 2016, 91, 1-4.	1.3	22
34	Virulence factors, antimicrobial resistance pattern and molecular analysis of Enterococcal strains isolated from burn patients. Microbial Pathogenesis, 2016, 90, 93-97.	1.3	49
35	Variable number of tandem repeat profiles and antimicrobial resistance patterns of Staphylococcus haemolyticus strains isolated from blood cultures in children. Infection, Genetics and Evolution, 2016, 38, 19-21.	1.0	5
36	Prevalence of Panton-Valentine leucocidin and phenotypic and genotypic characterization of biofilm formation among Staphylococcus aureus strains isolated from children with adenoid hypertrophy. Microbial Pathogenesis, 2015, 89, 150-153.	1.3	8

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37	Comparison of virulence factors and biofilm formation among Staphylococcus aureus strains isolated from human and bovine infections. Microbial Pathogenesis, 2015, 88, 73-77.	1.3	49
38	Molecular investigation of Staphylococcus aureus isolated from the patients, personnel, air and environment of an ICU in a hospital in Tehran. Journal of Infection and Public Health, 2015, 8, 202-206.	1.9	28
39	Prevalence of human papillomavirus in oral lichen planus in an Iranian cohort. Journal of Oral and Maxillofacial Pathology, 2015, 19, 170.	0.3	17
40	ISPpu22, a novel insertion sequence in the oprD porin gene of a carbapenem-resistant Pseudomonas aeruginosa isolate from a burn patient in Tehran, Iran. Iranian Journal of Microbiology, 2015, 7, 247-50.	0.8	5
41	Virulence factors, antimicrobial susceptibility and molecular characterization of Streptococcus agalactiae isolated from pregnant women. Acta Microbiologica Et Immunologica Hungarica, 2014, 61, 425-434.	0.4	29
42	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.	1.1	67
43	Reply to: Molecular methods require for confirmation blaAIM (Adelaide imipenemase) producing Pseudomonas aeruginosa. Burns, 2014, 40, 1419-1420.	1.1	Ο
44	Reply to: Differentiation between KPC and IMP carbapenemase need phenotypic and genotypic methods. Burns, 2014, 40, 1242-1243.	1.1	0
45	High Incidence of Macrolide and Tetracycline Resistance among Streptococcus Agalactiae Strains Isolated from Clinical Samples in Tehran, Iran. Mædica, 2014, 9, 157-61.	0.4	10
46	Detection of AmpC-β-lactamases producing isolates among carbapenem resistant P. aeruginosa isolated from burn patient. Iranian Journal of Microbiology, 2014, 6, 306-10.	0.8	27
47	The Modified Hodge Test for identification of Klebsiella pneumoniae carbapenemase producing isolates. Burns, 2013, 39, 370-371.	1.1	Ο
48	Genetic similarity between adenoid tissue and middle ear fluid isolates of Streptococcus pneumoniae, Haemophilus influenzae and Moraxella catarrhalis from Iranian children with otitis media with effusion. International Journal of Pediatric Otorhinolaryngology, 2013, 77, 1841-1845.	0.4	10
49	Molecular analysis of typical and atypical enteropathogenic Escherichia coli (EPEC) isolated from children with diarrhoea. Journal of Medical Microbiology, 2013, 62, 191-195.	0.7	30
50	Distribution of bacterial contamination in a teaching hospital in Tehran — A special focus on Staphylococcus aureus. Acta Microbiologica Et Immunologica Hungarica, 2012, 59, 1-11.	0.4	10
51	Characterization of Alloiococcus otitidis strains isolated from children with otitis media with effusion by Pulsed-Field Gel Electrophoresis. International Journal of Pediatric Otorhinolaryngology, 2012, 76, 1658-1660.	0.4	2
52	Frequency of Alloicoccus otitidis, Streptococcus pneumoniae, Moraxella catarrhalis and Haemophilus influenzae in children with otitis media with effusion (OME) in Iranian patients. Auris Nasus Larynx, 2012, 39, 369-373.	0.5	22
53	A high prevalence of mupirocin and macrolide resistance determinant among Staphylococcus aureus strains isolated from burnt patients. Burns, 2012, 38, 378-382.	1.1	45
54	Evaluation of biofilm production and characterization of genes encoding type III secretion system among Pseudomonas aeruginosa isolated from burn patients. Burns, 2012, 38, 1192-1197.	1.1	69

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55	Multiple-locus variable number of tandem repeats (VNTR) fingerprinting (MLVF) and antibacterial resistance profiles of extended spectrum beta lactamase (ESBL) producing Pseudomonas aeruginosa among burnt patients in Tehran. Burns, 2011, 37, 1202-1207.	1.1	30
56	Molecular characterization of Staphylococcus aureus isolated from children with adenoid hypertrophy: Emergence of new spa types t7685 and t7692. International Journal of Pediatric Otorhinolaryngology, 2011, 75, 1446-1449.	0.4	18
57	Phenotypic and genotypic evaluation of fluoroquinolone resistance in clinical isolates of Staphylococcus aureus in Tehran. Medical Science Monitor, 2011, 17, PH71-PH74.	0.5	5
58	Time-kill study and synergistic activity of cell-wall inhibitor antibiotics in combination with gentamicin against Enterococcus faecalis and Enterococcus faecium. Acta Microbiologica Et Immunologica Hungarica, 2011, 58, 219-226.	0.4	7
59	Molecular analysis and antimicrobial susceptibility of methicillin resistant Staphylococcus aureus in one of the hospitals of Tehran University of Medical Sciences: High prevalence of sequence type 239 (ST239) clone. Acta Microbiologica Et Immunologica Hungarica, 2011, 58, 31-39.	0.4	21
60	Multiple-Locus Variable Number of Tandem Repeats Fingerprinting (MLVF) and Virulence Factor Analysis of Methicillin Resistant Staphylococcus aureus SCCmec type III. Polish Journal of Microbiology, 2011, 60, 303-307.	0.6	10
61	Detection of VEB-1, OXA-10 and PER-1 genotypes in extended-spectrum β-lactamase-producing Pseudomonas aeruginosa strains isolated from burn patients. Burns, 2010, 36, 70-74.	1.1	98
62	Characterization of Phenotypic and Genotypic inducible Macrolide Resistance in Staphylococciin Tehran, Iran. Journal of Chemotherapy, 2009, 21, 595-597.	0.7	13
63	Molecular characterization of Staphylococcus aureus isolated from bovine mastitis in Iran. Veterinary Microbiology, 2009, 139, 207-208.	0.8	3
64	Characterisation of genes encoding aminoglycoside-modifying enzymes among meticillin-resistant Staphylococcus aureus isolated from two hospitals in Tehran, Iran. International Journal of Antimicrobial Agents, 2009, 33, 264-265.	1.1	27
65	Phenotypic and Genotypic Evaluation of Aminoglycoside Resistance in Clinical Isolates of Staphylococci in Tehran, Iran. Microbial Drug Resistance, 2009, 15, 129-132.	0.9	30
66	Relationship between human cytomegalovirus transcription and symptomatic apical periodontitis in Iran. Oral Microbiology and Immunology, 2008, 23, 510-514.	2.8	20
67	Emergence of High-Level Vancomycin-Resistant <i>Staphylococcus aureus</i> in the Imam Khomeini Hospital in Tehran. Medical Principles and Practice, 2008, 17, 432-434.	1.1	93
68	Isolation of vancomycin-resistant Staphylococcus aureus in a teaching hospital in Tehran. Journal of Hospital Infection, 2007, 66, 92-93.	1.4	24