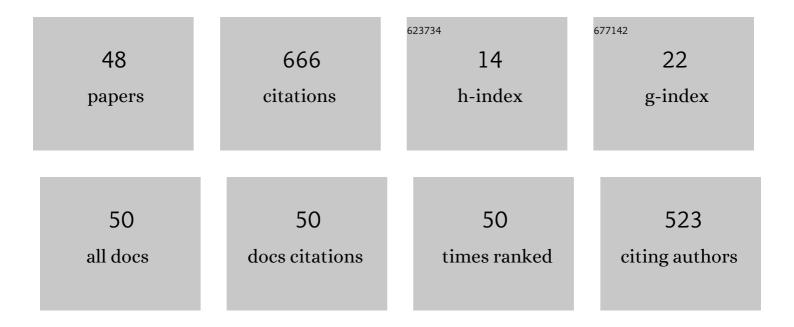
Morteza Saki

List of Publications by Year in descending order

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MODTEZA SAKI

#	Article	IF	CITATIONS
1	A review on mechanism of action, resistance, synergism, and clinical implications of mupirocin against Staphylococcus aureus. Biomedicine and Pharmacotherapy, 2019, 109, 1809-1818.	5.6	75
2	Black soldier fly larvae (BSFL) and their affinity for organic waste processing. Waste Management, 2022, 140, 1-13.	7.4	75
3	Isolation and identification of potential probiotic Lactobacillus species from feces of infants in southwest Iran. International Journal of Infectious Diseases, 2020, 96, 524-530.	3.3	33
4	Antibiotic resistance pattern and molecular characterization of extended-spectrum β-lactamase producing enteroaggregative Escherichia coli isolates in children from southwest Iran. Infection and Drug Resistance, 2018, Volume 11, 1097-1104.	2.7	30
5	Antibiotic resistance, biofilm production ability and genetic diversity of carbapenem-resistant Pseudomonas aeruginosa strains isolated from nosocomial infections in southwestern Iran. Molecular Biology Reports, 2022, 49, 3811-3822.	2.3	28
6	<p>Prevalence and antimicrobial resistance of Shigella species isolated from diarrheal patients in Ahvaz, southwest Iran</p> . Infection and Drug Resistance, 2019, Volume 12, 249-253.	2.7	27
7	Evaluation of the phytoconstituents of Auricularia auricula-judae mushroom and antimicrobial activity of its protein extract. European Journal of Integrative Medicine, 2020, 38, 101176.	1.7	26
8	Preliminary survey of extended-spectrum β-lactamases (ESBLs) in nosocomial uropathogen Klebsiella pneumoniae in north-central Iran. Heliyon, 2019, 5, e02349.	3.2	24
9	Prevalence of CRISPR-Cas Systems and Their Possible Association with Antibiotic Resistance in Enterococcus faecalis and Enterococcus faecium Collected from Hospital Wastewater. Infection and Drug Resistance, 2022, Volume 15, 1143-1154.	2.7	24
10	Investigation of SCC <i>mec</i> types I–IV in clinical isolates of methicillin-resistant coagulase-negative staphylococci in Ahvaz, Southwest Iran. Bioscience Reports, 2020, 40, .	2.4	21
11	Molecular Investigation of Staphylococcus aureus, coa and spa Genes in Ahvaz Hospitals, Staff Nose Compared With Patients Clinical Samples. Jundishapur Journal of Microbiology, 0, , .	0.5	19
12	Resistotyping and extended-spectrum beta-lactamase genes among Escherichia coli from wastewater treatment plants and recipient surface water for reuse in South Africa. New Microbes and New Infections, 2020, 38, 100803.	1.6	19
13	In vitro antibacterial properties of Cinnamomum zeylanicum essential oil against clinical extensively drug-resistant bacteria. European Journal of Integrative Medicine, 2020, 37, 101146.	1.7	19
14	<p>Prevalence of Extended-Spectrum Beta-Lactamase-Producing Enterobacteriaceae Causing Bloodstream Infections in Cancer Patients from Southwest of Iran</p> . Infection and Drug Resistance, 2020, Volume 13, 1319-1326.	2.7	19
15	Distribution of genes encoding resistance to macrolides, lincosamides, and streptogramins among methicillin-resistant Staphylococcus aureus strains isolated from burn patients. Acta Microbiologica Et Immunologica Hungarica, 2019, 66, 387-398.	0.8	18
16	Emergence of multidrug-resistant Shigella species harboring extended-spectrum beta-lactamase genes in pediatric patients with diarrhea from southwest of Iran. Molecular Biology Reports, 2020, 47, 7097-7106.	2.3	17
17	Detection of sul1 and sul2 genes in sulfonamide-resistant bacteria (SRB) from sewage, aquaculture sources, animal wastes and hospital wastewater in South-West Nigeria. Gene Reports, 2020, 20, 100742.	0.8	16
18	Lack of Association of Mouse Mammary Tumor Virus-Like Sequences in Iranian Breast Cancer Patients. Medical Principles and Practice, 2012, 21, 244-248.	2.4	14

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19	Distribution and antibiotic-resistance of different <i>Staphylococcus</i> species identified by matrix assisted laser desorption ionization-time of flight mass spectrometry (MALDI-TOF MS) isolated from the oral cavity. Journal of Oral Microbiology, 2021, 13, 1983322.	2.7	13
20	Metallo-β-lactamase and AmpC genes in Escherichia coli, Klebsiella pneumoniae, and Pseudomonas aeruginosa isolates from abattoir and poultry origin in Nigeria. BMC Microbiology, 2021, 21, 124.	3.3	12
21	Prevalence of enterotoxin-encoding genes among diverse Shigella strains isolated from patients with diarrhea, southwest Iran. Acta Microbiologica Et Immunologica Hungarica, 2018, 66, 91-101.	0.8	11
22	Detection of Streptococcus gallolyticus in colorectal cancer and inflammatory bowel disease patients compared to control group in southwest of Iran. Molecular Biology Reports, 2020, 47, 8361-8365.	2.3	11
23	Prevalence of bacterial vaginosis and aerobic vaginitis and their associated risk factors among pregnant women from northern Ethiopia: A cross-sectional study. PLoS ONE, 2022, 17, e0262692.	2.5	11
24	<p>Loop-mediated isothermal amplification for detection of Legionella pneumophila in respiratory specimens of hospitalized patients in Ahvaz, southwest Iran</p> . Infection and Drug Resistance, 2019, Volume 12, 529-534.	2.7	10
25	Occurrence of plasmid-mediated quinolone resistance genes in Pseudomonas aeruginosa strains isolated from clinical specimens in southwest Iran: a multicentral study. Scientific Reports, 2022, 12, 2296.	3.3	7
26	Occurrence of FOX AmpC gene among Pseudomonas aeruginosa isolates in abattoir samples from south-eastern Nigeria. Reviews in Medical Microbiology, 2020, 31, 99-103.	0.9	6
27	In vitro evaluation of the antibacterial effects of Cinnamomum zeylanicum essential oil against clinical multidrug-resistant Shigella isolates. Molecular Biology Reports, 2021, 48, 2583-2589.	2.3	6
28	Antibiotic resistance pattern and frequency of cagA and vacA genes in Helicobacter pylori strains isolated from patients in Tabriz city, Iran. BMC Research Notes, 2021, 14, 216.	1.4	6
29	Integron frequency of Escherichia coli strains from patients with urinary tract infection in Southwest of Iran. Journal of Acute Disease, 2019, 8, 113.	0.3	6
30	Identification of metallo-β-lactamases and AmpC production among Escherichia coli strains isolated from hemodialysis patients with urinary tract infection. Molecular Biology Reports, 2021, 48, 7883-7892.	2.3	6
31	Isolation and Identification of Legionella spp. in environmental water sources based on macrophage infectivity potentiator (mip) gene sequencing in southwest Iran. AIMS Microbiology, 2019, 5, 223-231.	2.2	6
32	The prevalence of plasmid-mediated quinolone resistance genes among Escherichia coli strains isolated from urinary tract infections in southwest Iran. Molecular Biology Reports, 2022, 49, 3757-3763.	2.3	6
33	Prevalence of carbapenemases and ESBL encoding genes among K. pneumoniae isolates obtained from an educational hospital in Ahvaz, Southwestern Iran. Gene Reports, 2021, 23, 101128.	0.8	5
34	Multiplex polymerase chain reaction detection of Streptococcus pneumoniae and Haemophilus influenzae and their antibiotic resistance in patients with community-acquired pneumonia from southwest Iran. BMC Microbiology, 2021, 21, 343.	3.3	5
35	Characterization of metallo-β-lactamases-encoding genes blaIMP-1 and blaVIM-1 amongst Klebsiella pneumoniae from abattoir samples of Ebonyi state, southeastern Nigeria. Gene Reports, 2019, 16, 100428.	0.8	4
36	Involvement of the AcrAB Efflux Pump in Ciprofloxacin Resistance in Clinical Klebsiella Pneumoniae Isolates. Infectious Disorders - Drug Targets, 2021, 21, 564-571.	0.8	4

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37	Effects of shallomin and podophyllin solution 25% for genital HPV warts in women: a randomized controlled trial. Journal of Acute Disease, 2019, 8, 118.	0.3	4
38	Study of aerobic and anaerobic bacterial profile of nosocomial infections and their antibiotic resistance in a referral center, Southwest Iran: A three year cross-sectional study. PLoS ONE, 2021, 16, e0259512.	2.5	4
39	Occurrence of Multiple-Drug Resistance Bacteria and Their Antimicrobial Resistance Patterns in Burn Infections from Southwest of Iran. Journal of Burn Care and Research, 2022, 43, 423-431.	0.4	3

40 Study of the D-dimer, C-reactive protein, and autoantibodies markers among HBV infected patients in