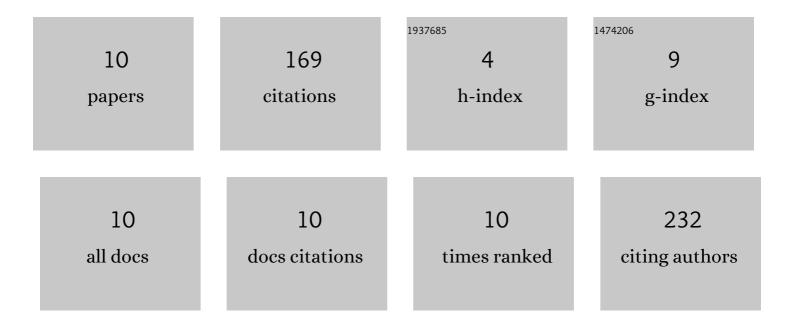
## Ali Bin-Thani

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2592672/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Facile synthesis of zero valent sulfur nanoparticles for catalytic detoxification of hexavalent chromium, cytotoxicity against microalgae and ultraviolet protection properties. Korean Journal of Chemical Engineering, 2021, 38, 2294.	2.7	3
2	Interference of novobiocin in the expression of a truncated blaCTX-M gene causes a phenotypic variation in the production of β-lactamases in Escherichia coli strain EC1091. Gene Reports, 2021, 25, 101390.	0.8	1
3	Patterns of antimicrobial resistance observed in the Middle East: Environmental and health care retrospectives. Science of the Total Environment, 2020, 740, 140089.	8.0	10
4	Characterization of previously identified novel DNA fragment associated with Pathogenicity Island III536 reveals new bla gene. Infection, Genetics and Evolution, 2019, 75, 103971.	2.3	4
5	A Retrospective Epidemiological Study of the Incidence and Risk Factors of Salmonellosis in Bahrain in Children During 2012–2016. Pathogens, 2019, 8, 51.	2.8	2
6	Detection of Novel GyrB Mutations Associated with <i>Escherichia coli </i> Clinical Isolates. Journal of Biomimetics, Biomaterials and Biomedical Engineering, 2018, 35, 88-95.	0.5	4
7	Identification of novel DNA sequence associated with pathogenicity island III536 locus in Uropathogenic Escherichia coli isolate and distribution of virulence determinants in β-lactam resistant isolates. Microbial Pathogenesis, 2018, 123, 393-397.	2.9	4
8	Evaluation of the mobile content in <i>Magnetospirillum magneticum</i> AMB-1 genome using bioinformatical approaches reveals a new genome size for the magnetosome island. Journal of the Association of Arab Universities for Basic and Applied Sciences, 2015, 18, 1-6.	1.0	0
9	MobilomeFINDER: web-based tools for in silico and experimental discovery of bacterial genomic islands. Nucleic Acids Research, 2007, 35, W97-W104.	14.5	74
10	A novel strategy for the identification of genomic islands by comparative analysis of the contents and contexts of tRNA sites in closely related bacteria. Nucleic Acids Research, 2006, 34, e3-e3.	14.5	67