Rania Siam

List of Publications by Year in descending order

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Version: 2024-02-01

304368 329751 1,452 48 22 37 citations h-index g-index papers 49 49 49 2468 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	The ocean sampling day consortium. GigaScience, 2015, 4, 27.	3.3	185
2	A global metagenomic map of urban microbiomes and antimicrobial resistance. Cell, 2021, 184, 3376-3393.e17.	13.5	164
3	Cell cycle regulator phosphorylation stimulates two distinct modes of binding at a chromosome replication origin. EMBO Journal, 2000, 19, 1138-1147.	3.5	65
4	Comparative genomics reveals adaptations of a halotolerant thaumarchaeon in the interfaces of brine pools in the Red Sea. ISME Journal, 2015, 9, 396-411.	4.4	60
5	Isolation and characterization of a heavy metal-resistant, thermophilic esterase from a Red Sea Brine Pool. Scientific Reports, 2013, 3, 3358.	1.6	55
6	Core-Shell Silver/Polymeric Nanoparticles-Based Combinatorial Therapy against Breast Cancer In-vitro. Scientific Reports, 2016, 6, 30729.	1.6	55
7	A Dual Binding Site for Integration Host Factor and the Response Regulator CtrA inside the Caulobacter crescentus Replication Origin. Journal of Bacteriology, 2003, 185, 5563-5572.	1.0	52
8	COVID-19 drug practices risk antimicrobial resistance evolution. Lancet Microbe, The, 2021, 2, e135-e136.	3.4	47
9	Unique Prokaryotic Consortia in Geochemically Distinct Sediments from Red Sea Atlantis II and Discovery Deep Brine Pools. PLoS ONE, 2012, 7, e42872.	1.1	45
10	Patterns of ecological specialization among microbial populations in the $<$ scp $>$ R $<$ /scp $>$ ed $<$ scp $>$ S $<$ /scp $>$ ea and diverse oligotrophic marine environments. Ecology and Evolution, 2013, 3, 1780-1797.	0.8	45
11	Antimicrobial and Wound-Healing Activities of Graphene-Reinforced Electrospun Chitosan/Gelatin Nanofibrous Nanocomposite Scaffolds. ACS Omega, 2022, 7, 1838-1850.	1.6	41
12	Conserved Response Regulator CtrA and IHF Binding Sites in the α-Proteobacteria Caulobacter crescentus and Rickettsia prowazekii Chromosomal Replication Origins. Journal of Bacteriology, 2002, 184, 5789-5799.	1.0	40
13	CtrA, a Global Response Regulator, Uses a Distinct Second Category of Weak DNA Binding Sites for Cell Cycle Transcription Control in <i>Caulobacter crescentus</i>). Journal of Bacteriology, 2009, 191, 5458-5470.	1.0	39
14	Distinct domains of Escherichia coli IgaA connect envelope stress sensing and down-regulation of the Rcs phosphorelay across subcellular compartments. PLoS Genetics, 2018, 14, e1007398.	1.5	38
15	Identification of novel conserved functional motifs across most Influenza A viral strains. Virology Journal, 2011, 8, 44.	1.4	36
16	A Novel Mercuric Reductase from the Unique Deep Brine Environment of Atlantis II in the Red Sea. Journal of Biological Chemistry, 2014, 289, 1675-1687.	1.6	36
17	First Insights into the Viral Communities of the Deep-sea Anoxic Brines of the Red Sea. Genomics, Proteomics and Bioinformatics, 2015, 13, 304-309.	3.0	33
18	Fabrication of Poly(vinyl alcohol)/Chitosan/ <i>Bidens pilosa</i> Composite Electrospun Nanofibers with Enhanced Antibacterial Activities. ACS Omega, 2019, 4, 8778-8785.	1.6	33

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19	Glutamate at the phosphorylation site of response regulator CtrA provides essential activities without increasing DNA binding. Nucleic Acids Research, 2003, 31, 1775-1779.	6.5	29
20	Aerobic methanotrophic communities at the Red Sea brine-seawater interface. Frontiers in Microbiology, 2014, 5, 487.	1.5	29
21	The Egyptian Red Sea coastal microbiome: A study revealing differential microbial responses to diverse anthropogenic pollutants. Environmental Pollution, 2016, 214, 892-902.	3.7	27
22	Conserved Gene Cluster at Replication Origins of the \hat{l}_{\pm} -Proteobacteria Caulobacter crescentus and Rickettsia prowazekii. Journal of Bacteriology, 2001, 183, 1824-1829.	1.0	26
23	Genome Sequence of Haloplasma contractile, an Unusual Contractile Bacterium from a Deep-Sea Anoxic Brine Lake. Journal of Bacteriology, 2011, 193, 4551-4552.	1.0	26
24	Transcriptional activation of the Lats1 tumor suppressor gene in tumors of CUX1 transgenic mice. Molecular Cancer, 2009, 8, 60.	7.9	21
25	Core Microbial Functional Activities in Ocean Environments Revealed by Global Metagenomic Profiling Analyses. PLoS ONE, 2014, 9, e97338.	1.1	20
26	Novel thermostable antibiotic resistance enzymes from the Atlantis II Deep Red Sea brine pool. Microbial Biotechnology, 2017, 10, 189-202.	2.0	20
27	Prevalence and antimicrobial resistance pattern of bacterial meningitis in Egypt. Annals of Clinical Microbiology and Antimicrobials, 2009, 8, 26.	1.7	19
28	Red Sea Atlantis II brine pool nitrilase with unique thermostability profile and heavy metal tolerance. BMC Biotechnology, 2016, 16, 14.	1.7	19
29	<i>In situ</i> polymerization of polyurethaneâ€silver nanocomposite foams with intact thermal stability, improved mechanical performance, and induced antimicrobial properties. Journal of Applied Polymer Science, 2016, 133, .	1.3	18
30	Antibacterial and anticancer activities of orphan biosynthetic gene clusters from Atlantis II Red Sea brine pool. Microbial Cell Factories, 2019, 18, 56.	1.9	18
31	Novel Inulin Electrospun Composite Nanofibers: Prebiotic and Antibacterial Activities. ACS Omega, 2020, 5, 3006-3015.	1.6	17
32	Egypt's Red Sea coast: phylogenetic analysis of cultured microbial consortia in industrialized sites. Frontiers in Microbiology, 2014, 5, 363.	1.5	16
33	Antibiotic Resistome: Improving Detection and Quantification Accuracy for Comparative Metagenomics. OMICS A Journal of Integrative Biology, 2016, 20, 229-238.	1.0	14
34	Insights into Red Sea Brine Pool Specialized Metabolism Gene Clusters Encoding Potential Metabolites for Biotechnological Applications and Extremophile Survival. Marine Drugs, 2019, 17, 273.	2.2	14
35	Viruses-to-mobile genetic elements skew in the deep Atlantis II brine pool sediments. Scientific Reports, 2016, 6, 32704.	1.6	11
36	Integration of tri-polar microelectrodes for performance enhancement of an impedance biosensor. Sensing and Bio-Sensing Research, 2020, 28, 100329.	2.2	10

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37	New genetic variants of LATS1 detected in urinary bladder and colon cancer. Frontiers in Genetics, 2015, 5, 425.	1.1	9
38	Correlating novel variable and conserved motifs in the Hemagglutinin protein with significant biological functions. Virology Journal, 2008, 5, 91.	1.4	4
39	Schizosaccharomyces pombeRad4/Cut5 Protein Modification and Chromatin Binding Changes in DNA Damage. DNA and Cell Biology, 2007, 26, 565-575.	0.9	3
40	Insertion sequences enrichment in extreme Red sea brine pool vent. Extremophiles, 2017, 21, 271-282.	0.9	3
41	Microfluidic Platform for Monitoring the Dielectric Parameters of U2OS Cells. , 2019, , .		2
42	The association of group IIB intron with integrons in hypersaline environments. Mobile DNA, 2021, 12, 8.	1.3	2
43	Evaluation of a Thermophilic, Psychrostable, and Heavy Metal-Resistant Red Sea Brine Pool Esterase. Marine Drugs, 2022, 20, 274.	2.2	2
44	Bioprospecting the microbiome of Red Sea Atlantis II brine pool for peptidases and biosynthetic genes with promising antibacterial activity. Microbial Cell Factories, 2022, 21, .	1.9	2
45	Molecular characterization of extended-spectrum-β-lactamases producing Klebsiella pneumoniae and Escherichia coli from hospitalized patients in Oman. BMC Proceedings, 2009, 3, O10.	1.8	1
46	An optimized protocol for high yield expression and purification of an extremophilic protein. Protein Expression and Purification, 2020, 169, 105585.	0.6	1
47	Abundance of integrons in halophilic bacteria. Canadian Journal of Microbiology, 2022, , 1-11.	0.8	0
48	A Developed Micro Electric Impedance Spectroscopy for Biological Cells' Electrical Parameters Extraction. , 2021, , .		0