

Mamoon Rashid

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7878523/publications.pdf>

Version: 2024-02-01

14
papers

2,269
citations

933447

10
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

6638
citing authors

#	ARTICLE	IF	CITATIONS
1	Interferon-induced transmembrane protein-3 genetic variant rs12252 is associated with COVID-19 mortality. <i>Genomics</i> , 2021, 113, 1733-1741.	2.9	39
2	Discovery of a novel potentially transforming somatic mutation in <i>CSF2RB</i> gene in breast cancer. <i>Cancer Medicine</i> , 2021, 10, 8138-8150.	2.8	6
3	SARS-CoV-2 ORF8 and SARS-CoV ORF8ab: Genomic Divergence and Functional Convergence. <i>Pathogens</i> , 2020, 9, 677.	2.8	44
4	Molecular classification of colorectal cancer using the gene expression profile of tumor samples. <i>Experimental Biology and Medicine</i> , 2019, 244, 1005-1016.	2.4	5
5	Metabolic traits of an uncultured archaeal lineage -MSBL1- from brine pools of the Red Sea. <i>Scientific Reports</i> , 2016, 6, 19181.	3.3	66
6	Contemporary molecular tools in microbial ecology and their application to advancing biotechnology. <i>Biotechnology Advances</i> , 2015, 33, 1755-1773.	11.7	31
7	Comparative genomics reveals adaptations of a halotolerant thaumarchaeon in the interfaces of brine pools in the Red Sea. <i>ISME Journal</i> , 2015, 9, 396-411.	9.8	60
8	The Population Structure of <i>Vibrio cholerae</i> from the Chandigarh Region of Northern India. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2981.	3.0	21
9	A promoter-level mammalian expression atlas. <i>Nature</i> , 2014, 507, 462-470.	27.8	1,838
10	Designing of peptides with desired half-life in intestine-like environment. <i>BMC Bioinformatics</i> , 2014, 15, 282.	2.6	73
11	Genomic differentiation among two strains of the PS1 clade isolated from geographically separated marine habitats. <i>FEMS Microbiology Ecology</i> , 2014, 89, 181-197.	2.7	22
12	READSCAN: a fast and scalable pathogen discovery program with accurate genome relative abundance estimation. <i>Bioinformatics</i> , 2013, 29, 391-392.	4.1	51
13	Complete Genome Sequence of <i>Mycobacterium xenopi</i> Type Strain RIVM700367. <i>Journal of Bacteriology</i> , 2012, 194, 3282-3283.	2.2	3
14	Complete Genome Sequence of <i>Mycobacterium phlei</i> Type Strain RIVM601174. <i>Journal of Bacteriology</i> , 2012, 194, 3284-3285.	2.2	10