

Samiul Islam

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7717162/publications.pdf>

Version: 2024-02-01

11
papers

165
citations

1477746

6
h-index

1372195

10
g-index

11
all docs

11
docs citations

11
times ranked

209
citing authors

#	ARTICLE	IF	CITATIONS
1	Virtual screening and molecular dynamics simulation study of plant-derived compounds to identify potential inhibitors of main protease from SARS-CoV-2. Briefings in Bioinformatics, 2021, 22, 1402-1414.	3.2	75
2	Identification and in silico molecular modelling study of newly isolated Bacillus subtilis SI-18 strain against S9 protein of Rhizoctonia solani. Arabian Journal of Chemistry, 2020, 13, 8600-8612.	2.3	22
3	Identification of host transcriptome-guided repurposable drugs for SARS-CoV-1 infections and their validation with SARS-CoV-2 infections by using the integrated bioinformatics approaches. PLoS ONE, 2022, 17, e0266124.	1.1	18
4	In vitro and in silico approach of fungal growth inhibition by Trichoderma asperellum HbGT6-07 derived volatile organic compounds. Arabian Journal of Chemistry, 2021, 14, 103290.	2.3	11
5	Isolation of Antimicrobial Genes from Oryza rufipogon Griff by Using a Bacillus subtilis Expression System with Potential Antimicrobial Activities. International Journal of Molecular Sciences, 2020, 21, 8722.	1.8	10
6	Characterization and biocontrol measures of Pseudomonas syringae pv. syringae associated with citrus blast disease. Vegetos, 2020, 33, 555-569.	0.8	8
7	Two antimicrobial genes from Aegilops tauschii Cosson identified by the Bacillus subtilis expression system. Scientific Reports, 2020, 10, 13346.	1.6	7
8	Isolation of Potato Endophytes and Screening of Chaetomium globosum Antimicrobial Genes. International Journal of Molecular Sciences, 2022, 23, 4611.	1.8	6
9	Antimicrobial Peptides from Plants: A cDNA-Library Based Isolation, Purification, Characterization Approach and Elucidating Their Modes of Action. International Journal of Molecular Sciences, 2021, 22, 8712.	1.8	5
10	The investigation of nonsynonymous SNPs of human SLC6A4 gene associated with depression: An in silico approach. Heliyon, 2021, 7, e07815.	1.4	2
11	Prevalence of Tuberculosis: Present Status and Overview of Its Control System in Bangladesh. , 2017, 3, 1471-1475.		1