

Muhamad Fahmi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9268660/publications.pdf>

Version: 2024-02-01

9
papers

132
citations

1683354

5
h-index

1473754

9
g-index

14
all docs

14
docs citations

14
times ranked

200
citing authors

#	ARTICLE	IF	CITATIONS
1	Nonstructural proteins NS7b and NS8 are likely to be phylogenetically associated with evolution of 2019-nCoV. <i>Infection, Genetics and Evolution</i> , 2020, 81, 104272.	1.0	48
2	Retrieval and Investigation of Data on SARS-CoV-2 and COVID-19 Using Bioinformatics Approach. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1318, 839-857.	0.8	23
3	Evolutionary Approach of Intrinsically Disordered CIP/KIP Proteins. <i>Scientific Reports</i> , 2019, 9, 1575.	1.6	16
4	In Silico Study of Rett Syndrome Treatment-Related Genes, MECP2, CDKL5, and FOXP1, by Evolutionary Classification and Disordered Region Assessment. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5593.	1.8	12
5	The Functional Classification of ORF8 in SARS-CoV-2 Replication, Immune Evasion, and Viral Pathogenesis Inferred through Phylogenetic Profiling. <i>Evolutionary Bioinformatics</i> , 2021, 17, 117693432110030.	0.6	9
6	In silico Analysis of SARS-CoV-2 ORF8-Binding Proteins Reveals the Involvement of ORF8 in Acquired-Immune and Innate-Immune Systems. <i>Frontiers in Medicine</i> , 2022, 9, 824622.	1.2	8
7	Genome-Wide Analysis of Whole Human Glycoside Hydrolases by Data-Driven Analysis in Silico. <i>International Journal of Molecular Sciences</i> , 2019, 20, 6290.	1.8	6
8	Gene Cascade Finder: A tool for identification of gene cascades and its application in <i>Caenorhabditis elegans</i> . <i>PLoS ONE</i> , 2019, 14, e0215187.	1.1	5
9	Overlapping and non-overlapping roles of the class-I histone deacetylase-1 corepressors LET-418, SIN-3, and SPR-1 in <i>Caenorhabditis elegans</i> embryonic development. <i>Genes and Genomics</i> , 2021, 43, 553-565.	0.5	3