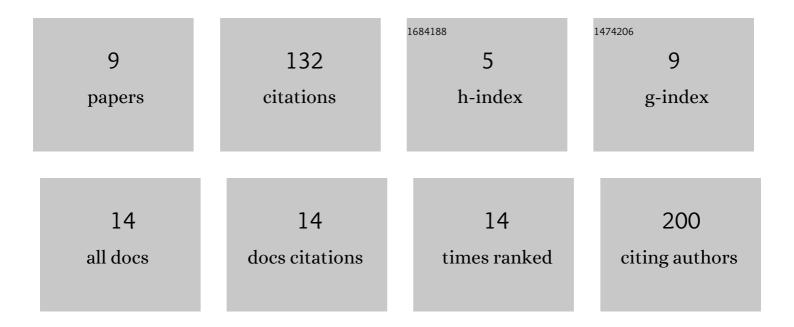
## Muhamad Fahmi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9268660/publications.pdf Version: 2024-02-01



Μιιμαμαρ Γλημι

#	Article	IF	CITATIONS
1	Nonstructural proteins NS7b and NS8 are likely to be phylogenetically associated with evolution of 2019-nCoV. Infection, Genetics and Evolution, 2020, 81, 104272.	2.3	48
2	Retrieval and Investigation of Data on SARS-CoV-2 and COVID-19 Using Bioinformatics Approach. Advances in Experimental Medicine and Biology, 2021, 1318, 839-857.	1.6	23
3	Evolutionary Approach of Intrinsically Disordered CIP/KIP Proteins. Scientific Reports, 2019, 9, 1575.	3.3	16
4	In Silico Study of Rett Syndrome Treatment-Related Genes, MECP2, CDKL5, and FOXG1, by Evolutionary Classification and Disordered Region Assessment. International Journal of Molecular Sciences, 2019, 20, 5593.	4.1	12
5	The Functional Classification of ORF8 in SARS-CoV-2 Replication, Immune Evasion, and Viral Pathogenesis Inferred through Phylogenetic Profiling. Evolutionary Bioinformatics, 2021, 17, 117693432110030.	1.2	9
6	In silico Analysis of SARS-CoV-2 ORF8-Binding Proteins Reveals the Involvement of ORF8 in Acquired-Immune and Innate-Immune Systems. Frontiers in Medicine, 2022, 9, 824622.	2.6	8
7	Genome-Wide Analysis of Whole Human Glycoside Hydrolases by Data-Driven Analysis in Silico. International Journal of Molecular Sciences, 2019, 20, 6290.	4.1	6
8	Gene Cascade Finder: A tool for identification of gene cascades and its application in Caenorhabditis elegans. PLoS ONE, 2019, 14, e0215187.	2.5	5
9	Overlapping and non-overlapping roles of the class-I histone deacetylase-1 corepressors LET-418, SIN-3, and SPR-1 in Caenorhabditis elegans embryonic development. Genes and Genomics, 2021, 43, 553-565.	1.4	3