Abdur Rehman

List of Publications by Year
in descending order

13	309	1040056	1125743
13 papers	309 citations	9 h-index	13 g-index
13 all docs	13 docs citations	13 times ranked	231 citing authors

#	Article	IF	Citations
1	The Screening of Phytochemicals Against NS5 Polymerase to Treat Zika Virus Infection: Integrated Computational Based Approach. Combinatorial Chemistry and High Throughput Screening, 2022, 25, 738-751.	1.1	22
2	Designing of a multi-epitopes-based peptide vaccine against rift valley fever virus and its validation through integrated computational approaches. Computers in Biology and Medicine, 2022, 141, 105151.	7.0	16
3	Integrating Network Pharmacology and Molecular Docking Approaches to Decipher the Multi-Target Pharmacological Mechanism of Abrus precatorius L. Acting on Diabetes. Pharmaceuticals, 2022, 15, 414.	3.8	32
4	Designing a multi-epitope vaccine against Chlamydia pneumoniae by integrating the core proteomics, subtractive proteomics and reverse vaccinology-based immunoinformatics approaches. Computers in Biology and Medicine, 2022, 145, 105507.	7.0	12
5	Network Pharmacology and Bioinformatics Approach Reveals the Multi-Target Pharmacological Mechanism of Fumaria indica in the Treatment of Liver Cancer. Pharmaceuticals, 2022, 15, 654.	3.8	17
6	Discovery of Rift Valley fever virus natural pan-inhibitors by targeting its multiple key proteins through computational approaches. Scientific Reports, 2022, 12, .	3.3	13
7	Immunoinformatics and Molecular Docking Studies Predicted Potential Multiepitope-Based Peptide Vaccine and Novel Compounds against Novel SARS-CoV-2 through Virtual Screening. BioMed Research International, 2021, 2021, 1-20.	1.9	26
8	Anti-aging Natural Compounds and their Role in the Regulation of Metabolic Pathways Leading to Longevity. Mini-Reviews in Medicinal Chemistry, 2021, 21, 2630-2656.	2.4	14
9	Integrated Core Proteomics, Subtractive Proteomics, and Immunoinformatics Investigation to Unveil a Potential Multi-Epitope Vaccine against Schistosomiasis. Vaccines, 2021, 9, 658.	4.4	30
10	Designing a Multi-Epitope Vaccine against Chlamydia trachomatis by Employing Integrated Core Proteomics, Immuno-Informatics and In Silico Approaches. Biology, 2021, 10, 997.	2.8	30
11	In Silico Core Proteomics and Molecular Docking Approaches for the Identification of Novel Inhibitors against Streptococcus pyogenes. International Journal of Environmental Research and Public Health, 2021, 18, 11355.	2.6	6
12	Development of a Candidate Multi-Epitope Subunit Vaccine against Klebsiella aerogenes: Subtractive Proteomics and Immuno-Informatics Approach. Vaccines, 2021, 9, 1373.	4.4	10
13	Designing of a next generation multiepitope based vaccine (MEV) against SARS-COV-2: Immunoinformatics and in silico approaches. PLoS ONE, 2020, 15, e0244176.	2.5	81