

Muhammad Naveed

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

33
papers

510
citations

9
h-index

22
g-index

41
ext. papers

1,009
ext. citations

3.4
avg, IF

3.71
L-index

#	Paper	IF	Citations
33	Causes of blindness and vision impairment in 2020 and trends over 30 years, and prevalence of avoidable blindness in relation to VISION 2020: the Right to Sight: an analysis for the Global Burden of Disease Study. <i>The Lancet Global Health</i> , 2021 , 9, e144-e160	13.6	253
32	L-asparaginase as a critical component to combat Acute Lymphoblastic Leukaemia (ALL): A novel approach to target ALL. <i>European Journal of Pharmacology</i> , 2016 , 771, 199-210	5.3	41
31	Comprehensive review on the molecular genetics of autosomal recessive primary microcephaly (MCPH). <i>Genetical Research</i> , 2018 , 100, e7	1.1	38
30	Community-Based Assessment of Knowledge, Attitude, Practices and Risk Factors Regarding COVID-19 Among Pakistanis Residents During a Recent Outbreak: A Cross-Sectional Survey. <i>Journal of Community Health</i> , 2021 , 46, 476-486	4	38
29	Protease α Versatile and Ecofriendly Biocatalyst with Multi-Industrial Applications: An Updated Review. <i>Catalysis Letters</i> , 2021 , 151, 307-323	2.8	25
28	Protease Production from <i>Cheatomium globusum</i> Through Central Composite Design Using Agricultural Wastes and Its Immobilization for Industrial Exploitation. <i>Waste and Biomass Valorization</i> , 2020 , 11, 6529-6539	3.2	12
27	In-silico analysis of non-synonymous-SNPs of STEAP2: To provoke the progression of prostate cancer. <i>Open Life Sciences</i> , 2016 , 11, 402-416	1.2	12
26	Targeting STAT-3 signaling pathway in cancer for development of novel drugs: Advancements and challenges. <i>Genetics and Molecular Biology</i> , 2020 , 43, e20180160	2	11
25	Structural and functional annotation of hypothetical proteins of human adenovirus: prioritizing the novel drug targets. <i>BMC Research Notes</i> , 2017 , 10, 706	2.3	10
24	The diversity of Rhizobia, Sinorhizobia and novel non-Rhizobial <i>Paenibacillus nodulating</i> wild herbaceous legumes. <i>Archives of Microbiology</i> , 2013 , 195, 647-53	3	9
23	Bioinformatics based structural characterization of glucose dehydrogenase (gdh) gene and growth promoting activity of <i>Leclercia</i> sp. QAU-66. <i>Brazilian Journal of Microbiology</i> , 2014 , 45, 603-11	2.2	9
22	Plasma Biomarkers: Potent Screeners of Alzheimer's Disease. <i>American Journal of Alzheimerts Disease and Other Dementias</i> , 2019 , 34, 290-301	2.5	7
21	Evaluation of Glucose Dehydrogenase and Pyrroloquinoline Quinine (pqq) Mutagenesis that Renders Functional Inadequacies in Host Plants. <i>Journal of Microbiology and Biotechnology</i> , 2015 , 25, 1349-60	3.3	7
20	Identification and characterization of rhizospheric microbial diversity by 16S ribosomal RNA gene sequencing. <i>Brazilian Journal of Microbiology</i> , 2014 , 45, 985-93	2.2	5
19	Clastogenic ROS and biophotonics in precancerous diagnosis. <i>Frontiers in Biology</i> , 2018 , 13, 103-122		4
18	Structural and Functional Annotation of Conserved Virulent Hypothetical Proteins in <i>Chlamydia Trachomatis</i> : An In-Silico Approach. <i>Current Bioinformatics</i> , 2019 , 14, 344-352	4.7	4
17	Design of a novel multiple epitope-based vaccine: An immunoinformatics approach to combat SARS-CoV-2 strains. <i>Journal of Infection and Public Health</i> , 2021 , 14, 938-946	7.4	3

16	A Vaccine Construction against COVID-19-Associated Mucormycosis Contrived with Immunoinformatics-Based Scavenging of Potential Mucoralean Epitopes. <i>Vaccines</i> , 2022 , 10, 664	5.3	3
15	functional and tumor suppressor role of hypothetical protein PCNXL2 with regulation of the Notch signaling pathway.. <i>RSC Advances</i> , 2018 , 8, 21414-21430	3.7	2
14	Computational Analysis and Polymorphism study of Tumor Suppressor Candidate Gene-3 for Non Syndromic Autosomal Recessive Mental Retardation 2016 , 05,		2
13	Geographical, Molecular, and Computational Analysis of Migraine-Causing Genes. <i>Journal of Computational Biophysics and Chemistry</i> , 2021 , 20, 391-403		2
12	In-Silico analysis of missense SNPs in Human HPPD gene associated with Tyrosinemia type III and Hawkinsinuria. <i>Computational Biology and Chemistry</i> , 2019 , 80, 284-291	3.6	1
11	Swimming with Death as the Major Outbreak in Pakistan. <i>Iranian Journal of Public Health</i> , 2019 , 48, 1741-1742	1.742	1
10	Major natural sinks for harboring microorganisms with altered antibiotic resistance versus major human contributing sources of antibiotic resistance: a detailed insight 2020 , 70-98		1
9	Antibiotics resistance mechanism 2020 , 292-312		1
8	Neuroproteomics in Paving the Pathway for Drug Abuse Research. <i>Current Proteomics</i> , 2019 , 16, 256-266	0.7	0
7	Characterization of Mercury-Resistant Rhizobacteria for Plant Growth Promotion: An In Vitro and In Silico Approach. <i>Current Microbiology</i> , 2021 , 78, 3968-3979	2.4	0
6	Bio-Molecular analysis of selected food derived Lactiplantibacillus strains for CLA production reveals possibly a complex mechanism.. <i>Food Research International</i> , 2022 , 154, 111031	7	0
5	Dengue death tolls: A nightmare for Khyber Pakhtunkhwa, Pakistan. <i>Journal of Infection and Public Health</i> , 2018 , 11, 898-899	7.4	
4	Computational and Pathway Analysis of nsSNPs of MED23 Gene Involved in Human Congenital Diseases. <i>Current Biotechnology</i> , 2018 , 7, 132-147	0.6	
3	Geographical, Molecular, and Computational Analysis of Migraine-Causing Genes. <i>Journal of Computational Biophysics and Chemistry</i> , 2021 , 20, 557-557		
2	The syringes catastrophe of the HIV outbreak in Faisalabad, Punjab, Pakistan. <i>Future Virology</i> , 2019 , 14, 571-572	2.4	
1	TULP3 NLS inhibition: an study to hamper cargo transport to nucleus.. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022 , 1-9	3.6	