Muhammad Naveed

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4470734/publications.pdf

Version: 2024-02-01

41 papers 1,787 citations

623574 14 h-index 34 g-index

41 all docs

41 docs citations

41 times ranked

1760 citing authors

#	Article	IF	CITATIONS
1	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. The Lancet Global Health, 2021, 9, e144-e160.	2.9	1,148
2	Proteaseâ€"A Versatile and Ecofriendly Biocatalyst with Multi-Industrial Applications: An Updated Review. Catalysis Letters, 2021, 151, 307-323.	1.4	84
3	Community-Based Assessment of Knowledge, Attitude, Practices and Risk Factors Regarding COVID-19 Among Pakistanis Residents During a Recent Outbreak: A Cross-Sectional Survey. Journal of Community Health, 2021, 46, 476-486.	1.9	59
4	L-asparaginase as a critical component to combat Acute Lymphoblastic Leukaemia (ALL): A novel approach to target ALL. European Journal of Pharmacology, 2016, 771, 199-210.	1.7	57
5	Comprehensive review on the molecular genetics of autosomal recessive primary microcephaly (MCPH). Genetical Research, 2018, 100, e7.	0.3	54
6	Role of Mitochondrial Membrane Potential and Lactate Dehydrogenase A in Apoptosis. Anti-Cancer Agents in Medicinal Chemistry, 2022, 22, 2048-2062.	0.9	50
7	Green Synthesis of Silver Nanoparticles Using the Plant Extract of Acer oblongifolium and Study of Its Antibacterial and Antiproliferative Activity via Mathematical Approaches. Molecules, 2022, 27, 4226.	1.7	37
8	Design of a novel multiple epitope-based vaccine: An immunoinformatics approach to combat SARS-CoV-2 strains. Journal of Infection and Public Health, 2021, 14, 938-946.	1.9	31
9	Targeting STAT-3 signaling pathway in cancer for development of novel drugs: Advancements and challenges. Genetics and Molecular Biology, 2020, 43, e20180160.	0.6	25
10	In-silico analysis of non-synonymous-SNPs of STEAP2: To provoke the progression of prostate cancer. Open Life Sciences, 2016, 11, 402-416.	0.6	22
11	A Vaccine Construction against COVID-19-Associated Mucormycosis Contrived with Immunoinformatics-Based Scavenging of Potential Mucoralean Epitopes. Vaccines, 2022, 10, 664.	2.1	22
12	Self-Medication as an Important Risk Factor for Antibiotic Resistance: A Multi-Institutional Survey among Students. Antibiotics, 2022, 11, 842.	1.5	22
13	Bioinformatics based structural characterization of glucose dehydrogenase (gdh) gene and growth promoting activity of Leclercia sp. QAU-66. Brazilian Journal of Microbiology, 2014, 45, 603-611.	0.8	20
14	Structural and functional annotation of hypothetical proteins of human adenovirus: prioritizing the novel drug targets. BMC Research Notes, 2017, 10, 706.	0.6	20
15	Protease Production from Cheotomium globusum Through Central Composite Design Using Agricultural Wastes and Its Immobilization for Industrial Exploitation. Waste and Biomass Valorization, 2020, 11, 6529-6539.	1.8	15
16	Bio-Molecular analysis of selected food derived Lactiplantibacillus strains for CLA production reveals possibly a complex mechanism. Food Research International, 2022, 154, 111031.	2.9	15
17	Identification and characterization of rhizospheric microbial diversity by 16S ribosomal RNA gene sequencing. Brazilian Journal of Microbiology, 2014, 45, 985-993.	0.8	14
18	Execution and Design of an Anti HPIV-1 Vaccine with Multiple Epitopes Triggering Innate and Adaptive Immune Responses: An Immunoinformatic Approach. Vaccines, 2022, 10, 869.	2.1	14

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19	The diversity of Rhizobia, Sinorhizobia and novel non-Rhizobial Paenibacillus nodulating wild herbaceous legumes. Archives of Microbiology, 2013, 195, 647-653.	1.0	10
20	Plasma Biomarkers: Potent Screeners of Alzheimer's Disease. American Journal of Alzheimer's Disease and Other Dementias, 2019, 34, 290-301.	0.9	10
21	Evaluation of Glucose Dehydrogenase and Pyrroloquinoline Quinine (pqq) Mutagenesis that Renders Functional Inadequacies in Host Plants. Journal of Microbiology and Biotechnology, 2015, 25, 1349-1360.	0.9	9
22	<i>In silico</i> functional and tumor suppressor role of hypothetical protein PCNXL2 with regulation of the Notch signaling pathway. RSC Advances, 2018, 8, 21414-21430.	1.7	7
23	In-Silico analysis of missense SNPs in Human HPPD gene associated with Tyrosinemia type III and Hawkinsinuria. Computational Biology and Chemistry, 2019, 80, 284-291.	1.1	6
24	Geographical, Molecular, and Computational Analysis of Migraine-Causing Genes. Journal of Computational Biophysics and Chemistry, 2021, 20, 391-403.	1.0	6
25	Evaluation of Bi-Lateral Co-Infections and Antibiotic Resistance Rates among COVID-19 Patients in Lahore, Pakistan. Medicina (Lithuania), 2022, 58, 904.	0.8	6
26	Structural and Functional Annotation of Conserved Virulent Hypothetical Proteins in Chlamydia Trachomatis: An In-Silico Approach. Current Bioinformatics, 2019, 14, 344-352.	0.7	5
27	Clastogenic ROS and biophotonics in precancerous diagnosis. Frontiers in Biology, 2018, 13, 103-122.	0.7	4
28	Major natural sinks for harboring microorganisms with altered antibiotic resistance versus major human contributing sources of antibiotic resistance: a detailed insight., 2020,, 70-98.		3
29	Antibiotics resistance mechanism. , 2020, , 292-312.		2
30	Characterization of Mercury-Resistant Rhizobacteria for Plant Growth Promotion: An In Vitro and In Silico Approach. Current Microbiology, 2021, 78, 3968-3979.	1.0	2
31	Computational Analysis and Polymorphism study of Tumor Suppressor Candidate Gene-3 for Non Syndromic Autosomal Recessive Mental Retardation. Journal of Applied Bioinformatics & Computational Biology, 2016, 05, .	0.2	2
32	The Virulent Hypothetical Proteins: The Potential Drug Target Involved in Bacterial Pathogenesis. Mini-Reviews in Medicinal Chemistry, 2022, 22, 2608-2623.	1.1	2
33	Dengue death tolls: A nightmare for Khyber Pakhtunkhwa, Pakistan. Journal of Infection and Public Health, 2018, 11, 898-899.	1.9	1
34	Neuroproteomics in Paving the Pathway for Drug Abuse Research. Current Proteomics, 2019, 16, 256-266.	0.1	1
35	Swimming with Death as the Major Outbreak in Pakistan. Iranian Journal of Public Health, 2019, 48, 1741-1742.	0.3	1
36	Computational Drug Shifting towards Drug-Drug Conjugates and Monoclonal Antibody Conjugates in the Contradictory Excursion of Asthma. Letters in Drug Design and Discovery, 2022, 19, .	0.4	1

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37	The syringes catastrophe of the HIV outbreak in Faisalabad, Punjab, Pakistan. Future Virology, 2019, 14, 571-572.	0.9	O
38	Geographical, Molecular, and Computational Analysis of Migraine-Causing Genes. Journal of Computational Biophysics and Chemistry, 2021, 20, 557-557.	1.0	0
39	Computational and Pathway Analysis of nsSNPs of MED23 Gene Involved in Human Congenital Diseases. Current Biotechnology, 2018, 7, 132-147.	0.2	O
40	Naegleria fowleri: Swimming with Death as the Major Outbreak in Pakistan. Iranian Journal of Public Health, 0, , .	0.3	0
41	TULP3 NLS inhibition: an <i>in silico</i> study to hamper cargo transport to nucleus. Journal of Biomolecular Structure and Dynamics, 2022, , 1-9.	2.0	0