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Nucleic Acids Research

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Using a public database of <i>Neisseria gonorrhoeae</i> genomes to detect mutations associated with zoliflodacin resistance. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2847-2849.	1.3	13
2	The Ensembl COVID-19 resource: ongoing integration of public SARS-CoV-2 data. <i>Nucleic Acids Research</i> , 2022, 50, D765-D770.	6.5	10
3	ProTG4: A Web Server to Approximate the Sequence of a Generic Protein From an in Silico Library of Translatable G-Quadruplex (TG4)-Mapped Peptides. <i>Bioinformatics and Biology Insights</i> , 2021, 15, 117793222110458.	1.0	0
4	Cold-Shock Domains' Abundance, Structure, Properties, and Nucleic-Acid Binding. <i>Cancers</i> , 2021, 13, 190.	1.7	35
5	Influence of charge configuration on substrate binding to SARS-CoV-2 main protease. <i>Chemical Communications</i> , 2021, 57, 5314-5317.	2.2	14
6	PathBIX' a web server for network-based pathway annotation with adaptive null models. <i>Bioinformatics Advances</i> , 2021, 1, .	0.9	5
7	ProNAB: database for binding affinities of protein' nucleic acid complexes and their mutants. <i>Nucleic Acids Research</i> , 2022, 50, D1528-D1534.	6.5	20
9	Signalink3: a multi-layered resource to uncover tissue-specific signaling networks. <i>Nucleic Acids Research</i> , 2022, 50, D701-D709.	6.5	19
10	BIONDA: a free database for a fast information on published biomarkers. <i>Bioinformatics Advances</i> , 2021, 1, .	0.9	5
11	Modelling the active SARS-CoV-2 helicase complex as a basis for structure-based inhibitor design. <i>Chemical Science</i> , 2021, 12, 13492-13505.	3.7	6
12	NPCDR: natural product-based drug combination and its disease-specific molecular regulation. <i>Nucleic Acids Research</i> , 2022, 50, D1324-D1333.	6.5	21
13	Identifying protein subcellular locations with embeddings-based node2loc. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2021, PP, 1-1.	1.9	26
14	Representation learning applications in biological sequence analysis. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 3198-3208.	1.9	42
15	Gut Microbiome-Based Analysis of Lipid A Biosynthesis in Individuals with Autism Spectrum Disorder: An In Silico Evaluation. <i>Nutrients</i> , 2021, 13, 688.	1.7	16
18	Effect of Syringopicroside Extracted from <i>Syringa oblata</i> Lindl on the Biofilm Formation of <i>Streptococcus suis</i> . <i>Molecules</i> , 2021, 26, 1295.	1.7	10
19	BAGET 2.0: an updated web tool for the effortless retrieval of prokaryotic gene context and sequence. <i>Bioinformatics</i> , 2021, 37, 2750-2752.	1.8	1
23	Quantitative Proteomic Approach Reveals Altered Metabolic Pathways in Response to the Inhibition of Lysine Deacetylases in A549 Cells under Normoxia and Hypoxia. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3378.	1.8	3
24	Genetics in the Host' Mycobacterium ulcerans interaction. <i>Immunological Reviews</i> , 2021, 301, 222-241.	2.8	0

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31	PrecisionProDB: improving the proteomics performance for precision medicine. <i>Bioinformatics</i> , 2021, 37, 3361-3363.	1.8	4
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40	Regulation of MST complexes and activity via SARAH domain modifications. <i>Biochemical Society Transactions</i> , 2021, 49, 675-683.	1.6	9
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43	The First Insight into Polyhydroxyalkanoates Accumulation in Multi-Extremophilic <i>Rubrobacter xylanophilus</i> and <i>Rubrobacter spartanus</i> . <i>Microorganisms</i> , 2021, 9, 909.	1.6	28
44	The Protein Interactome of Glycolysis in <i>Escherichia coli</i> . <i>Proteomes</i> , 2021, 9, 16.	1.7	3
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61	PredictProtein - Predicting Protein Structure and Function for 29 Years. <i>Nucleic Acids Research</i> , 2021, 49, W535-W540.	6.5	135

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64	AniProtDB: A Collection of Consistently Generated Metazoan Proteomes for Comparative Genomics Studies. <i>Molecular Biology and Evolution</i> , 2021, 38, 4628-4633.	3.5	5
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79	MTR3D: identifying regions within protein tertiary structures under purifying selection. <i>Nucleic Acids Research</i> , 2021, 49, W438-W445.	6.5	17
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