

Niema Moshiri

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8238959/publications.pdf>

Version: 2024-02-01

15
papers

507
citations

1307594
7
h-index

1281871
11
g-index

28
all docs

28
docs citations

28
times ranked

698
citing authors

#	ARTICLE	IF	CITATIONS
1	Timing the SARS-CoV-2 index case in Hubei province. <i>Science</i> , 2021, 372, 412-417.	12.6	109
2	TreeCluster: Clustering biological sequences using phylogenetic trees. <i>PLoS ONE</i> , 2019, 14, e0221068.	2.5	99
3	Ten simple rules for writing and sharing computational analyses in Jupyter Notebooks. <i>PLoS Computational Biology</i> , 2019, 15, e1007007.	3.2	86
4	ViralMSA: massively scalable reference-guided multiple sequence alignment of viral genomes. <i>Bioinformatics</i> , 2021, 37, 714-716.	4.1	55
5	FAVITES: simultaneous simulation of transmission networks, phylogenetic trees and sequences. <i>Bioinformatics</i> , 2019, 35, 1852-1861.	4.1	45
6	GenieHD: Efficient DNA Pattern Matching Accelerator Using Hyperdimensional Computing. , 2020, , .		34
7	TreeSwift: A massively scalable Python tree package. <i>SoftwareX</i> , 2020, 11, 100436.	2.6	23
8	Accelerators for Classical Molecular Dynamics Simulations of Biomolecules. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 4047-4069.	5.3	15
9	A Two-State Model of Tree Evolution and Its Applications to Alu Retrotransposition. <i>Systematic Biology</i> , 2018, 67, 475-489.	5.6	5
10	HIV Care Prioritization Using Phylogenetic Branch Length. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2021, 86, 626-637.	2.1	5
11	NiemaGraphGen: A memory-efficient global-scale contact network simulation toolkit. <i>GigaByte</i> , 0, 2022, 1-11.	0.0	2
12	An Evaluation of Phylogenetic Workflows in Viral Molecular Epidemiology. <i>Viruses</i> , 2022, 14, 774.	3.3	2
13	FPGA Acceleration of Protein Back-Translation and Alignment. , 2021, , .		1
14	Ten simple rules for attending your first conference. <i>PLoS Computational Biology</i> , 2021, 17, e1009133.	3.2	1
15	SEPIA: simulation-based evaluation of prioritization algorithms. <i>BMC Medical Informatics and Decision Making</i> , 2021, 21, 177.	3.0	0