Cátia Vaz

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/514016/publications.pdf

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14	1,629	7	11
papers	citations	h-index	g-index
15	15	15	2622
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	GrapeTree: visualization of core genomic relationships among 100,000 bacterial pathogens. Genome Research, 2018, 28, 1395-1404.	5.5	553
2	PHYLOViZ: phylogenetic inference and data visualization for sequence based typing methods. BMC Bioinformatics, 2012, 13, 87.	2.6	492
3	PHYLOViZ 2.0: providing scalable data integration and visualization for multiple phylogenetic inference methods. Bioinformatics, 2017, 33, 128-129.	4.1	336
4	PHYLOViZ Online: web-based tool for visualization, phylogenetic inference, analysis and sharing of minimum spanning trees. Nucleic Acids Research, 2016, 44, W246-W251.	14.5	152
5	On the Expressive Power of Primitives for Compensation Handling. Lecture Notes in Computer Science, 2010, , 366-386.	1.3	26
6	Dynamic Recovering of Long Running Transactions. Lecture Notes in Computer Science, 2009, , 201-215.	1.3	20
7	Fast phylogenetic inference from typing data. Algorithms for Molecular Biology, 2018, 13, 4.	1.2	14
8	Distance-based phylogenetic inference from typing data: a unifying view. Briefings in Bioinformatics, 2021, 22, .	6.5	8
9	TypOn: the microbial typing ontology. Journal of Biomedical Semantics, 2014, 5, 43.	1.6	7
10	On the analysis of compensation correctness. The Journal of Logic and Algebraic Programming, 2012, 81, 585-605.	1.4	5
11	Towards Compensation Correctness in Interactive Systems. Lecture Notes in Computer Science, 2010, , 161-177.	1.3	4
12	Dynamic Phylogenetic Inference for Sequence-based Typing Data. , 2017, , .		2
13	NGSPipes., 2017,,.		0
14	An Ontology and a REST API for Sequence Based Microbial Typing Data. Lecture Notes in Computer Science, 2012, , 21-28.	1.3	0