Daniel Barker

List of Publications by Year in descending order

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1684188 1058476 17 220 5 14 citations h-index g-index papers 19 19 19 408 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Constrained models of evolution lead to improved prediction of functional linkage from correlated gain and loss of genes. Bioinformatics, 2007, 23, 14-20.	4.1	94
2	Xenolog classification. Bioinformatics, 2017, 33, 640-649.	4.1	42
3	4273Ï€: Bioinformatics education on low cost ARM hardware. BMC Bioinformatics, 2013, 14, 243.	2.6	19
4	On simulated annealing phase transitions in phylogeny reconstruction. Molecular Phylogenetics and Evolution, 2016, 101, 46-55.	2.7	11
5	Using sound to understand protein sequence data: new sonification algorithms for protein sequences and multiple sequence alignments. BMC Bioinformatics, 2021, 22, 456.	2.6	7
6	Seeing the wood for the trees: philosophical aspects of classical, Bayesian and likelihood approaches in statistical inference and some implications for phylogenetic analysis. Biology and Philosophy, 2015, 30, 505-525.	1.4	6
7	Comparison of the protein-coding genomes of three deep-sea, sulfur-oxidising bacteria: "Candidatus Ruthia magnificaâ€, "Candidatus Vesicomyosocius okutanii―and Thiomicrospira crunogena. BMC Research Notes, 2017, 10, 296.	1.4	5
8	DNA sonification for public engagement in bioinformatics. BMC Research Notes, 2021, 14, 273.	1.4	5
9	Genomic insights into the biosynthesis and physiology of the cyanobacterial neurotoxin 2,4-diaminobutanoic acid (2,4-DAB). Phytochemistry, 2021, 192, 112953.	2.9	5
10	Genomic insights into the biosynthesis and physiology of the cyanobacterial neurotoxin 3-N-methyl-2,3-diaminopropanoic acid (BMAA). Phytochemistry, 2022, 200, 113198.	2.9	5
11	Comparison of the protein-coding gene content of Chlamydia trachomatis and Protochlamydia amoebophila using a Raspberry Pi computer. BMC Research Notes, 2015, 8, 561.	1.4	4
12	Design, delivery and evaluation of a bioinformatics education workshop for 13-16-year-olds. Journal of Biological Education, 2022, 56, 570-580.	1.5	4
13	A comparison of the protein-coding genomes of two green sulphur bacteria, Chlorobium tepidum TLS and Pelodictyon phaeoclathratiforme BU-1. BMC Research Notes, 2015, 8, 565.	1.4	2
14	Bringing computational science to the public. SpringerPlus, 2016, 5, 259.	1.2	2
15	Bringing bioinformatics to schools with the 4273pi project. PLoS Computational Biology, 2022, 18, e1009705.	3.2	2
16	Cluster Analysis of p53 Binding Site Sequences Reveals Subsets with Different Functions. Cancer Informatics, 2016, 15, CIN.S39968.	1.9	1
17	A tribute to Peter Sneath. Taxon, 2012, 61, 481-483.	0.7	O