

Andre J Aberer

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10731830/publications.pdf>

Version: 2024-02-01

12
papers

4,949
citations

1040056

9
h-index

1372567

10
g-index

12
all docs

12
docs citations

12
times ranked

7622
citing authors

#	ARTICLE	IF	CITATIONS
1	An Efficient Independence Sampler for Updating Branches in Bayesian Markov chain Monte Carlo Sampling of Phylogenetic Trees. <i>Systematic Biology</i> , 2016, 65, 161-176.	5.6	11
2	ExaML version 3: a tool for phylogenomic analyses on supercomputers. <i>Bioinformatics</i> , 2015, 31, 2577-2579.	4.1	209
3	Phylogenomic analyses data of the avian phylogenomics project. <i>GigaScience</i> , 2015, 4, 4.	6.4	72
4	ExaBayes: Massively Parallel Bayesian Tree Inference for the Whole-Genome Era. <i>Molecular Biology and Evolution</i> , 2014, 31, 2553-2556.	8.9	423
5	Whole-genome analyses resolve early branches in the tree of life of modern birds. <i>Science</i> , 2014, 346, 1320-1331.	12.6	1,583
6	Phylogenomics resolves the timing and pattern of insect evolution. <i>Science</i> , 2014, 346, 763-767.	12.6	2,096
7	Decisive Data Sets in Phylogenomics: Lessons from Studies on the Phylogenetic Relationships of Primarily Wingless Insects. <i>Molecular Biology and Evolution</i> , 2014, 31, 239-249.	8.9	127
8	Novel Parallelization Schemes for Large-Scale Likelihood-based Phylogenetic Inference. , 2013, , .		29
9	Pruning Rogue Taxa Improves Phylogenetic Accuracy: An Efficient Algorithm and Webservice. <i>Systematic Biology</i> , 2013, 62, 162-166.	5.6	327
10	A Simple and Accurate Method for Rogue Taxon Identification. , 2011, , .		29
11	Uncovering Hidden Phylogenetic Consensus in Large Data Sets. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2011, 8, 902-911.	3.0	32
12	Parallelized phylogenetic post-analysis on multi-core architectures. <i>Journal of Computational Science</i> , 2010, 1, 107-114.	2.9	11