

Joseph Heled

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3546455/publications.pdf>

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14
papers

11,748
citations

687363

13
h-index

1125743

13
g-index

16
all docs

16
docs citations

16
times ranked

16892
citing authors

#	ARTICLE	IF	CITATIONS
1	Stay in Command: Optimal Play for Two Person Generala. <i>Recreational Mathematics Magazine</i> , 2020, 7, 53-70.	0.2	0
2	BEAST 2.5: An advanced software platform for Bayesian evolutionary analysis. <i>PLoS Computational Biology</i> , 2019, 15, e1006650.	3.2	2,484
3	Computational Performance and Statistical Accuracy of *BEAST and Comparisons with Other Methods. <i>Systematic Biology</i> , 2016, 65, 381-396.	5.6	107
4	Calibrated Birth-Death Phylogenetic Time-Tree Priors for Bayesian Inference. <i>Systematic Biology</i> , 2015, 64, 369-383.	5.6	48
5	Evaluating a multigene environmental DNA approach for biodiversity assessment. <i>GigaScience</i> , 2015, 4, 46.	6.4	122
6	BEAST 2: A Software Platform for Bayesian Evolutionary Analysis. <i>PLoS Computational Biology</i> , 2014, 10, e1003537.	3.2	5,301
7	Looking for trees in the forest: summary tree from posterior samples. <i>BMC Evolutionary Biology</i> , 2013, 13, 221.	3.2	131
8	Simulating gene trees under the multispecies coalescent and time-dependent migration. <i>BMC Evolutionary Biology</i> , 2013, 13, 44.	3.2	55
9	The Pipid Root. <i>Systematic Biology</i> , 2012, 61, 913-926.	5.6	49
10	Calibrated Tree Priors for Relaxed Phylogenetics and Divergence Time Estimation. <i>Systematic Biology</i> , 2012, 61, 138-149.	5.6	275
11	Sequence diversity under the multispecies coalescent with Yule process and constant population size. <i>Theoretical Population Biology</i> , 2012, 81, 97-101.	1.1	18
12	Bayesian Inference of Species Trees from Multilocus Data. <i>Molecular Biology and Evolution</i> , 2010, 27, 570-580.	8.9	2,246
13	Bayesian inference of population size history from multiple loci. <i>BMC Evolutionary Biology</i> , 2008, 8, 289.	3.2	658
14	The perils of plenty: what are we going to do with all these genes?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2008, 363, 3893-3902.	4.0	81