Joëlle Barido-Sottani

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4894955/publications.pdf

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1039880 1372474 2,857 11 9 10 citations h-index g-index papers 23 23 23 5296 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	BEAST 2.5: An advanced software platform for Bayesian evolutionary analysis. PLoS Computational Biology, 2019, 15, e1006650.	1.5	2,484
2	Taming the BEAST—A Community Teaching Material Resource for BEAST 2. Systematic Biology, 2018, 67, 170-174.	2.7	79
3	Ignoring stratigraphic age uncertainty leads to erroneous estimates of species divergence times under the fossilized birth–death process. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20190685.	1.2	50
4	A Multitype Birth–Death Model for Bayesian Inference of Lineage-Specific Birth and Death Rates. Systematic Biology, 2020, 69, 973-986.	2.7	44
5	Ignoring Fossil Age Uncertainty Leads to Inaccurate Topology and Divergence Time Estimates in Time Calibrated Tree Inference. Frontiers in Ecology and Evolution, 2020, 8, .	1.1	35
6	RevGadgets: An R package for visualizing Bayesian phylogenetic analyses from RevBayes. Methods in Ecology and Evolution, 2022, 13, 314-323.	2.2	33
7	<scp>FossilSim /scp>: An <scp>r /scp> package for simulating fossil occurrence data under mechanistic models of preservation and recovery. Methods in Ecology and Evolution, 2019, 10, 835-840.</scp></scp>	2.2	27
8	Detection of HIV transmission clusters from phylogenetic trees using a multi-state birth–death model. Journal of the Royal Society Interface, 2018, 15, 20180512.	1.5	22
9	Seven rules for simulations in paleobiology. Paleobiology, 2020, 46, 435-444.	1.3	15
10	The Occurrence Birth–Death Process for Combined-Evidence Analysis in Macroevolution and Epidemiology. Systematic Biology, 2022, 71, 1440-1452.	2.7	10
11	Lessons learned from organizing and teaching virtual phylogenetics workshops. , 2022, 1, .		O