## Joëlle Barido-Sottani

List of Publications by Year in descending order

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

Version: 2024-02-01

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	RevGadgets: An R package for visualizing Bayesian phylogenetic analyses from RevBayes. Methods in Ecology and Evolution, 2022, 13, 314-323.	2.2	33
2	The Occurrence Birth–Death Process for Combined-Evidence Analysis in Macroevolution and Epidemiology. Systematic Biology, 2022, 71, 1440-1452.	2.7	10
3	Lessons learned from organizing and teaching virtual phylogenetics workshops. , 2022, 1, .		O
4	Seven rules for simulations in paleobiology. Paleobiology, 2020, 46, 435-444.	1.3	15
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	A Multitype Birth–Death Model for Bayesian Inference of Lineage-Specific Birth and Death Rates. Systematic Biology, 2020, 69, 973-986.	2.7	44
7	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
8	<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.	2.2	27
9	BEAST 2.5: An advanced software platform for Bayesian evolutionary analysis. PLoS Computational Biology, 2019, 15, e1006650.	1.5	2,484
10	Taming the BEAST—A Community Teaching Material Resource for BEAST 2. Systematic Biology, 2018, 67, 170-174.	2.7	79
11	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