## Beatriz Mello

## List of Publications by Year in descending order

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1051969 721071 24 618 10 23 citations h-index g-index papers 26 26 26 1081 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Molecular dating of the blood pigment hemocyanin provides new insight into the origin of animals. Geobiology, 2022, 20, 333-345.	1.1	5
2	The performance of outgroup-free rooting under evolutionary radiations. Molecular Phylogenetics and Evolution, 2022, 169, 107434.	1.2	3
3	Molecular dating for phylogenies containing a mix of populations and species by using Bayesian and RelTime approaches. Molecular Ecology Resources, 2021, 21, 122-136.	2.2	18
4	Evolution of a key enzyme of aerobic metabolism reveals Proterozoic functional subunit duplication events and an ancient origin of animals. Scientific Reports, 2021, 11, 15744.	1.6	4
5	Reliable Confidence Intervals for RelTime Estimates of Evolutionary Divergence Times. Molecular Biology and Evolution, 2020, 37, 280-290.	3.5	36
6	A phylogenomic study of Steganinae fruit flies (Diptera: Drosophilidae): strong gene tree heterogeneity and evidence for monophyly. BMC Evolutionary Biology, 2020, 20, 141.	3.2	4
7	Employing statistical learning to derive speciesâ€level genetic diversity for mammalian species. Mammal Review, 2020, 50, 240-251.	2.2	10
8	The Estimated Pacemaker for Great Apes Supports the Hominoid Slowdown Hypothesis. Evolutionary Bioinformatics, 2019, 15, 117693431985598.	0.6	3
9	Comparative evaluation of macroevolutionary regimes of Ruminantia and selected mammalian lineages. Biological Journal of the Linnean Society, 2018, 123, 814-824.	0.7	1
10	Mode and Rate of Evolution of Haemosporidian Mitochondrial Genomes: Timing the Radiation of Avian Parasites. Molecular Biology and Evolution, 2018, 35, 383-403.	3.5	122
11	Impact of longâ€term chromosomal shuffling on the multispecies coalescent analysis of two anthropoid primate lineages. Ecology and Evolution, 2018, 8, 1206-1216.	0.8	2
12	Conservation phylogenetics and computational species delimitation of Neotropical primates. Biological Conservation, 2018, 217, 397-406.	1.9	11
13	Comparative evaluation of maximum parsimony and Bayesian phylogenetic reconstruction using empirical morphological data. Journal of Evolutionary Biology, 2018, 31, 1477-1484.	0.8	47
14	Multispecies coalescent analysis confirms standing phylogenetic instability in Hexapoda. Journal of Evolutionary Biology, 2018, 31, 1623-1631.	0.8	7
15	Estimating TimeTrees with MEGA and the TimeTree Resource. Molecular Biology and Evolution, 2018, 35, 2334-2342.	3.5	92
16	Fast and Accurate Estimates of Divergence Times from Big Data. Molecular Biology and Evolution, 2017, 34, 45-50.	3.5	52
17	Performance of Hidden Markov Models in Recovering the Standard Classification of Glycoside Hydrolases. Evolutionary Bioinformatics, 2017, 13, 117693431770340.	0.6	7
18	Analysis of Adaptive Evolution in Lyssavirus Genomes Reveals Pervasive Diversifying Selection during Species Diversification. Viruses, 2014, 6, 4465-4478.	1.5	4

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19	Sigmodontine rodents diversified in South America prior to the complete rise of the Panamanian Isthmus. Journal of Zoological Systematics and Evolutionary Research, 2014, 52, 249-256.	0.6	31
20	Assignment of Calibration Information to Deeper Phylogenetic Nodes is More Effective in Obtaining Precise and Accurate Divergence Time Estimates. Evolutionary Bioinformatics, 2014, 10, EBO.S13908.	0.6	18
21	Combining fossil and molecular data to date the diversification of New World Primates. Journal of Evolutionary Biology, 2013, 26, 2438-2446.	0.8	35
22	Phylogenetic analysis and a time tree for a large drosophilid data set (Diptera: Drosophilidae). Zoological Journal of the Linnean Society, 2013, 169, 765-775.	1.0	86
23	Phylogenetic analysis and a time tree for a large drosophilid data set (Diptera: Drosophilidae). Zoological Journal of the Linnean Society, 2013, , .	1.0	8
24	Incorrect handling of calibration information in divergence time inference: an example from volcanic islands. Ecology and Evolution, 2012, 2, 493-500.	0.8	10