

Ariadna E Morales

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

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1039406

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727
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene losses in the common vampire bat illuminate molecular adaptations to blood feeding. <i>Science Advances</i> , 2022, 8, eabm6494.	4.7	24
2	A New Dichromatic Species of <i>Myotis</i> (Chiroptera: Vespertilionidae) from the Nimba Mountains, Guinea. <i>American Museum Novitates</i> , 2021, 2020, .	0.2	1
3	Comment on "Population genetics reveal <i>Myotis keenii</i> (Keen's myotis) and <i>Myotis evotis</i> (long-eared myotis) to be a single species" Canadian Journal of Zoology, 2021, 99, 415-422.	0.4	5
4	Diversification rates have no effect on the convergent evolution of foraging strategies in the most speciose genus of bats, <i>Myotis</i> *. <i>Evolution; International Journal of Organic Evolution</i> , 2019, 73, 2263-2280.	1.1	40
5	Phylogeography, Population Structure, and Species Delimitation in Rockhopper Penguins (<i>Eudyptes</i>) Tj ETQq1 1 0.784314 rgBT /Overbo	1.0	3
6	Evidence that <i>Myotis lucifugus</i> "Subspecies" are Five Nonsister Species, Despite Gene Flow. <i>Systematic Biology</i> , 2018, 67, 756-769.	2.7	37
7	A global analysis of bats using automated comparative phylogeography uncovers a surprising impact of Pleistocene glaciation. <i>Journal of Biogeography</i> , 2018, 45, 1795-1805.	1.4	26
8	Genome-specific histories of divergence and introgression between an allopolyploid unisexual salamander lineage and two ancestral sexual species. <i>Evolution; International Journal of Organic Evolution</i> , 2018, 72, 1689-1700.	1.1	6
9	Spatial and environmental factors predict skull variation and genetic structure in the cosmopolitan bat <i>Tadarida brasiliensis</i> . <i>Journal of Biogeography</i> , 2018, 45, 1529-1540.	1.4	10
10	Species Delimitation with Gene Flow. <i>Systematic Biology</i> , 2017, 66, syw117.	2.7	118
11	Speciation with Gene Flow in North American <i>Myotis</i> Bats. <i>Systematic Biology</i> , 2017, 66, syw100.	2.7	50
12	PHRAPL: Phylogeographic Inference Using Approximate Likelihoods. <i>Systematic Biology</i> , 2017, 66, 1045-1053.	2.7	59
13	Objective choice of phylogeographic models. <i>Molecular Phylogenetics and Evolution</i> , 2017, 116, 136-140.	1.2	13
14	Environmental niche drives genetic and morphometric structure in a widespread bat. <i>Journal of Biogeography</i> , 2016, 43, 1057-1068.	1.4	17