

Nathan D Jackson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11822768/publications.pdf>

Version: 2024-02-01

15
papers

916
citations

759233

12
h-index

996975

15
g-index

15
all docs

15
docs citations

15
times ranked

1646
citing authors

#	ARTICLE	IF	CITATIONS
1	What determines the spatial extent of landscape effects on species?. <i>Landscape Ecology</i> , 2016, 31, 1177-1194.	4.2	194
2	Relative effects of road mortality and decreased connectivity on population genetic diversity. <i>Biological Conservation</i> , 2011, 144, 3143-3148.	4.1	169
3	Species Delimitation with Gene Flow. <i>Systematic Biology</i> , 2017, 66, syw117.	5.6	118
4	Habitat amount, not habitat configuration, best predicts population genetic structure in fragmented landscapes. <i>Landscape Ecology</i> , 2016, 31, 951-968.	4.2	97
5	Landscape context affects genetic diversity at a much larger spatial extent than population abundance. <i>Ecology</i> , 2014, 95, 871-881.	3.2	67
6	PHRAPL: Phylogeographic Inference Using Approximate Likelihoods. <i>Systematic Biology</i> , 2017, 66, 1045-1053.	5.6	59
7	Speciation with Gene Flow in North American <i>Myotis</i> Bats. <i>Systematic Biology</i> , 2017, 66, syw100.	5.6	50
8	THE COMBINED EFFECTS OF RIVERS AND REFUGIA GENERATE EXTREME CRYPTIC FRAGMENTATION WITHIN THE COMMON GROUND SKINK (<i>SCINCELLA LATERALIS</i>). <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 409-428.	2.3	49
9	Genome-Wide Analysis Reveals Mucociliary Remodeling of the Nasal Airway Epithelium Induced by Urban PM _{2.5} . <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2020, 63, 172-184.	2.9	32
10	Nasal airway transcriptome-wide association study of asthma reveals genetically driven mucus pathobiology. <i>Nature Communications</i> , 2022, 13, 1632.	12.8	24
11	The bioinvasion of Guam: inferring geographic origin, pace, pattern and process of an invasive lizard (<i>Carlia</i>) in the Pacific using multi-locus genomic data. <i>Biological Invasions</i> , 2011, 13, 1951-1967.	2.4	18
12	Inferring the evolutionary history of divergence despite gene flow in a lizard species, <i>Scincella lateralis</i> (Scincidae), composed of cryptic lineages. <i>Biological Journal of the Linnean Society</i> , 2012, 107, 192-209.	1.6	13
13	Objective choice of phylogeographic models. <i>Molecular Phylogenetics and Evolution</i> , 2017, 116, 136-140.	2.7	13
14	Testing the Role of Meander Cutoff in Promoting Gene Flow across a Riverine Barrier in Ground Skinks (<i>Scincella lateralis</i>). <i>PLoS ONE</i> , 2013, 8, e62812.	2.5	12
15	Microsatellites isolated from the North American ground skink (<i>Scincella lateralis</i>). <i>Conservation Genetics Resources</i> , 2011, 3, 95-97.	0.8	1