

# 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

758635

12  
h-index

996533

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.	1.9	194
2	Relative effects of road mortality and decreased connectivity on population genetic diversity. <i>Biological Conservation</i> , 2011, 144, 3143-3148.	1.9	169
3	Species Delimitation with Gene Flow. <i>Systematic Biology</i> , 2017, 66, syw117.	2.7	118
4	Habitat amount, not habitat configuration, best predicts population genetic structure in fragmented landscapes. <i>Landscape Ecology</i> , 2016, 31, 951-968.	1.9	97
5	Landscape context affects genetic diversity at a much larger spatial extent than population abundance. <i>Ecology</i> , 2014, 95, 871-881.	1.5	67
6	PHRAPL: Phylogeographic Inference Using Approximate Likelihoods. <i>Systematic Biology</i> , 2017, 66, 1045-1053.	2.7	59
7	Speciation with Gene Flow in North American <i>Myotis</i> Bats. <i>Systematic Biology</i> , 2017, 66, syw100.	2.7	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.	1.1	49
9	Genome-Wide Analysis Reveals Mucociliary Remodeling of the Nasal Airway Epithelium Induced by Urban PM <sub>2.5</sub> . <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2020, 63, 172-184.	1.4	32
10	Nasal airway transcriptome-wide association study of asthma reveals genetically driven mucus pathobiology. <i>Nature Communications</i> , 2022, 13, 1632.	5.8	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.	1.2	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.	0.7	13
13	Objective choice of phylogeographic models. <i>Molecular Phylogenetics and Evolution</i> , 2017, 116, 136-140.	1.2	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.	1.1	12
15	Microsatellites isolated from the North American ground skink ( <i>Scincella lateralis</i> ). <i>Conservation Genetics Resources</i> , 2011, 3, 95-97.	0.4	1