Tobias Andermann

List of Publications by Year
in descending order

16	1,027	949033 11	1051228 16
papers	citations	h-index	g-index

27	27	27	2006
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Target sequence capture of Barnadesioideae (Compositae) demonstrates the utility of low coverage loci in phylogenomic analyses. Molecular Phylogenetics and Evolution, 2022, 169, 107432.	1.2	9
2	IUCNN â€“ Deep learning approaches to approximate species' extinction risk. Diversity and Distributions, 2022, 28, 227-241.	1.9	19
3	Global Estimation and Mapping of the Conservation Status of Tree Species Using Artificial Intelligence. Frontiers in Plant Science, 2022, 13, 839792.	1.7	13
4	Recent and local diversification of Central American understorey palms. Global Ecology and Biogeography, 2022, 31, 1513-1525.	2.7	3
5	iucn_sim: a new program to simulate future extinctions based on IUCN threat status. Ecography, 2021, 44, 162-176.	2.1	17
6	Genomic and niche divergence in an Amazonian palm species complex. Botanical Journal of the Linnean Society, 2021, 197, 498-512.	0.8	8
7	Phylogenomics of the tropical plant family Ochnaceae using targeted enrichment of nuclear genes and 250+ taxa. Taxon, 2021, 70, 48-71.	0.4	14
8	A bioinformatic platform to integrate target capture and whole genome sequences of various read depths for phylogenomics. Molecular Ecology, 2021, 30, 6021-6035.	2.0	10
9	The past and future human impact on mammalian diversity. Science Advances, 2020, 6, .	4.7	91
10	Phylogenomics and biogeography of the world's thrushes (Aves,Turdus): new evidence for a more parsimonious evolutionary history. Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20192400.	1.2	16
11	Allele Phasing Greatly Improves the Phylogenetic Utility of Ultraconserved Elements. Systematic Biology, 2019, 68, 32-46.	2.7	74
12	<scp>CoordinateCleaner</scp>: Standardized cleaning of occurrence records from biological collection databases. Methods in Ecology and Evolution, 2019, 10, 744-751.	2.2	473
13	A Guide to Carrying Out a Phylogenomic Target Sequence Capture Project. Frontiers in Genetics, 2019, 10, 1407.	1.1	76
14	Estimating Age-Dependent Extinction: Contrasting Evidence from Fossils and Phylogenies. Systematic Biology, 2018, 67, 458-474.	2.7	32
15	SECAPRâ€”a bioinformatics pipeline for the rapid and user-friendly processing of targeted enriched Illumina sequences, from raw reads to alignments. PeerJ, 2018, 6, e5175.	0.9	52
16	Conceptual and empirical advances in Neotropical biodiversity research. PeerJ, 2018, 6, e5644.	0.9	107