## Deren A R Eaton

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5230270/publications.pdf

Version: 2024-02-01

394421 477307 3,356 29 19 29 citations h-index g-index papers 37 37 37 3914 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	PyRAD: assembly of <i>de novo</i> RADseq loci for phylogenetic analyses. Bioinformatics, 2014, 30, 1844-1849.	4.1	708
2	Inferring Phylogeny and Introgression using RADseq Data: An Example from Flowering Plants (Pedicularis: Orobanchaceae). Systematic Biology, 2013, 62, 689-706.	5 <b>.</b> 6	482
3	ipyrad: Interactive assembly and analysis of RADseq datasets. Bioinformatics, 2020, 36, 2592-2594.	4.1	479
4	A Framework Phylogeny of the American Oak Clade Based on Sequenced RAD Data. PLoS ONE, 2014, 9, e93975.	2.5	215
5	Historical introgression among the American live oaks and the comparative nature of tests for introgression. Evolution; International Journal of Organic Evolution, 2015, 69, 2587-2601.	2.3	193
6	Misconceptions on Missing Data in RAD-seq Phylogenetics with a Deep-scale Example from Flowering Plants. Systematic Biology, 2017, 66, syw092.	5.6	167
7	Phylogeny and biogeography of the American live oaks ( <i>Quercus</i> subsection <i>Virentes</i> ): a genomic and population genetics approach. Molecular Ecology, 2015, 24, 3668-3687.	3.9	165
8	Practical considerations for plant phylogenomics. Applications in Plant Sciences, 2018, 6, e1038.	2.1	165
9	Genotyping-by-sequencing as a tool to infer phylogeny and ancestral hybridization: A case study in Carex (Cyperaceae). Molecular Phylogenetics and Evolution, 2014, 79, 359-367.	2.7	115
10	Paths to selection on life history loci in different natural environments across the native range of <i><i>&gt;Arabidopsis thalianaMolecular Ecology, 2013, 22, 3552-3566.</i></i>	3.9	101
11	Floral diversity and community structure in <i>Pedicularis</i> (Orobanchaceae). Ecology, 2012, 93, S182.	3.2	96
12	Coral hybridization or phenotypic variation? Genomic data reveal gene flow between Porites lobata and P. Compressa. Molecular Phylogenetics and Evolution, 2017, 111, 132-148.	2.7	59
13	Identification of SNP markers for inferring phylogeny in temperate bamboos (Poaceae: Bambusoideae) using RAD sequencing. Molecular Ecology Resources, 2013, 13, 938-945.	4.8	53
14	Toytree: A minimalist tree visualization and manipulation library for Python. Methods in Ecology and Evolution, 2020, 11, 187-191.	5.2	48
15	Parallel ddRAD and Genome Skimming Analyses Reveal a Radiative and Reticulate Evolutionary History of the Temperate Bamboos. Systematic Biology, 2021, 70, 756-773.	5 <b>.</b> 6	38
16	Restriction-Site-Associated DNA Sequencing Reveals a Cryptic <b><i>Viburnum</i></b> Species on the North American Coastal Plain. Systematic Biology, 2019, 68, 187-203.	5.6	36
17	Phylogeny of Hawaiian Melicope (Rutaceae): RAD-seq Resolves Species Relationships and Reveals Ancient Introgression. Frontiers in Plant Science, 2019, 10, 1074.	3.6	35
18	Joint Phylogenetic Estimation of Geographic Movements and Biome Shifts during the Global Diversification of <i>Viburnum </i> . Systematic Biology, 2021, 70, 67-85.	5 <b>.</b> 6	33

#	Article	IF	CITATIONS
19	Inferring processes of coevolutionary diversification in a community of Panamanian strangler figs and associated pollinating wasps*. Evolution; International Journal of Organic Evolution, 2019, 73, 2295-2311.	2.3	30
20	Genome-Wide Assessment of Diversity and Divergence Among Extant Galapagos Giant Tortoise Species. Journal of Heredity, 2018, 109, 611-619.	2.4	22
21	Differences in flowering time maintain species boundaries in a continental radiation of <i>Viburnum</i> . American Journal of Botany, 2019, 106, 833-849.	1.7	19
22	The potential of genome-wide RAD sequences for resolving rapid radiations: a case study in Cactaceae. Molecular Phylogenetics and Evolution, 2020, 151, 106896.	2.7	16
23	Reconciling species diversity in a tropical plant clade (Canarium, Burseraceae). PLoS ONE, 2018, 13, e0198882.	2.5	13
24	Replicated radiation of a plant clade along a cloud forest archipelago. Nature Ecology and Evolution, 2022, 6, 1318-1329.	7.8	11
25	Resolved phylogenetic relationships in the <i>Ocotea</i> complex ( <i>Supraocotea</i> ) facilitate phylogenetic classification and studies of character evolution. American Journal of Botany, 2021, 108, 664-679.	1.7	10
26	Tracking the xeric biomes of South America: The spatiotemporal diversification of Mandacaru cactus. Journal of Biogeography, 2021, 48, 3085-3103.	3.0	10
27	Sterile marginal flowers increase visitation and fruit set in the hobblebush ( <i>Viburnum) Tj ETQq1 1 0.784314 r</i>	gBŢ ĮOver	lock 10 Tf 50
28	Selection on the gametophyte: Modeling alternation of generations in plants. Applications in Plant Sciences, 2022, 10, e11472.	2.1	7
29	ipcoal: an interactive Python package for simulating and analyzing genealogies and sequences on a species tree or network. Bioinformatics, 2020, 36, 4193-4196.	4.1	6