Paul S Manos

List of Publications by Year
in descending order

43 papers	5,143 citations	136950 32 h-index	254184 43 g-index
47 all docs	47 docs citations	47 times ranked	5094 citing authors

#	Article	IF	Citations
1	Phylogenomic analyses highlight innovation and introgression in the continental radiations of Fagaceae across the Northern Hemisphere. Nature Communications, 2022, 13, 1320.	12.8	43
2	An Updated Infrageneric Classification of the North American Oaks (Quercus Subgenus Quercus): Review of the Contribution of Phylogenomic Data to Biogeography and Species Diversity. Forests, 2021, 12, 786.	2.1	27
3	Uncovering the genomic signature of ancient introgression between white oak lineages (Quercus). New Phytologist, 2020, 226, 1158-1170.	7.3	63
4	Genomic landscape of the global oak phylogeny. New Phytologist, 2020, 226, 1198-1212.	7.3	186
5	Comment on â€œEocene Fagaceae from Patagonia and Gondwanan legacy in Asian rainforestsâ€• Science, 2019, 366, .	12.6	3
6	Genomic Identity of White Oak Species in an Eastern North American Syngameon. Annals of the Missouri Botanical Garden, 2019, 104, 455-477.	1.3	22
7	The role of diversification in community assembly of the oaks (Quercus L.) across the continental U.S.. American Journal of Botany, 2018, 105, 565-586.	1.7	50
8	Geographic range velocity and its association with phylogeny and life history traits in North American woody plants. Ecology and Evolution, 2018, 8, 2632-2644.	1.9	9
9	Pervasive migration across rainforest and sandy coastal plain Aechmea nudicaulis (Bromeliaceae) populations despite contrasting environmental conditions. Molecular Ecology, 2018, 27, 1261-1272.	3.9	8
10	Sympatric parallel diversification of major oak clades in the Americas and the origins of Mexican species diversity. New Phytologist, 2018, 217, 439-452.	7.3	216
11	Phylogeny and biogeography of East Asian evergreen oaks (Quercus section Cyclobalanopsis;) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T Molecular Phylogenetics and Evolution, 2018, 119, 170-181.	2.7	99
12	A time and a place for everything: phylogenetic history and geography as joint predictors of oak plastome phylogeny. Genome, 2017, 60, 720-732.	2.0	64
13	A genetic legacy of introgression confounds phylogeny and biogeography in oaks. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20170300.	2.6	105
14	Phylogenomic inferences from reference-mapped and de novo assembled short-read sequence data using RADseq sequencing of California white oaks (Quercus section Quercus). Genome, 2017, 60, 743-755.	2.0	50
15	The evolution and diversification of the red oaks of the California Floristic Province (Quercus) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	1.7	27
16	Phylogenomics reveals a complex evolutionary history of lobed-leaf white oaks in western North America. Genome, 2017, 60, 733-742.	2.0	39
17	An Updated Infrageneric Classification of the Oaks: Review of Previous Taxonomic Schemes and Synthesis of Evolutionary Patterns. Tree Physiology, 2017, , 13-38.	2.5	117
18	Evidence for genetic erosion of a California native tree, Platanus racemosa, via recent, ongoing introgressive hybridization with an introduced ornamental species. Conservation Genetics, 2016, 17, 593-602.	1.5	14

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19	Phylogeny and biogeography of the American live oaks (Quercus subsection Virentes): a genomic and population genetics approach. Molecular Ecology, 2015, 24, 3668-3687.	3.9	165
20	Biogeographic analysis of the woody plants of the Southern Appalachians: Implications for the origins of a regional flora. American Journal of Botany, 2015, 102, 780-804.	1.7	39
21	A Framework Phylogeny of the American Oak Clade Based on Sequenced RAD Data. PLoS ONE, 2014, 9, e93975.	2.5	215
22	Genomics of Fagaceae. Tree Genetics and Genomes, 2012, 8, 583-610.	1.6	109
23	Diversification ofCeanothus(Rhamnaceae) in the California Floristic Province. International Journal of Plant Sciences, 2011, 172, 1137-1164.	1.3	36
24	Host density drives the postglacial migration of the tree parasite, Epifagus virginiana. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 17035-17040.	7.1	40
25	Natural history, distribution, phylogenetic relationships, and conservation of Central American black walnuts (Juglans sect. Rhysocaryon)1. Journal of the Torrey Botanical Society, 2009, 136, 1-25.	0.3	32
26	IS FLORAL SPECIALIZATION AN EVOLUTIONARY DEADâ€END? POLLINATION SYSTEM TRANSITIONS INRUELLIA(ACANTHACEAE). Evolution; International Journal of Organic Evolution, 2008, 62, 1712-1737.	2.3	199
27	Inferring longâ€distance dispersal and topographic barriers during postâ€glacial colonization from the genetic structure of red maple (Acer rubrum L.) in New England. Journal of Biogeography, 2008, 35, 1665-1673.	3.0	18
28	Phylogenetic Relationships and Taxonomic Status Of the Paleoendemic Fagaceae Of Western North America: Recognition Of A New Genus, Notholithocarpus. MadroÃ±o, 2008, 55, 181-190.	0.4	92
29	Phylogeny of Extant and Fossil Juglandaceae Inferred from the Integration of Molecular and Morphological Data Sets. Systematic Biology, 2007, 56, 412-430.	5.6	127
30	Comparative phylogeography of unglaciated eastern North America. Molecular Ecology, 2006, 15, 4261-4293.	3.9	843
31	Phylogeny and Historical Biogeography of the Genus <I>Platanus</I> as Inferred From Nuclear and Chloroplast DNA. Systematic Botany, 2005, 30, 786-799.	0.5	45
32	MOLECULAR INDICATORS OF TREE MIGRATION CAPACITY UNDER RAPID CLIMATE CHANGE. Ecology, 2005, 86, 2088-2098.	3.2	502
33	Phylogenetic Relationships in Fagales Based on DNA Sequences fromÂThree Genomes. International Journal of Plant Sciences, 2004, 165, 311-324.	1.3	110
34	Phylogeography of the Southeast Asian stone oaks (Lithocarpus). Journal of Biogeography, 2003, 30, 211-226.	3.0	122
35	The Historical Biogeography of Fagaceae: Tracking the Tertiary History of Temperate and Subtropical Forests of the Northern Hemisphere. International Journal of Plant Sciences, 2001, 162, S77-S93.	1.3	237
36	Molecular and morphological analyses of Bryopsis (Bryopsidales, Chlorophyta) from the western North Atlantic and Caribbean. Phycologia, 2001, 40, 330-339.	1.4	18

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37	Evolution, Phylogeny, and Systematics of the Juglandaceae. Annals of the Missouri Botanical Garden, 2001, 88, 231.	1.3	110
38	Combining and Comparing Morphometric Shape Descriptors with a Molecular Phylogeny: The Case of Fruit Type Evolution in Bornean Lithocarpus (Fagaceae). Systematic Biology, 2001, 50, 860-880.	5.6	48
39	Phylogeny and patterns of floral diversity in the genus Piper (Piperaceae). American Journal of Botany, 2001, 88, 706-716.	1.7	189
40	Phylogeny, Biogeography, and Processes of Molecular Differentiation in Quercus Subgenus Quercus (Fagaceae). Molecular Phylogenetics and Evolution, 1999, 12, 333-349.	2.7	353
41	Phylogenetic analyses of â€œhigherâ€•Hamamelididae based on plastid sequence data. American Journal of Botany, 1997, 84, 1407-1419.	1.7	130
42	Systematics of Nothofagus (Nothofagaceae) based on rDNA spacer sequences (ITS): taxonomic congruence with morphology and plastid sequences. American Journal of Botany, 1997, 84, 1137-1155.	1.7	152
43	Allozyme Variation in Populations of Six Northeastern American Red Oaks (Fagaceae: Quercus Subg.) Tj ETQq1 1 0.784314 rgBT /Ove	0.5	56