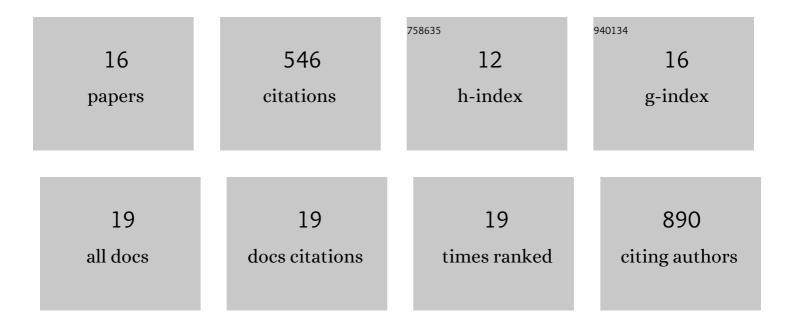
Richard G J Hodel

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3782488/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Concordance-Based Approaches for the Inference of Relationships and Molecular Rates with Phylogenomic Data Sets. Systematic Biology, 2022, 71, 943-958.	2.7	11
2	<i>Amborella</i> gene presence/absence variation is associated with abiotic stress responses that may contribute to environmental adaptation. New Phytologist, 2022, 233, 1548-1555.	3.5	16
3	Phylogenomic conflict analyses in the apple genus <i>Malus</i> s.l. reveal widespread hybridization and allopolyploidy driving diversification, with insights into the complex biogeographic history in the Northern Hemisphere. Journal of Integrative Plant Biology, 2022, 64, 1020-1043.	4.1	31
4	Hybrid enrichment of adaptive variation revealed by genotype–environment associations in montane sedges. Molecular Ecology, 2022, 31, 3722-3737.	2.0	7
5	A phylogenomic approach resolves the backbone of Prunus (Rosaceae) and identifies signals of hybridization and allopolyploidy. Molecular Phylogenetics and Evolution, 2021, 160, 107118.	1.2	30
6	Capturing singleâ€copy nuclear genes, organellar genomes, and nuclear ribosomal DNA from deep genome skimming data for plant phylogenetics: A case study in Vitaceae. Journal of Systematics and Evolution, 2021, 59, 1124-1138.	1.6	43
7	Testing which axes of species differentiation underlie covariance of phylogeographic similarity among montane sedge species. Evolution; International Journal of Organic Evolution, 2021, 75, 349-364.	1.1	8
8	Synthesis of Nuclear and Chloroplast Data Combined With Network Analyses Supports the Polyploid Origin of the Apple Tribe and the Hybrid Origin of the Maleae—Gillenieae Clade. Frontiers in Plant Science, 2021, 12, 820997.	1.7	16
9	Evolutionary history of a relict conifer, Pseudotaxus chienii (Taxaceae), in south-east China during the late Neogene: old lineage, young populations. Annals of Botany, 2020, 125, 105-117.	1.4	27
10	Linking genome signatures of selection and adaptation in non-model plants: exploring potential and limitations in the angiosperm Amborella. Current Opinion in Plant Biology, 2018, 42, 81-89.	3.5	4
11	Terrestrial species adapted to sea dispersal: Differences in propagule dispersal of two Caribbean mangroves. Molecular Ecology, 2018, 27, 4612-4626.	2.0	25
12	Dispersal corridors for plant species in the Poyang Lake Basin of southeast China identified by integration of phylogeographic and geospatial data. Ecology and Evolution, 2017, 7, 5140-5148.	0.8	16
13	Adding loci improves phylogeographic resolution in red mangroves despite increased missing data: comparing microsatellites and RAD-Seq and investigating loci filtering. Scientific Reports, 2017, 7, 17598.	1.6	99
14	Comparative phylogeography of black mangroves (<i>Avicennia germinans</i>) and red mangroves (<i>Rhizophora mangle</i>) in Florida: Testing the maritime discontinuity in coastal plants. American Journal of Botany, 2016, 103, 730-739.	0.8	24
15	The report of my death was an exaggeration: A review for researchers using microsatellites in the 21st century. Applications in Plant Sciences, 2016, 4, 1600025.	0.8	155
16	A new resource for the development of SSR markers: Millions of loci from a thousand plant transcriptomes. Applications in Plant Sciences, 2016, 4, 1600024.	0.8	29