Jim Provan

List of Publications by Year in descending order

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687363 610901 2,068 24 13 citations h-index papers

g-index 24 24 24 3567 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	Strong spatial structuring of clonal genetic diversity within blackthorn (Prunus spinosa) hedgerows and woodlands. Tree Genetics and Genomes, 2022, 18 , 1 .	1.6	1
2	Hierarchical structuring of genetic variation at differing geographic scales in the cultivated sugar kelp Saccharina latissima. Marine Environmental Research, 2018, 142, 108-115.	2.5	9
3	Evidence for facultative protocarnivory in Capsella bursa-pastoris seeds. Scientific Reports, 2018, 8, 10120.	3.3	6
4	Low genetic diversity and potential inbreeding in an isolated population of alder buckthorn (Frangula) Tj ETQq0	0 0 rgBT /0	Overlock 10 Tf
5	Using genetic monitoring to inform best practice in a captive breeding programme: inbreeding and potential genetic rescue in the freshwater pearl mussel Margaritifera margaritifera. Conservation Genetics, 2016, 17, 1323-1332.	1.5	6
6	High-resolution genetic analysis reveals extensive gene flow within the jellyfish <i>Pelagia noctiluca</i> (Scyphozoa) in the North Atlantic and Mediterranean Sea. Biological Journal of the Linnean Society, 2016, 117, 252-263.	1.6	7
7	Broad-scale genetic homogeneity in natural populations of common hazel (Corylus avellana) in Ireland. Tree Genetics and Genomes, 2016, 12, 1.	1.6	10
8	Pure species in a continuum of genetic and morphological variation: sympatric oaks at the edge of their range. Annals of Botany, 2016, 117, 541-549.	2.9	21
9	Genetic analyses reveal high levels of seed and pollen flow in hawthorn (Crataegus monogyna Jacq.), a key component of hedgerows. Tree Genetics and Genomes, 2016, 12, 1.	1.6	16
10	Population genetic analyses reveal distinct geographical blooms of the jellyfish <i>Rhizostoma octopus </i> (Scyphozoa). Biological Journal of the Linnean Society, 2015, 116, 582-592.	1.6	12
11	Lack of genetic structure and evidence for long-distance dispersal in ash (Fraxinus excelsior) populations under threat from an emergent fungal pathogen: implications for restorative planting. Tree Genetics and Genomes, 2015, 11, 1.	1.6	18
12	Genetic provenance and best practice woodland management: a case study in native alder (Alnus) Tj ETQq0 0 0	rgBT/Ovei	rlock 10 Tf 50
13	Cryptic introgression into the kidney saxifrage (Saxifraga hirsuta) from its more abundant sympatric congener Saxifraga spathularis, and the potential risk of genetic assimilation. Annals of Botany, 2015, 115, 179-186.	2.9	13
14	Phylogeographical analyses of shellfish viruses: inferring a geographical origin for ostreid herpesviruses OsHV-1 (Malacoherpesviridae). Marine Biology, 2015, 162, 181-192.	1.5	31
15	The not-so-Irish spurge:Euphorbia hyberna(Euphorbiaceae) and the Littletonian plant â€~steeplechase'. Biological Journal of the Linnean Society, 2015, 114, 249-259.	1.6	6
16	Understanding macroalgal dispersal in a complex hydrodynamic environment: a combined population genetic and physical modelling approach. Journal of the Royal Society Interface, 2014, 11, 20140197.	3.4	25
17	Phylogeographical analysis of two coldâ€tolerant plants with disjunct Lusitanian distributions does not support <i>in situ</i> survival during the last glaciation. Journal of Biogeography, 2014, 41, 2185-2193.	3.0	23
18	Retrospective genetic monitoring of the threatened Yellow marsh saxifrage (<i><scp>S</scp>axifraga) Tj ETQq0</i>	0 0 rgBT / 4.1	Overlock 10 T 6

Diversity and Distributions, 2014, 20, 529-537.

#	Article	IF	CITATIONS
19	Restricted gene flow in fragmented populations of a wind-pollinated tree. Conservation Genetics, 2008, 9, 1521-1532.	1.5	61
20	Phylogeographic insights into cryptic glacial refugia. Trends in Ecology and Evolution, 2008, 23, 564-571.	8.7	930
21	Universal plastid primers for Chlorophyta and Rhodophyta. European Journal of Phycology, 2004, 39, 43-50.	2.0	40
22	Chloroplast microsatellites: new tools for studies in plant ecology and evolution. Trends in Ecology and Evolution, 2001, 16, 142-147.	8.7	587
23	Analysis of the genusZea (Poaceae) using polymorphic chloroplast simple sequence repeats. Plant Systematics and Evolution, 1999, 218, 245-256.	0.9	18
24	A Low Mutation Rate For Chloroplast Microsatellites. Genetics, 1999, 153, 943-947.	2.9	197