

Andrew J Lowe

List of Publications by Year in descending order

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Version: 2024-02-01

138
papers

10,264
citations

50276

46
h-index

38395

95
g-index

145
all docs

145
docs citations

145
times ranked

14021
citing authors

#	ARTICLE	IF	CITATIONS
1	Adaptive evolution in invasive species. <i>Trends in Plant Science</i> , 2008, 13, 288-294.	8.8	724
2	Something in the way you move: dispersal pathways affect invasion success. <i>Trends in Ecology and Evolution</i> , 2009, 24, 136-144.	8.7	680
3	Building evolutionary resilience for conserving biodiversity under climate change. <i>Evolutionary Applications</i> , 2011, 4, 326-337.	3.1	617
4	Identification of refugia and post-glacial colonisation routes of European white oaks based on chloroplast DNA and fossil pollen evidence. <i>Forest Ecology and Management</i> , 2002, 156, 49-74.	3.2	577
5	Seed supply for broadscale restoration: maximizing evolutionary potential. <i>Evolutionary Applications</i> , 2008, 1, 587-597.	3.1	495
6	Chloroplast DNA variation in European white oaks. <i>Forest Ecology and Management</i> , 2002, 156, 5-26.	3.2	424
7	Value of long-term ecological studies. <i>Austral Ecology</i> , 2012, 37, 745-757.	1.5	326
8	The extent of forest in dryland biomes. <i>Science</i> , 2017, 356, 635-638.	12.6	300
9	Which provenance and where? Seed sourcing strategies for revegetation in a changing environment. <i>Conservation Genetics</i> , 2013, 14, 1-10.	1.5	290
10	Rise of the machines – recommendations for ecologists when using next generation sequencing for microsatellite development. <i>Molecular Ecology Resources</i> , 2011, 11, 1093-1101.	4.8	228
11	Introducing BASE: the Biomes of Australian Soil Environments soil microbial diversity database. <i>GigaScience</i> , 2016, 5, 21.	6.4	204
12	Effective Seed Dispersal Across a Fragmented Landscape. <i>Science</i> , 2006, 311, 628-628.	12.6	201
13	Origins, establishment and evolution of new polyploid species: <i>Senecio cambrensis</i> and <i>S. jefeboracensis</i> in the British Isles. <i>Biological Journal of the Linnean Society</i> , 2004, 82, 467-474.	1.6	189
14	Forensic timber identification: It's time to integrate disciplines to combat illegal logging. <i>Biological Conservation</i> , 2015, 191, 790-798.	4.1	176
15	Leaf morphological differentiation between <i>Quercus robur</i> and <i>Quercus petraea</i> is stable across western European mixed oak stands. <i>Annals of Forest Science</i> , 2002, 59, 777-787.	2.0	161
16	Improving biodiversity monitoring. <i>Austral Ecology</i> , 2012, 37, 285-294.	1.5	130
17	Urban habitat restoration provides a human health benefit through microbiome rewilding: the Microbiome Rewilding Hypothesis. <i>Restoration Ecology</i> , 2017, 25, 866-872.	2.9	129
18	Leaf morphology shift linked to climate change. <i>Biology Letters</i> , 2012, 8, 882-886.	2.3	127

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19	Constraints to and conservation implications for climate change adaptation in plants. <i>Conservation Genetics</i> , 2016, 17, 305-320.	1.5	122
20	HISTORICAL AND CONTEMPORARY MATING PATTERNS IN REMNANT POPULATIONS OF THE FOREST TREE <i>FRAXINUS EXCELSIOR</i> L.. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 979-990.	2.3	114
21	Refugia within refugia: the case study of a canopy tree (<i>Eurycorymbus cavaleriei</i>) in subtropical China. <i>Journal of Biogeography</i> , 2009, 36, 2156-2164.	3.0	111
22	Genomic evidence for the parallel evolution of coastal forms in the <i>Senecio laetus</i> complex. <i>Molecular Ecology</i> , 2013, 22, 2941-2952.	3.9	109
23	Bridging the gap: a genetic assessment framework for population-level threatened plant conservation prioritization and decision-making. <i>Diversity and Distributions</i> , 2016, 22, 174-188.	4.1	105
24	Transfer of environmental microbes to the skin and respiratory tract of humans after urban green space exposure. <i>Environment International</i> , 2020, 145, 106084.	10.0	103
25	Is there a correlation between chloroplastic and nuclear divergence, or what are the roles of history and selection on genetic diversity in European oaks?. <i>Forest Ecology and Management</i> , 2002, 156, 75-87.	3.2	101
26	Plant DNA Barcodes Can Accurately Estimate Species Richness in Poorly Known Floras. <i>PLoS ONE</i> , 2011, 6, e26841.	2.5	100
27	Comparative phylogeography of African rain forest trees: A review of genetic signatures of vegetation history in the Guineo-Congolian region. <i>Comptes Rendus - Geoscience</i> , 2013, 345, 284-296.	1.2	94
28	Changing perspectives on the biogeography of the tropical South Pacific: influences of dispersal, vicariance and extinction. <i>Journal of Biogeography</i> , 2009, 36, 1035-1054.	3.0	91
29	Transferability and genome specificity of a new set of microsatellite primers among Brassica species of the U triangle. <i>Molecular Ecology Notes</i> , 2002, 2, 7-11.	1.7	90
30	Leaf evolution in Southern Hemisphere conifers tracks the angiosperm ecological radiation. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 341-348.	2.6	88
31	Priority Actions to Improve Provenance Decision-Making. <i>BioScience</i> , 2018, 68, 510-516.	4.9	87
32	High-throughput eDNA monitoring of fungi to track functional recovery in ecological restoration. <i>Biological Conservation</i> , 2018, 217, 113-120.	4.1	81
33	Leaf nitrogen from first principles: field evidence for adaptive variation with climate. <i>Biogeosciences</i> , 2017, 14, 481-495.	3.3	75
34	High Nuclear Genetic Diversity, High Levels of Outcrossing and Low Differentiation Among Remnant Populations of <i>Quercus petraea</i> at the Margin of its Range in Ireland. <i>Annals of Botany</i> , 2004, 93, 691-697.	2.9	74
35	A landscape genetics approach for quantifying the relative influence of historic and contemporary habitat heterogeneity on the genetic connectivity of a rainforest bird. <i>Molecular Ecology</i> , 2009, 18, 2945-2960.	3.9	70
36	Revegetation rewilds the soil bacterial microbiome of an old field. <i>Molecular Ecology</i> , 2017, 26, 2895-2904.	3.9	68

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37	CONVERGENCE AND DIVERGENCE DURING THE ADAPTATION TO SIMILAR ENVIRONMENTS BY AN AUSTRALIAN GROUNDSEL. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 2515-2529.	2.3	66
38	Testing the role of genetic factors across multiple independent invasions of the shrub Scotch broom (<i>Cytisus scoparius</i>). <i>Molecular Ecology</i> , 2007, 16, 4662-4673.	3.9	64
39	Relating Urban Biodiversity to Human Health With the "Holobiont"™ Concept. <i>Frontiers in Microbiology</i> , 2019, 10, 550.	3.5	64
40	Opportunities for Improved Transparency in the Timber Trade through Scientific Verification. <i>BioScience</i> , 2016, 66, 990-998.	4.9	60
41	Pollen diversity matters: revealing the neglected effect of pollen diversity on fitness in fragmented landscapes. <i>Molecular Ecology</i> , 2012, 21, 5955-5968.	3.9	57
42	Genetic diversity and structure of the Australian flora. <i>Diversity and Distributions</i> , 2017, 23, 41-52.	4.1	56
43	Did Kauri (<i>Agathis</i> : <i>Araucariaceae</i>) Really Survive the Oligocene Drowning of New Zealand?. <i>Systematic Biology</i> , 2010, 59, 594-602.	5.6	55
44	Shifts in reproductive assurance strategies and inbreeding costs associated with habitat fragmentation in Central American mahogany. <i>Ecology Letters</i> , 2012, 15, 444-452.	6.4	55
45	Advancing DNA Barcoding and Metabarcoding Applications for Plants Requires Systematic Analysis of Herbarium Collections—An Australian Perspective. <i>Frontiers in Ecology and Evolution</i> , 2018, 6, .	2.2	55
46	Bacterial natural product biosynthetic domain composition in soil correlates with changes in latitude on a continent-wide scale. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 11615-11620.	7.1	53
47	The founding charter of the Genomic Observatories Network. <i>GigaScience</i> , 2014, 3, 2.	6.4	51
48	Using phylogenetic diversity to identify ancient rain forest refugia and diversification zones in a biodiversity hotspot. <i>Diversity and Distributions</i> , 2015, 21, 279-289.	4.1	50
49	Landscape biodiversity correlates with respiratory health in Australia. <i>Journal of Environmental Management</i> , 2018, 206, 113-122.	7.8	50
50	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 June 2010 – 31 July 2010. <i>Molecular Ecology Resources</i> , 2010, 10, 1106-1108.	4.8	48
51	Predicting reproductive success of insect- versus bird-pollinated scattered trees in agricultural landscapes. <i>Biological Conservation</i> , 2009, 142, 888-898.	4.1	45
52	Anthropogenic landscape change promotes asymmetric dispersal and limits regional patch occupancy in a spatially structured bird population. <i>Journal of Animal Ecology</i> , 2012, 81, 940-952.	2.8	44
53	The Application of DNA methods to Timber Tracking and Origin Verification. <i>IAWA Journal</i> , 2011, 32, 251-262.	2.7	43
54	Networked and embedded scientific experiments will improve restoration outcomes. <i>Frontiers in Ecology and the Environment</i> , 2018, 16, 288-294.	4.0	43

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55	Revegetation of urban green space rewilds soil microbiotas with implications for human health and urban design. <i>Restoration Ecology</i> , 2020, 28, S322.	2.9	43
56	Testing Putative African Tropical Forest Refugia Using Chloroplast and Nuclear DNA Phylogeography. <i>Tropical Plant Biology</i> , 2010, 3, 50-58.	1.9	40
57	Palaeodistribution modelling and genetic evidence highlight differential post-glacial range shifts of a rain forest conifer distributed across a latitudinal gradient. <i>Journal of Biogeography</i> , 2012, 39, 2292-2302.	3.0	40
58	Identifying Centres of Plant Biodiversity in South Australia. <i>PLoS ONE</i> , 2016, 11, e0144779.	2.5	40
59	Genetic Consequences of Multigenerational and Landscape Colonisation Bottlenecks for a Neotropical Forest Pioneer Tree, <i>Vochysia ferruginea</i> . <i>Tropical Plant Biology</i> , 2010, 3, 14-27.	1.9	39
60	The biodiversity impacts of non-native species should not be extrapolated from biased single-species studies. <i>Biodiversity and Conservation</i> , 2018, 27, 785-790.	2.6	36
61	Evolutionary Diversification of New Caledonian <i>Araucaria</i> . <i>PLoS ONE</i> , 2014, 9, e110308.	2.5	36
62	Routes of origin of two recently evolved hybrid taxa: <i>Senecio vulgaris</i> var. <i>hibernicus</i> and York radiate groundsel (<i>Asteraceae</i>). <i>American Journal of Botany</i> , 2000, 87, 1159-1167.	1.7	35
63	Molecular Markers and the Management of Tropical Trees: the Case of Indigenous Fruits. <i>Tropical Plant Biology</i> , 2009, 2, 1-12.	1.9	35
64	A DNA Method to Verify the Integrity of Timber Supply Chains; Confirming the Legal Sourcing of Merbau Timber From Logging Concession to Sawmill. <i>Silvae Genetica</i> , 2010, 59, 263-268.	0.8	34
65	Massively parallel sequencing and analysis of expressed sequence tags in a successful invasive plant. <i>Annals of Botany</i> , 2010, 106, 1009-1017.	2.9	33
66	Clarifying climate change adaptation responses for scattered trees in modified landscapes. <i>Journal of Applied Ecology</i> , 2011, 48, 637-641.	4.0	32
67	Chloroplast DNA Microsatellites Reveal Contrasting Phylogeographic Structure in Mahogany (<i>Swietenia macrophylla</i> King, <i>Meliaceae</i>) from Amazonia and Central America. <i>Tropical Plant Biology</i> , 2010, 3, 40-49.	1.9	31
68	A georeferenced implementation of weighted endemism. <i>Methods in Ecology and Evolution</i> , 2015, 6, 845-852.	5.2	31
69	An investigation into effects of long-distance seed dispersal on organelle population genetic structure and colonization rate: a model analysis. <i>Heredity</i> , 2004, 93, 566-576.	2.6	30
70	Opportunities for Integrated Ecological Analysis across Inland Australia with Standardised Data from Ausplots Rangelands. <i>PLoS ONE</i> , 2017, 12, e0170137.	2.5	30
71	Biogeographic concepts define invasion biology. <i>Trends in Ecology and Evolution</i> , 2009, 24, 586-586.	8.7	29
72	Local maladaptation in a foundation tree species: Implications for restoration. <i>Biological Conservation</i> , 2016, 203, 226-232.	4.1	29

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73	Bioclimatic transect networks: Powerful observatories of ecological change. <i>Ecology and Evolution</i> , 2017, 7, 4607-4619.	1.9	29
74	Weed abundance is positively correlated with native plant diversity in grasslands of southern Australia. <i>PLoS ONE</i> , 2017, 12, e0178681.	2.5	29
75	A Vegetation and Soil Survey Method for Surveillance Monitoring of Rangeland Environments. <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	2.2	28
76	Diversification history and hybridisation of <i>Dacrydium</i> (Podocarpaceae) in remote Oceania. <i>Australian Journal of Botany</i> , 2011, 59, 262.	0.6	27
77	“Sum of inverse range-sizes”™ (SIR), a biodiversity metric with many names and interpretations. <i>Biodiversity and Conservation</i> , 2015, 24, 2877-2882.	2.6	27
78	Origins of the New Allopolyploid Species <i>Senecio cambrensis</i> (Asteraceae) and its Relationship to the Canary Islands Endemic <i>Senecio teneriffae</i> . <i>American Journal of Botany</i> , 1996, 83, 1365.	1.7	26
79	Finding needles in a genomic haystack: targeted capture identifies clear signatures of selection in a nonmodel plant species. <i>Molecular Ecology</i> , 2016, 25, 4216-4233.	3.9	25
80	Higher Levels of Multiple Paternities Increase Seedling Survival in the Long-Lived Tree <i>Eucalyptus gracilis</i> . <i>PLoS ONE</i> , 2014, 9, e90478.	2.5	25
81	Herbarium Collections and Photographic Images: Alternative Data Sources for Phenological Research. , 2010, , 425-461.		24
82	Leaf trait associations with environmental variation in the wide-ranging shrub <i>Dodonaea viscosa</i> subsp. <i>angustissima</i> (Sapindaceae). <i>Austral Ecology</i> , 2017, 42, 553-561.	1.5	24
83	Spatial modelling of species turnover identifies climate ecotones, climate change tipping points and vulnerable taxonomic groups. <i>Ecography</i> , 2013, 36, 1086-1096.	4.5	23
84	Targeted capture to assess neutral genomic variation in the narrow-leaf hopbush across a continental biodiversity refugium. <i>Scientific Reports</i> , 2017, 7, 41367.	3.3	23
85	DOMESTICATION OF INDIGENOUS FRUIT AND NUT TREES FOR AGROFORESTRY IN THE SOLOMON ISLANDS. <i>Forests Trees and Livelihoods</i> , 2010, 19, 269-287.	1.2	22
86	No consistent association between changes in genetic diversity and adaptive responses of Australian acacias in novel ranges. <i>Evolutionary Ecology</i> , 2012, 26, 1345-1360.	1.2	22
87	Combining population genetics, species distribution modelling and field assessments to understand a species vulnerability to climate change. <i>Austral Ecology</i> , 2014, 39, 17-28.	1.5	22
88	AusPlots Rangelands field data collection and publication: Infrastructure for ecological monitoring. <i>Future Generation Computer Systems</i> , 2016, 56, 537-549.	7.5	21
89	Standardized genetic diversity-life history correlates for improved genetic resource management of Neotropical trees. <i>Diversity and Distributions</i> , 2018, 24, 730-741.	4.1	21
90	Consequences of long- and short-term fragmentation on the genetic diversity and differentiation of a late successional rainforest conifer. <i>Australian Journal of Botany</i> , 2011, 59, 351.	0.6	20

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91	Forest reference emission level and carbon sequestration in Cambodia. <i>Global Ecology and Conservation</i> , 2016, 7, 82-96.	2.1	20
92	Publish openly but responsibly. <i>Science</i> , 2017, 357, 141-141.	12.6	20
93	Response to Comments on "The global tree restoration potential". <i>Science</i> , 2019, 366, .	12.6	20
94	Contrasting levels of connectivity and localised persistence characterise the latitudinal distribution of a wind-dispersed rainforest canopy tree. <i>Genetica</i> , 2014, 142, 251-264.	1.1	19
95	Multi-species distribution modelling highlights the Adelaide Geosyncline, South Australia, as an important continental-scale arid-zone refugium. <i>Austral Ecology</i> , 2013, 38, 427-435.	1.5	18
96	Understanding population structure and historical demography in a conservation context: population genetics of an endangered fern. <i>Diversity and Distributions</i> , 2008, 14, 799-807.	4.1	17
97	Foundations for the future: A long-term plan for Australian ecosystem science. <i>Austral Ecology</i> , 2014, 39, 739-748.	1.5	17
98	Spatially designed revegetation—why the spatial arrangement of plants should be as important to revegetation as they are to natural systems. <i>Restoration Ecology</i> , 2018, 26, 446-455.	2.9	17
99	Plant and ant assemblages predicted to decouple under climate change. <i>Diversity and Distributions</i> , 2019, 25, 551-567.	4.1	17
100	Significant population genetic structure detected for a new and highly restricted species of <i>Atriplex</i> (Chenopodiaceae) from Western Australia, and implications for conservation management. <i>Australian Journal of Botany</i> , 2012, 60, 32.	0.6	16
101	Genetic Bottlenecks in Time and Space: Reconstructing Invasions from Contemporary and Historical Collections. <i>PLoS ONE</i> , 2014, 9, e106874.	2.5	16
102	A spatially predictive baseline for monitoring multivariate species occurrences and phylogenetic shifts in mediterranean southern Australia. <i>Journal of Vegetation Science</i> , 2014, 25, 338-348.	2.2	16
103	Population Genetics and Spatial Autocorrelation in an Unmanaged Stand of <i>Quercus petraea</i> in Denmark. <i>Scandinavian Journal of Forest Research</i> , 2003, 18, 295-304.	1.4	15
104	Can a seed bank provide demographic and genetic rescue in a declining population of the endangered shrub <i>Acacia pinguifolia</i> ?. <i>Conservation Genetics</i> , 2011, 12, 669-678.	1.5	15
105	Building a Plant DNA Barcode Reference Library for a Diverse Tropical Flora: An Example from Queensland, Australia. <i>Diversity</i> , 2016, 8, 5.	1.7	15
106	When macroecological transitions are a fiction of sampling: comparing herbarium records to plot-based species inventory data. <i>Ecography</i> , 2018, 41, 1864-1875.	4.5	15
107	The utility and limitations of chloroplast DNA analysis for identifying native British oak stands and for guiding replanting strategy. <i>Forestry</i> , 2004, 77, 335-347.	2.3	14
108	Genetic variation among <i>Helicoverpa armigera</i> populations as assessed by microsatellites: a cautionary tale about accurate allele scoring. <i>Bulletin of Entomological Research</i> , 2010, 100, 445-450.	1.0	14

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109	Restoration: 'Garden of Eden' unrealistic. <i>Nature</i> , 2016, 533, 469-469.	27.8	14
110	Mapping phylogenetic endemism in R using georeferenced branch extents. <i>SoftwareX</i> , 2015, 3-4, 22-26.	2.6	13
111	Assessment of carbon stocks of semi-evergreen forests in Cambodia. <i>Global Ecology and Conservation</i> , 2016, 5, 34-47.	2.1	13
112	Leaf morphology shift: new data and analysis support climate link. <i>Biology Letters</i> , 2013, 9, 20120860.	2.3	12
113	Applicability of chloroplast DNA barcodes for wood identification between <i>Santalum album</i> and its adulterants. <i>Holzforschung</i> , 2019, 73, 209-218.	1.9	12
114	Using the ancient past for establishing current threat in poorly inventoried regions. <i>Biological Conservation</i> , 2012, 147, 153-162.	4.1	11
115	Response to Comment on "The extent of forest in dryland biomes". <i>Science</i> , 2017, 358, .	12.6	11
116	Response to Comment on "The extent of forest in dryland biomes". <i>Science</i> , 2017, 358, 881-881.	12.6	11
117	Floristic and structural assessment of Australian rangeland vegetation with standardized plot-based surveys. <i>PLoS ONE</i> , 2018, 13, e0202073.	2.5	11
118	Systematic monitoring of heathy woodlands in a Mediterranean climate—a practical assessment of methods. <i>Environmental Monitoring and Assessment</i> , 2013, 185, 3959-3975.	2.7	10
119	Transcriptome sequencing, annotation and polymorphism detection in the hop bush, <i>Dodonaea viscosa</i> . <i>BMC Genomics</i> , 2015, 16, 803.	2.8	9
120	Hybrid swarms: catalysts for multiple evolutionary events in <i>Senecio</i> in the British Isles. <i>Plant Ecology and Diversity</i> , 2015, 8, 449-463.	2.4	9
121	Response to Comment on "The extent of forest in dryland biomes". <i>Science</i> , 2017, 358, .	12.6	9
122	Global change community ecology beyond species sorting: a quantitative framework based on mediterranean biome examples. <i>Global Ecology and Biogeography</i> , 2014, 23, 1062-1072.	5.8	8
123	Population genetics of neotropical trees focus issue. <i>Heredity</i> , 2005, 95, 243-245.	2.6	6
124	Protecting evolutionary significant units for the remnant populations of <i>Berchemiella wilsonii</i> var. <i>pubipetiolata</i> (Rhamnaceae). <i>Conservation Genetics</i> , 2007, 8, 465-473.	1.5	6
125	Genetic analysis of the dry forest timber tree <i>Sideroxylon capiri</i> in Costa Rica using AFLP. <i>Plant Systematics and Evolution</i> , 2015, 301, 15-23.	0.9	6
126	Clumped planting arrangements improve seed production in a revegetated eucalypt woodland. <i>Restoration Ecology</i> , 2019, 27, 638-646.	2.9	6

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127	Height differences in two eucalypt provenances with contrasting levels of aridity. <i>Restoration Ecology</i> , 2016, 24, 471-478.	2.9	5
128	Measuring genome-wide genetic variation to reassess subspecies classifications in <i>Dodonaea viscosa</i> (Sapindaceae). <i>Australian Journal of Botany</i> , 2018, 66, 287.	0.6	5
129	Isolation and characterization of polymorphic microsatellite loci for the invasive plant <i>Cytisus scoparius</i> . <i>Molecular Ecology Notes</i> , 2006, 7, 100-102.	1.7	4
130	Polymorphic Microsatellite Loci for <i>Virola sebifera</i> (Myristicaceae) Derived from Shotgun 454 Pyrosequencing. <i>Applications in Plant Sciences</i> , 2013, 1, 1200295.	2.1	3
131	Effective application of next-generation sequencing (NGS) approaches in systematics and population genetics: case studies in <i>Eucalyptus</i> and <i>Acacia</i> . <i>Australian Systematic Botany</i> , 2016, 29, 235.	0.9	3
132	Functional acclimation across microgeographic scales in <i>Dodonaea viscosa</i> . <i>AoB PLANTS</i> , 2018, 10, ply029.	2.3	3
133	AusPlots Rangelands Field Data Collection and Publication: Infrastructure for Ecological Monitoring. , 2014, , .		2
134	Rare genera differentiate urban green space soil bacterial communities in three cities across the world. <i>Access Microbiology</i> , 2022, 4, 000320.	0.5	2
135	Plants, position and pollination: Planting arrangement and pollination limitation in a revegetated eucalypt woodland. <i>Ecological Management and Restoration</i> , 2019, 20, 222-230.	1.5	1
136	Occasional hybridization between a native and invasive <i>Senecio</i> species in Australia is unlikely to contribute to invasive success. <i>PeerJ</i> , 2017, 5, e3630.	2.0	1
137	Interspecific hybridization and the origin of new plant taxa in Scotland. <i>Botanical Journal of Scotland</i> , 1997, 49, 247-256.	0.3	0
138	Isolation via 454 sequencing, and characterisation of microsatellites for <i>Drymodes brunneopygia</i> , southern scrub-robin (Aves: Petroicidae): a species at risk due to substantial habitat loss and climate change. <i>Conservation Genetics Resources</i> , 2012, 4, 331-333.	0.8	0