## Leo Joseph

## List of Publications by Year in descending order

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101384 74018 6,804 164 36 75 h-index citations g-index papers 171 171 171 7876 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Declining body size: a third universal response to warming?. Trends in Ecology and Evolution, 2011, 26, 285-291.	4.2	845
2	Birth of a biome: insights into the assembly and maintenance of the Australian arid zone biota. Molecular Ecology, 2008, 17, 4398-4417.	2.0	580
3	Genome 10K: A Proposal to Obtain Whole-Genome Sequence for 10 000 Vertebrate Species. Journal of Heredity, 2009, 100, 659-674.	1.0	504
4	Decline of a biome: evolution, contraction, fragmentation, extinction and invasion of the Australian mesic zone biota. Journal of Biogeography, 2011, 38, 1635-1656.	1.4	324
5	Biogeography of the Australian monsoon tropics. Journal of Biogeography, 2010, 37, 201-216.	1.4	277
6	Integrative taxonomy, or iterative taxonomy?. Systematic Entomology, 2011, 36, 209-217.	1.7	254
7	Earth history and the passerine superradiation. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 7916-7925.	3.3	238
8	A framework for incorporating evolutionary genomics into biodiversity conservation and management. Climate Change Responses, 2015, 2, .	2.6	175
9	Positive and purifying selection in mitochondrial genomes of a bird with mitonuclear discordance. Molecular Ecology, 2015, 24, 2820-2837.	2.0	112
10	Biogeography and speciation of terrestrial fauna in the southâ€western Australian biodiversity hotspot. Biological Reviews, 2015, 90, 762-793.	4.7	107
11	Impacts of recent climate change on terrestrial flora and fauna: Some emerging Australian examples. Austral Ecology, 2019, 44, 3-27.	0.7	105
12	PERCHED AT THE MITO-NUCLEAR CROSSROADS: DIVERGENT MITOCHONDRIAL LINEAGES CORRELATE WITH ENVIRONMENT IN THE FACE OF ONGOING NUCLEAR GENE FLOW IN AN AUSTRALIAN BIRD. Evolution; International Journal of Organic Evolution, 2013, 67, 3412-3428.	1.1	97
13	Shifting latitudinal clines in avian body size correlate with global warming in Australian passerines. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 3845-3852.	1.2	95
14	Phylogeography: its development and impact in Australo-Papuan ornithology with special reference to paraphyly in Australian birds. Emu, 2009, 109, 1-23.	0.2	89
15	A revised nomenclature and classification for family-group taxa of parrots (Psittaciformes). Zootaxa, 2012, 3205, 26.	0.2	83
16	Where and when does a ring start and end? Testing the ring-species hypothesis in a species complex of Australian parrots. Proceedings of the Royal Society B: Biological Sciences, 2008, 275, 2431-2440.	1.2	78
17	Multilocus analysis of honeyeaters (Aves: Meliphagidae) highlights spatio-temporal heterogeneity in the influence of biogeographic barriers in the Australian monsoonal zone. Molecular Ecology, 2010, 19, 2980-2994.	2.0	76
18	Implications of the 2019–2020 megafires for the biogeography and conservation of Australian vegetation. Nature Communications, 2021, 12, 1023.	5.8	68

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19	Phylogeny and evolution of the Meliphagoidea, the largest radiation of Australasian songbirds. Molecular Phylogenetics and Evolution, 2010, 55, 1087-1102.	1.2	65
20	Australia's arid-adapted butcherbirds experienced range expansions during Pleistocene glacial maxima. Nature Communications, 2014, 5, 3994.	5.8	65
21	Permanent Genetic Resources added to Molecular Ecology Resources database 1 January 2009–30 April 2009. Molecular Ecology Resources, 2009, 9, 1375-1379.	2.2	64
22	Mitochondrial-Dna Phylogeography of Birds in Eastern Australian Rain-Forests - First Fragments. Australian Journal of Zoology, 1994, 42, 385.	0.6	59
23	Reconciling genetic expectations from host specificity with historical population dynamics in an avian brood parasite, Horsfield's Bronze-Cuckoo Chalcites basalis of Australia. Molecular Ecology, 2002, 11, 829-837.	2.0	56
24	A Species Tree for the Australo-Papuan Fairy-wrens and Allies (Aves: Maluridae). Systematic Biology, 2012, 61, 253.	2.7	55
25	Phylogeny and biogeography in the evolution of migration: shorebirds of the Charadrius complex. Journal of Biogeography, 1999, 26, 329-342.	1.4	52
26	Independent evolution of migration on the South American landscape in a long-distance temperate-tropical migratory bird, Swainson's flycatcher (Myiarchus swainsoni). Journal of Biogeography, 2003, 30, 925-937.	1.4	52
27	A species assemblage approach to comparative phylogeography of birds in southern Australia. Ecology and Evolution, 2012, 2, 354-369.	0.8	52
28	Molecular systematics of the world's most polytypic bird: the <i>Pachycephala pectoralis</i> /i>/wesi-pachycephalidae) species complex. Zoological Journal of the Linnean Society, 2014, 170, 566-588.	1.0	48
29	Phylogeny and historical aspects of the ecology of eastern Australian scrubwrens <i>Sericornis</i> spp. â€" evidence from mitochondrial DNA. Molecular Ecology, 1993, 2, 161-170.	2.0	47
30	A new synthesis of the molecular systematics and biogeography of honeyeaters (Passeriformes:) Tj ETQq0 0 0 r	gBT /Overlo 0.7	ock 10 Tf 50 3 46
31	Zoologica Scripta, 2014, 43, 235-248.  Evolutionary history of birds across southern Australia: structure, history and taxonomic implications of mitochondrial DNA diversity in an ecologically diverse suite of species. Emu, 2015, 115, 35-48.	0.2	45
32	Resolving a phylogenetic hypothesis for parrots: implications from systematics to conservation. Emu, 2018, 118, 7-21.	0.2	45
33	Museum collections in ornithology: today's record of avian biodiversity for tomorrow's world. Emu, 2011, 111, i-xii.	0.2	44
34	Climateâ€driven mitochondrial selection: A test in Australian songbirds. Molecular Ecology, 2018, 27, 898-918.	2.0	43
35	Molecular resolution of population history, systematics and historical biogeography of the Australian ringneck parrotsBarnardius: are we there yet?. Emu, 2006, 106, 49-62.	0.2	41
36	Molecular phylogenetics suggests a New Guinean origin and frequent episodes of founder-event speciation in the nectarivorous lories and lorikeets (Aves: Psittaciformes). Molecular Phylogenetics and Evolution, 2015, 90, 34-48.	1.2	41

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37	A cryptic and critically endangered species revealed by mitochondrial DNA analyses: the Western Ground Parrot. Conservation Genetics, 2011, 12, 595-600.	0.8	40
38	Current geography masks dynamic history of gene flow during speciation in northern Australian birds. Molecular Ecology, 2019, 28, 630-643.	2.0	40
39	Inferring the phylogeography and evolutionary history of the splendid fairyâ€wren <i>Malurus splendens</i> from mitochondrial DNA and spectrophotometry. Journal of Avian Biology, 2009, 40, 7-17.	0.6	38
40	Testing the effect of transient Plio-Pleistocene barriers in monsoonal Australo-Papua: did mangrove habitats maintain genetic connectivity in the Black Butcherbird?. Molecular Ecology, 2011, 20, 5042-5059.	2.0	38
41	Lack of phylogeographic structure in three widespread Australian birds reinforces emerging challenges in Australian historical biogeography. Journal of Biogeography, 2007, 34, 612-624.	1.4	36
42	Towards a phylogenetic framework for the evolution of shakes, rattles, and rolls in Myiarchus tyrant-flycatchers (Aves: Passeriformes: Tyrannidae). Molecular Phylogenetics and Evolution, 2004, 31, 139-152.	1.2	35
43	Museum specimens provide reliable SNP data for population genomic analysis of a widely distributed but threatened cockatoo species. Molecular Ecology Resources, 2019, 19, 1578-1592.	2.2	35
44	Remarks on the phylogeny and structure of fatty acid binding proteins from parasitic platyhelminths. International Journal for Parasitology, 1997, 27, 1013-1023.	1.3	33
45	Molecular Systematics and Patterns of Diversification in Pyrrhura (Psittacidae), with Special Reference to the Picta-Leucotis Complex. Auk, 2006, 123, 660-680.	0.7	33
46	Implications of mitochondrial DNA polyphyly in two ecologically undifferentiated but morphologically distinct migratory birds, the masked and white-browed woodswallows Artamus spp. of inland Australia. Journal of Avian Biology, 2006, 37, 625-636.	0.6	33
47	Spatial variation in avian bill size is associated with humidity in summer among Australian passerines. Climate Change Responses, 2016, 3, .	2.6	33
48	Does the ring species concept predict vocal variation in the crimson rosella, Platycercus elegans, complex?. Animal Behaviour, 2009, 77, 581-593.	0.8	30
49	The impact of Pleistocene changes of climate and landscape on Australian birds: a test using the Pied Butcherbird (Cracticus nigrogularis). Emu, 2010, 110, 285-295.	0.2	30
50	Phylogeography of a â€~great speciator' (Aves: <i>Edolisoma tenuirostre</i> ) reveals complex dispersal and diversification dynamics across the Indoâ€Pacific. Journal of Biogeography, 2018, 45, 826-837.	1.4	30
51	Cryptic diversity in an endemic rainforest skink (Gnypetoscincus queenslandiae). Biodiversity and Conservation, 1993, 2, 412-425.	1.2	29
52	Molecular systematics and phylogeography of New Guinean logrunners (Orthonychidae). Emu, 2001, 101, 273-280.	0.2	29
53	Learned Vocal Variation Is Associated with Abrupt Cryptic Genetic Change in a Parrot Species Complex. PLoS ONE, 2012, 7, e50484.	1.1	29
54	<p class="HeadingRunIn"><strong>Higher classification of New World parrots (Psittaciformes; Arinae), with diagnoses of tribes</strong></p> . Zootaxa, 2013, 3691, 591.	0.2	28

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55	Perpendicular axes of differentiation generated by mitochondrial introgression. Molecular Ecology, 2017, 26, 3241-3255.	2.0	28
56	Molecular systematics of two enigmatic genera Psittacella and Pezoporus illuminate the ecological radiation of Australo-Papuan parrots (Aves: Psittaciformes). Molecular Phylogenetics and Evolution, 2011, 59, 675-684.	1.2	27
57	Uncovering cryptic evolutionary diversity in extant and extinct populations of the southern Australian arid zone Western and Thick-billed Grasswrens (Passeriformes: Maluridae: Amytornis). Conservation Genetics, 2013, 14, 1173-1184.	0.8	27
58	Recovering the evolutionary history of crowned pigeons (Columbidae: Goura): Implications for the biogeography and conservation of New Guinean lowland birds. Molecular Phylogenetics and Evolution, 2018, 120, 248-258.	1.2	27
59	A multilocus coalescent analysis of the speciational history of the Australo-Papuan butcherbirds and their allies. Molecular Phylogenetics and Evolution, 2013, 66, 941-952.	1.2	25
60	Complex histories of gene flow and a mitochondrial capture event in a nonsister pair of birds. Molecular Ecology, 2021, 30, 2087-2103.	2.0	25
61	Are natural history collections coming to an end as timeâ€series?. Frontiers in Ecology and the Environment, 2014, 12, 436-438.	1.9	24
62	Recent speciation and elevated Z hromosome differentiation between sexually monochromatic and dichromatic species of Australian teals. Journal of Avian Biology, 2016, 47, 92-102.	0.6	24
63	Genome of an iconic Australian bird: Highâ€quality assembly and linkage map of the superb fairyâ€wren ( <i>Malurus cyaneus</i> ). Molecular Ecology Resources, 2020, 20, 560-578.	2.2	24
64	MOLECULAR SYSTEMATICS AND PATTERNS OF DIVERSIFICATION IN PYRRHURA (PSITTACIDAE), WITH SPECIAL REFERENCE TO THE PICTA-LEUCOTIS COMPLEX. Auk, 2006, 123, 660.	0.7	23
65	A tangled tale of two teal: population history of the grey <i>Anas gracilis</i> and chestnut teal <i>A. castanea</i> of Australia. Journal of Avian Biology, 2009, 40, 430-439.	0.6	23
66	Multi-locus phylogeny clarifies the systematics of the Australo-Papuan robins (Family Petroicidae,) Tj ETQq0 0 0 r	gBT /Over	·lock 10 Tf 50
67	Evolution of arid zone birds in Australia: Leapfrog distribution patterns and mesic-arid connections in quail-thrush (Cinclosoma, Cinclosomatidae). Molecular Phylogenetics and Evolution, 2012, 62, 286-295.	1.2	23
68	Systematic dismantlement of <i>Lichenostomus </i> inproves the basis for understanding relationships within the honeyeaters (Meliphagidae) and the historical development of Australo-Papuan bird communities. Emu, 2011, 111, 202-211.	0.2	22
69	Neutral and selective drivers of colour evolution in a widespread Australian passerine. Journal of Biogeography, 2017, 44, 522-536.	1.4	21
70	Ultraconserved elements resolve genus-level relationships in a major Australasian bird radiation (Aves: Meliphagidae). Emu, 2019, 119, 218-232.	0.2	21
71	The Glossy Black-Cockatoo on Kangaroo Island. Emu, 1982, 82, 46-49.	0.2	20
72	Will Wallace's Line Save Australia from Avian Influenza?. Ecology and Society, 2008, 13, .	1.0	20

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73	Speciation in chestnut-shouldered fairy-wrens (Malurus spp.) and rapid phenotypic divergence in variegated fairy-wrens (Malurus lamberti): A multilocus approach. Molecular Phylogenetics and Evolution, 2012, 63, 668-678.	1.2	20
74	Australian songbird body size tracks climate variation: 82 species over 50 years. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20192258.	1.2	20
75	Ecological Divergence, Adaptive Diversification, and the Evolution of Social Signaling Traits: An Empirical Study in Arid Australian Lizards. American Naturalist, 2015, 186, E144-E161.	1.0	19
76	A genomic approach reinforces a hypothesis of mitochondrial capture in eastern Australian rosellas. Auk, 2017, 134, 181-192.	0.7	18
77	Complete subspecies-level phylogeny of the Oriolidae (Aves: Passeriformes): Out of Australasia and return. Molecular Phylogenetics and Evolution, 2019, 137, 200-209.	1.2	18
78	Evolution in Australasian Mangrove Forests: Multilocus Phylogenetic Analysis of the Gerygone Warblers (Aves: Acanthizidae). PLoS ONE, 2012, 7, e31840.	1.1	16
79	The Maluridae: inferring avian biology and evolutionary history from DNA sequences. Emu, 2013, 113, 195-207.	0.2	16
80	Phylogenetic analysis of the Australian rosella parrots (Platycercus) reveals discordance among molecules and plumage. Molecular Phylogenetics and Evolution, 2015, 91, 150-159.	1.2	16
81	Hybridisation Between the White-browed and Atherton Scrubwrens: Detection With Mitochondrial DNA. Emu, 1993, 93, 93-99.	0.2	16
82	A review of evolutionary research on birds of the New Guinean savannas and closely associated habitats of riparian rainforests, mangroves and grasslands. Emu, 2019, 119, 317-330.	0.2	15
83	Parallel Evolution of Bower-Building Behavior in Two Groups of Bowerbirds Suggested by Phylogenomics. Systematic Biology, 2020, 69, 820-829.	2.7	15
84	A taxonomic framework for interpreting evolution within the <i>Amytornis textilisâ€"modestus</i> complex of grasswrens. Emu, 2010, 110, 358-363.	0.2	14
85	Biogeographic models of gene flow in two waterfowl of the Australoâ€Papuan tropics. Ecology and Evolution, 2012, 2, 2803-2814.	0.8	14
86	Observations on breeding Night Parrots ( <i>Pezoporus occidentalis</i> ) in western Queensland. Emu, 2017, 117, 107-113.	0.2	14
87	Phylogeography and the Role of Hybridization in Speciation. Fascinating Life Sciences, 2018, , 165-194.	0.5	14
88	Evolutionary History. , 2018, , 45-75.		14
89	Comparative phylogeography of Australo-Papuan mangrove-restricted and mangrove-associated avifaunas. Biological Journal of the Linnean Society, 2013, 109, 574-598.	0.7	13
90	Multi-locus sequence data illuminate demographic drivers of Pleistocene speciation in semi-arid southern Australian birds (Cinclosoma spp.). BMC Evolutionary Biology, 2016, 16, 226.	3.2	13

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91	Norfolk Island Robins are a distinct endangered species: ancient DNA unlocks surprising relationships and phenotypic discordance within the Australo-Pacific Robins. Conservation Genetics, 2016, 17, 321-335.	0.8	13
92	Phylogeography and geno-phenotypic discordance in a widespread Australian bird, the Variegated Fairy-wren, Malurus lamberti (Aves: Maluridae). Biological Journal of the Linnean Society, 2017, 121, 655-669.	0.7	13
93	Characterizing divergence through three adjacent Australian avian transition zones. Journal of Biogeography, 2017, 44, 2247-2258.	1.4	13
94	Resources for phylogenomic analyses of Australian terrestrial vertebrates. Molecular Ecology Resources, 2017, 17, 869-876.	2.2	13
95	De-novo emergence of SINE retroposons during the early evolution of passerine birds. Mobile DNA, 2017, 8, 21.	1.3	13
96	Mitogenomic and nuclear diversity in the Mulga Parrot of the Australian arid zone: cryptic subspecies and tests for selection. Emu, 2018, 118, 22-35.	0.2	13
97	Pleistocene-dated biogeographic barriers drove divergence within the Australo-Papuan region in a sex-specific manner: an example in a widespread Australian songbird. Heredity, 2019, 123, 608-621.	1.2	13
98	Oligocene divergence of frogmouth birds (Podargidae) across Wallace's Line. Biology Letters, 2020, 16, 20200040.	1.0	13
99	The Red-Tailed Black-Cockatoo in South-Eastern Australia. Emu, 1982, 82, 42-45.	0.2	12
100	Critical Assessment of the Conservation Status of Red-tailed Black-Cockatoos in South-eastern Australia with Special Reference to Nesting Requirements. Emu, 1991, 91, 46-50.	0.2	12
101	Phylogeography and taxonomy of the Little Bronze-Cuckoo (Chalcites minutillus) in Australia's monsoon tropics. Emu, 2011, 111, 113-119.	0.2	12
102	Multilocus phylogeography of Australian teals ( <i>Anas</i> spp.): a case study of the relationship between vagility and genetic structure. Journal of Avian Biology, 2013, 44, 169-178.	0.6	12
103	Interisland gene flow among populations of the buffâ€banded rail (Aves: Rallidae) and its implications for insular endemism in Oceania. Journal of Avian Biology, 2017, 48, 679-690.	0.6	12
104	Another piece in an Australian ornithological puzzleâ€"a second Night Parrot is found dead in Queensland. Emu, 2009, 109, 198-203.	0.2	11
105	Dissection by genomic and plumage variation of a geographically complex hybrid zone between two Australian non-sister parrot species, Platycercus adscitus and Platycercus eximius. Heredity, 2019, 122, 402-416.	1.2	11
106	No Signs of Genetic Erosion in a 19th Century Genome of the Extinct Paradise Parrot (Psephotellus) Tj ETQq0 0	O rgBT /O\	verlock 10 Tf !
107	Robbery in progress: Historical museum collections bring to light a mitochondrial capture within a bird species widespread across southern Australia, the Copperback Quailâ€thrush Cinclosoma clarum. Ecology and Evolution, 2020, 10, 6785-6793.	0.8	11
108	Microscopic characteristics of the plumulaceous feathers of Australian birds: a preliminary analysis of taxonomic discrimination for forensic purposes. Australian Journal of Forensic Sciences, 2016, 48, 421-444.	0.7	10

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109	Improved systematics of lorikeets reflects their evolutionary history and frames conservation priorities. Emu, 2020, 120, 201-215.	0.2	10
110	Reassessment of a possible case of intraspecific gene flow across Australia's Great Dividing Range in the variegated fairy wren, Malurus lamberti (Aves: Maluridae), and its systematic consequences. Biological Journal of the Linnean Society, 2017, 122, 210-223.	0.7	9
111	Ancient DNA reveals that the †extinct†Hunter Island penguin (Tasidyptes hunteri) is not a distinct taxon. Zoological Journal of the Linnean Society, 2018, 182, 459-464.	1.0	9
112	Species limits in birds: Australian perspectives on interrelated challenges of allopatry, introgression of mitochondrial DNA, recent speciation, and selection. Auk, 2021, 138, .	0.7	9
113	Genetic barcoding of museum eggshell improves data integrity of avian biological collections. Scientific Reports, 2021, 11, 1605.	1.6	9
114	The origin of exotic pet sugar gliders ( <i>Petaurus breviceps</i> ) kept in the United States of America. PeerJ, 2019, 7, e6180.	0.9	9
115	Birds of Iwokrama Forest. Proceedings of the Academy of Natural Sciences of Philadelphia, 2005, 154, 109-121.	1.3	8
116	A quantum leap in avian biology. Emu, 2015, 115, 1-5.	0.2	8
117	Aberrantly plumaged orioles from the Trans-Fly savannas of New Guinea and their ecological and evolutionary significance. Emu, 2019, 119, 264-273.	0.2	8
118	Phylogeography of the iconic Australian red-tailed black-cockatoo (Calyptorhynchus banksii) and implications for its conservation. Heredity, 2020, 125, 85-100.	1,2	8
119	Descriptions of known vocalisations of the Night Parrot Pezoporus occidentalis., 0, 36, 79-88.		8
120	Advancing Genetic Methods in the Study of Parrot Biology and Conservation. Diversity, 2021, 13, 521.	0.7	8
121	A review of the conservation status of Australian parrots in 1987. Biological Conservation, 1988, 46, 261-280.	1.9	7
122	Rise (and demise?) of subspecies in the Galah (Eolophus roseicapilla), a widespread and abundant Australian cockatoo. Emu, 2015, 115, 289-301.	0.2	7
123	Parrots move to centre stage in conservation and evolution. Emu, 2018, 118, 1-6.	0.2	7
124	How far east can a Western Whistler go? Genomic data reveal large eastward range extension, taxonomic and nomenclatural change, and reassessment of conservation needs. Emu, 2021, 121, 90-101.	0.2	7
125	Benchmarking Taxonomic and Genetic Diversity After the Fact: Lessons Learned From the Catastrophic $2019\hat{a}\in 2020$ Australian Bushfires. Frontiers in Ecology and Evolution, 2021, 9, .	1.1	7
126	When DNA throws a spanner in the taxonomic works: testing for monophyly in the Dusky-capped Flycatcher, Myiarchus tuberculifer, and its South American subspecies, M. t. atriceps. Emu, 2004, 104, 197-204.	0.2	6

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127	Genetic analysis of the Australian whipbirds and wedgebills illuminates the evolution of their plumage and vocal diversity. Emu, 2013, 113, 359-366.	0.2	6
128	Taxonomic consequences of cryptic speciation in the Golden Whistler Pachycephala pectoralis complex in mainland southern Australia. Zootaxa, 2014, 3900, 294-300.	0.2	6
129	Relationships of the New Guinean subspecies, <i>Gymnorhina tibicen papuana</i> , of the Australian Magpie: an assessment from DNA sequence data. Emu, 2017, 117, 305-315.	0.2	6
130	Introgression between non-sister species of honeyeaters (Aves: Meliphagidae) several million years after speciation. Biological Journal of the Linnean Society, 2019, 128, 583-591.	0.7	6
131	Complex mosaic of sexual dichromatism and monochromatism in Pacific robins results from both gains and losses of elaborate coloration. Journal of Avian Biology, 2020, 51, .	0.6	6
132	A case for realigning species limits in the southern Australian whipbirds long recognised as the Western Whipbird (Psophodes nigrogularis). Emu, 2017, 117, 254-263.	0.2	5
133	Phylogeography of the blue-winged kookaburra <i>Dacelo leachii</i> across tropical northern Australia and New Guinea. Emu, 2020, 120, 33-45.	0.2	5
134	A taxonomic revision of the Striated Grasswren <i>Amytornis striatus</i> complex (Aves: Maluridae) after analysis of phylogenetic and phenotypic data. Emu, 2020, 120, 191-200.	0.2	5
135	Using anecdotal reports to clarify the distribution and status of a near mythical species: Australia's Night Parrot (Pezoporus occidentalis). Emu, 2021, 121, 239-249.	0.2	5
136	Diversification of <i>Petroica</i> robins across the Australo-Pacific region: first insights into the phylogenetic affinities of New Guinea's highland robin species. Emu, 2019, 119, 205-217.	0.2	4
137	Ultraconserved elements put the final nail in the coffin of traditional use of the genus Meliphaga (Aves: Meliphagidae). Zoologica Scripta, 2019, 48, 411-418.	0.7	4
138	Nuclear introns help unravel the diversification history of the Australo-Pacific Petroica robins. Molecular Phylogenetics and Evolution, 2019, 131, 48-54.	1.2	4
139	A review of specimens of Buff-breasted Button-quail <i>Turnix olivii</i> suggests serious concern for its conservation outlook. Emu, 2022, 122, 121-130.	0.2	4
140	Phylogeography of the iconic Australian pink cockatoo, <i>Lophochroa leadbeateri </i> Journal of the Linnean Society, 2021, 132, 704-723.	0.7	3
141	A multi-locus approach to discern conservation units and adaptive potential of Pacific Black Ducks across Australia and surrounding islands. Emu, 2021, 121, 124-135.	0.2	3
142	Disparate origins for endemic bird taxa from the â€~Gondwana Rainforests' of Central Eastern Australia. Biological Journal of the Linnean Society, 2021, 134, 40-56.	0.7	3
143	Two new but threatened subspecies of Rufous Grasswren Amytornis whitei (Maluridae). Bulletin of the British Ornithologists' Club, 2020, 140, 151.	0.1	3
144	Mechanisms and consequences of changing body size: reply to Bickford et al. and McCauley and Mabry. Trends in Ecology and Evolution, 2011, 26, 555-556.	4.2	2

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145	Male sex-ratio bias in the endangered South Australian Glossy Black-CockatooCalyptorhynchus lathami halmaturinus. Emu, 2015, 115, 356-359.	0.2	2
146	Ancient DNA resolves the subspecific identity of the holotype of the GalahEolophus roseicapilla, a widespread Australian cockatoo. Emu, 2016, 116, 472-475.	0.2	2
147	Ornithology of New Guinea and the Indo-Pacific Islands: introduction to the special issue of Emu – Austral Ornithology and a dedication to Paul Igag. Emu, 2019, 119, 191-195.	0.2	2
148	Taxonomic implications of recent molecular analyses of Spectacled (Symposiachrus trivirgatus) and Spot-winged (S. guttula) Monarchs (Passeriformes: Monarchidae). Emu, $0$ , , $1$ -7.	0.2	2
149	Systematics and biogeography of the whistlers (Aves: Pachycephalidae) inferred from ultraconserved elements and ancestral area reconstruction. Molecular Phylogenetics and Evolution, 2022, 168, 107379.	1.2	2
150	The Identities of Two Clutches of Calyptorhynchid Eggs from Western Victoria. Emu, 1991, 91, 51-53.	0.2	1
151	Perspectives from parrots on biological invasions. , 0, , 58-82.		1
152	Divergence and gene flow in the globally distributed blueâ€winged ducks. Journal of Avian Biology, 2017, 48, 640-649.	0.6	1
153	Circumscription, diagnosis and description of a subfamily of Australo-Papuan robins. Zootaxa, 2011, 3106, 67.	0.2	1
154	The generic taxonomy of the Australian Magpie and Australo-Papuan butcherbirds is not all black-and-white. Bulletin of the British Ornithologists' Club, 2018, 138, 346.	0.1	1
155	Genomic data show little geographical structure across the naturally fragmented range of the purple-gaped honeyeater. Australian Journal of Zoology, 2019, 67, 226.	0.6	1
156	Shane A. Parker 1943–1992. Emu, 1993, 93, 297-298.	0.2	0
157	Clarification of the nomenclatural status of a recently introduced genus-group name for some honeyeaters (Meliphagidae). Emu, 2012, 112, 173-173.	0.2	0
158	A new subspecies of the Australian Ringneck (Barnardius zonarius) from the Upper Cooper Creek Drainage, Lake Eyre Basin, central Australia. Emu, 2016, 116, 440-444.	0.2	0
159	Considerations on the type specimens and type locality of Pyrrhura roseifrons (Gray, 1859) (Psittacidae). Zootaxa, 2016, 4179, 107.	0.2	0
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