

Leo Joseph

List of Publications by Year in descending order

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Version: 2024-02-01

164
papers

6,804
citations

101384

36
h-index

74018

75
g-index

171
all docs

171
docs citations

171
times ranked

7876
citing authors

#	ARTICLE	IF	CITATIONS
1	Population connectivity across a highly fragmented distribution: Phylogeography of the Chalcophaps doves. <i>Molecular Phylogenetics and Evolution</i> , 2022, 166, 107333.	1.2	0
2	Systematics and biogeography of the whistlers (Aves: Pachycephalidae) inferred from ultraconserved elements and ancestral area reconstruction. <i>Molecular Phylogenetics and Evolution</i> , 2022, 168, 107379.	1.2	2
3	A review of specimens of Buff-breasted Button-quail (<i>Turnix olivii</i>) suggests serious concern for its conservation outlook. <i>Emu</i> , 2022, 122, 121-130.	0.2	4
4	How far east can a Western Whistler go? Genomic data reveal large eastward range extension, taxonomic and nomenclatural change, and reassessment of conservation needs. <i>Emu</i> , 2021, 121, 90-101.	0.2	7
5	Phylogeography of the iconic Australian pink cockatoo, <i>Lophochroa leadbeateri</i> . <i>Biological Journal of the Linnean Society</i> , 2021, 132, 704-723.	0.7	3
6	Implications of the 2019–2020 megafires for the biogeography and conservation of Australian vegetation. <i>Nature Communications</i> , 2021, 12, 1023.	5.8	68
7	Complex histories of gene flow and a mitochondrial capture event in a nonsister pair of birds. <i>Molecular Ecology</i> , 2021, 30, 2087-2103.	2.0	25
8	A multi-locus approach to discern conservation units and adaptive potential of Pacific Black Ducks across Australia and surrounding islands. <i>Emu</i> , 2021, 121, 124-135.	0.2	3
9	Species limits in birds: Australian perspectives on interrelated challenges of allopatry, introgression of mitochondrial DNA, recent speciation, and selection. <i>Auk</i> , 2021, 138, .	0.7	9
10	Benchmarking Taxonomic and Genetic Diversity After the Fact: Lessons Learned From the Catastrophic 2019–2020 Australian Bushfires. <i>Frontiers in Ecology and Evolution</i> , 2021, 9, .	1.1	7
11	Disparate origins for endemic bird taxa from the “Gondwana Rainforests” of Central Eastern Australia. <i>Biological Journal of the Linnean Society</i> , 2021, 134, 40-56.	0.7	3
12	Using anecdotal reports to clarify the distribution and status of a near mythical species: Australia’s Night Parrot (<i>Pezoporus occidentalis</i>). <i>Emu</i> , 2021, 121, 239-249.	0.2	5
13	Collection, curation and the use of humidification to restore nest shape in a research museum bird nest collection. <i>Avian Research</i> , 2021, 12, .	0.5	0
14	Genetic barcoding of museum eggshell improves data integrity of avian biological collections. <i>Scientific Reports</i> , 2021, 11, 1605.	1.6	9
15	Advancing Genetic Methods in the Study of Parrot Biology and Conservation. <i>Diversity</i> , 2021, 13, 521.	0.7	8
16	Emu’s first 120 years: landmark papers of change in austral ornithology. <i>Emu</i> , 2021, 121, 284-291.	0.2	0
17	Phylogeography of the blue-winged kookaburra (<i>Dacelo leachii</i>) across tropical northern Australia and New Guinea. <i>Emu</i> , 2020, 120, 33-45.	0.2	5
18	Genome of an iconic Australian bird: High-quality assembly and linkage map of the superb fairywren (<i>Malurus cyaneus</i>). <i>Molecular Ecology Resources</i> , 2020, 20, 560-578.	2.2	24

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19	A taxonomic revision of the Striated Grasswren <i>Amytornis striatus</i> complex (Aves: Maluridae) after analysis of phylogenetic and phenotypic data. <i>Emu</i> , 2020, 120, 191-200.	0.2	5
20	Improved systematics of lorikeets reflects their evolutionary history and frames conservation priorities. <i>Emu</i> , 2020, 120, 201-215.	0.2	10
21	Robbery in progress: Historical museum collections bring to light a mitochondrial capture within a bird species widespread across southern Australia, the Copperback Quail <i>Cinclosoma clarum</i> . <i>Ecology and Evolution</i> , 2020, 10, 6785-6793.	0.8	11
22	Oligocene divergence of frogmouth birds (Podargidae) across Wallace's Line. <i>Biology Letters</i> , 2020, 16, 20200040.	1.0	13
23	Parallel Evolution of Bower-Building Behavior in Two Groups of Bowerbirds Suggested by Phylogenomics. <i>Systematic Biology</i> , 2020, 69, 820-829.	2.7	15
24	Phylogeography of the iconic Australian red-tailed black-cockatoo (<i>Calyptorhynchus banksii</i>) and implications for its conservation. <i>Heredity</i> , 2020, 125, 85-100.	1.2	8
25	Complex mosaic of sexual dichromatism and monochromatism in Pacific robins results from both gains and losses of elaborate coloration. <i>Journal of Avian Biology</i> , 2020, 51, .	0.6	6
26	Biotic and abiotic drivers of evolution in some Australian thornbills (Passeriformes: Acanthiza) in allopatry, sympatry, and parapatry including a case of character displacement. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 1290-1302.	0.6	0
27	Two new but threatened subspecies of Rufous Grasswren <i>Amytornis whitei</i> (Maluridae). <i>Bulletin of the British Ornithologists' Club</i> , 2020, 140, 151.	0.1	3
28	Dissection by genomic and plumage variation of a geographically complex hybrid zone between two Australian non-sister parrot species, <i>Platycercus adscitus</i> and <i>Platycercus eximius</i> . <i>Heredity</i> , 2019, 122, 402-416.	1.2	11
29	Diversification of <i>Petroica</i> robins across the Australo-Pacific region: first insights into the phylogenetic affinities of New Guinea's highland robin species. <i>Emu</i> , 2019, 119, 205-217.	0.2	4
30	Museum specimens provide reliable SNP data for population genomic analysis of a widely distributed but threatened cockatoo species. <i>Molecular Ecology Resources</i> , 2019, 19, 1578-1592.	2.2	35
31	Ornithology of New Guinea and the Indo-Pacific Islands: introduction to the special issue of <i>Emu</i> "Austral Ornithology and a dedication to Paul Iqag". <i>Emu</i> , 2019, 119, 191-195.	0.2	2
32	Introgression between non-sister species of honeyeaters (Aves: Meliphagidae) several million years after speciation. <i>Biological Journal of the Linnean Society</i> , 2019, 128, 583-591.	0.7	6
33	A review of evolutionary research on birds of the New Guinean savannas and closely associated habitats of riparian rainforests, mangroves and grasslands. <i>Emu</i> , 2019, 119, 317-330.	0.2	15
34	Aberrantly plumaged orioles from the Trans-Fly savannas of New Guinea and their ecological and evolutionary significance. <i>Emu</i> , 2019, 119, 264-273.	0.2	8
35	No Signs of Genetic Erosion in a 19th Century Genome of the Extinct Paradise Parrot (<i>Psephotellus</i>) <i>Tj ETQq1 1 0.784314 rgBT /Overloc</i>	0.7	11
36	Ultraconserved elements resolve genus-level relationships in a major Australasian bird radiation (Aves: Meliphagidae). <i>Emu</i> , 2019, 119, 218-232.	0.2	21

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37	Complete subspecies-level phylogeny of the Oriolidae (Aves: Passeriformes): Out of Australasia and return. <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 200-209.	1.2	18
38	Pleistocene-dated biogeographic barriers drove divergence within the Australo-Papuan region in a sex-specific manner: an example in a widespread Australian songbird. <i>Heredity</i> , 2019, 123, 608-621.	1.2	13
39	Ultraconserved elements put the final nail in the coffin of traditional use of the genus <i>Meliphaga</i> (Aves: Meliphagidae). <i>Zoologica Scripta</i> , 2019, 48, 411-418.	0.7	4
40	Earth history and the passerine superradiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 7916-7925.	3.3	238
41	Australian songbird body size tracks climate variation: 82 species over 50 years. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20192258.	1.2	20
42	Current geography masks dynamic history of gene flow during speciation in northern Australian birds. <i>Molecular Ecology</i> , 2019, 28, 630-643.	2.0	40
43	Impacts of recent climate change on terrestrial flora and fauna: Some emerging Australian examples. <i>Austral Ecology</i> , 2019, 44, 3-27.	0.7	105
44	Nuclear introns help unravel the diversification history of the Australo-Pacific <i>Petroica</i> robins. <i>Molecular Phylogenetics and Evolution</i> , 2019, 131, 48-54.	1.2	4
45	Major emergency orthopaedic surgery in patients with a concomitant acute coronary event following trauma. <i>International Orthopaedics</i> , 2019, 43, 1793-1798.	0.9	0
46	The origin of exotic pet sugar gliders (<i>Petaurus breviceps</i>) kept in the United States of America. <i>PeerJ</i> , 2019, 7, e6180.	0.9	9
47	Genomic data show little geographical structure across the naturally fragmented range of the purple-gaped honeyeater. <i>Australian Journal of Zoology</i> , 2019, 67, 226.	0.6	1
48	Parrots move to centre stage in conservation and evolution. <i>Emu</i> , 2018, 118, 1-6.	0.2	7
49	Climate-driven mitochondrial selection: A test in Australian songbirds. <i>Molecular Ecology</i> , 2018, 27, 898-918.	2.0	43
50	Phylogeography of a "great speciator" (Aves: <i>Edolisoma tenuirostre</i>) reveals complex dispersal and diversification dynamics across the Indo-Pacific. <i>Journal of Biogeography</i> , 2018, 45, 826-837.	1.4	30
51	Mitogenomic and nuclear diversity in the Mulga Parrot of the Australian arid zone: cryptic subspecies and tests for selection. <i>Emu</i> , 2018, 118, 22-35.	0.2	13
52	Resolving a phylogenetic hypothesis for parrots: implications from systematics to conservation. <i>Emu</i> , 2018, 118, 7-21.	0.2	45
53	Recovering the evolutionary history of crowned pigeons (Columbidae: <i>Goura</i>): Implications for the biogeography and conservation of New Guinean lowland birds. <i>Molecular Phylogenetics and Evolution</i> , 2018, 120, 248-258.	1.2	27
54	Ancient DNA reveals that the "extinct" Hunter Island penguin (<i>Tasidyptes hunteri</i>) is not a distinct taxon. <i>Zoological Journal of the Linnean Society</i> , 2018, 182, 459-464.	1.0	9

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55	Phylogeography and the Role of Hybridization in Speciation. <i>Fascinating Life Sciences</i> , 2018, , 165-194.	0.5	14
56	Evolutionary History. , 2018, , 45-75.		14
57	The generic taxonomy of the Australian Magpie and Australo-Papuan butcherbirds is not all black-and-white. <i>Bulletin of the British Ornithologists' Club</i> , 2018, 138, 346.	0.1	1
58	A case for realigning species limits in the southern Australian whipbirds long recognised as the Western Whipbird (<i>Psophodes nigrogularis</i>). <i>Emu</i> , 2017, 117, 254-263.	0.2	5
59	Relationships of the New Guinean subspecies, <i>Gymnorhina tibicen papuana</i> , of the Australian Magpie: an assessment from DNA sequence data. <i>Emu</i> , 2017, 117, 305-315.	0.2	6
60	Phylogeography and geno-phenotypic discordance in a widespread Australian bird, the Variegated Fairy-wren, <i>Malurus lamberti</i> (Aves: Maluridae). <i>Biological Journal of the Linnean Society</i> , 2017, 121, 655-669.	0.7	13
61	Observations on breeding Night Parrots (<i>Pezoporus occidentalis</i>) in western Queensland. <i>Emu</i> , 2017, 117, 107-113.	0.2	14
62	Perpendicular axes of differentiation generated by mitochondrial introgression. <i>Molecular Ecology</i> , 2017, 26, 3241-3255.	2.0	28
63	A genomic approach reinforces a hypothesis of mitochondrial capture in eastern Australian rosellas. <i>Auk</i> , 2017, 134, 181-192.	0.7	18
64	Neutral and selective drivers of colour evolution in a widespread Australian passerine. <i>Journal of Biogeography</i> , 2017, 44, 522-536.	1.4	21
65	Reassessment of a possible case of intraspecific gene flow across Australia's Great Dividing Range in the variegated fairy wren, <i>Malurus lamberti</i> (Aves: Maluridae), and its systematic consequences. <i>Biological Journal of the Linnean Society</i> , 2017, 122, 210-223.	0.7	9
66	Characterizing divergence through three adjacent Australian avian transition zones. <i>Journal of Biogeography</i> , 2017, 44, 2247-2258.	1.4	13
67	Resources for phylogenomic analyses of Australian terrestrial vertebrates. <i>Molecular Ecology Resources</i> , 2017, 17, 869-876.	2.2	13
68	Interisland gene flow among populations of the buff-banded rail (Aves: Rallidae) and its implications for insular endemism in Oceania. <i>Journal of Avian Biology</i> , 2017, 48, 679-690.	0.6	12
69	Divergence and gene flow in the globally distributed blue-winged ducks. <i>Journal of Avian Biology</i> , 2017, 48, 640-649.	0.6	1
70	De-novo emergence of SINE retroposons during the early evolution of passerine birds. <i>Mobile DNA</i> , 2017, 8, 21.	1.3	13
71	Multi-locus sequence data illuminate demographic drivers of Pleistocene speciation in semi-arid southern Australian birds (<i>Cinclosoma</i> spp.). <i>BMC Evolutionary Biology</i> , 2016, 16, 226.	3.2	13
72	Ancient DNA resolves the subspecific identity of the holotype of the Galah <i>Eolophus roseicapilla</i> , a widespread Australian cockatoo. <i>Emu</i> , 2016, 116, 472-475.	0.2	2

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73	Recent speciation and elevated Z chromosome differentiation between sexually monochromatic and dichromatic species of Australian teals. <i>Journal of Avian Biology</i> , 2016, 47, 92-102.	0.6	24
74	A new subspecies of the Australian Ringneck (<i>Barnardius zonarius</i>) from the Upper Cooper Creek Drainage, Lake Eyre Basin, central Australia. <i>Emu</i> , 2016, 116, 440-444.	0.2	0
75	Considerations on the type specimens and type locality of <i>Pyrrhura roseifrons</i> (Gray, 1859) (Psittacidae). <i>Zootaxa</i> , 2016, 4179, 107.	0.2	0
76	Spatial variation in avian bill size is associated with humidity in summer among Australian passerines. <i>Climate Change Responses</i> , 2016, 3, .	2.6	33
77	Norfolk Island Robins are a distinct endangered species: ancient DNA unlocks surprising relationships and phenotypic discordance within the Australo-Pacific Robins. <i>Conservation Genetics</i> , 2016, 17, 321-335.	0.8	13
78	Microscopic characteristics of the plumulaceous feathers of Australian birds: a preliminary analysis of taxonomic discrimination for forensic purposes. <i>Australian Journal of Forensic Sciences</i> , 2016, 48, 421-444.	0.7	10
79	Phylogenetic analysis of the Australian rosella parrots (<i>Platycercus</i>) reveals discordance among molecules and plumage. <i>Molecular Phylogenetics and Evolution</i> , 2015, 91, 150-159.	1.2	16
80	Evolutionary history of birds across southern Australia: structure, history and taxonomic implications of mitochondrial DNA diversity in an ecologically diverse suite of species. <i>Emu</i> , 2015, 115, 35-48.	0.2	45
81	A framework for incorporating evolutionary genomics into biodiversity conservation and management. <i>Climate Change Responses</i> , 2015, 2, .	2.6	175
82	Positive and purifying selection in mitochondrial genomes of a bird with mitonuclear discordance. <i>Molecular Ecology</i> , 2015, 24, 2820-2837.	2.0	112
83	A quantum leap in avian biology. <i>Emu</i> , 2015, 115, 1-5.	0.2	8
84	Molecular phylogenetics suggests a New Guinean origin and frequent episodes of founder-event speciation in the nectarivorous lorries and lorikeets (Aves: Psittaciformes). <i>Molecular Phylogenetics and Evolution</i> , 2015, 90, 34-48.	1.2	41
85	Male sex-ratio bias in the endangered South Australian Glossy Black-Cockatoo <i>Calyptorhynchus lathami halmaturinus</i> . <i>Emu</i> , 2015, 115, 356-359.	0.2	2
86	Rise (and demise?) of subspecies in the Galah (<i>Eolophus roseicapilla</i>), a widespread and abundant Australian cockatoo. <i>Emu</i> , 2015, 115, 289-301.	0.2	7
87	Ecological Divergence, Adaptive Diversification, and the Evolution of Social Signaling Traits: An Empirical Study in Arid Australian Lizards. <i>American Naturalist</i> , 2015, 186, E144-E161.	1.0	19
88	Biogeography and speciation of terrestrial fauna in the south-western Australian biodiversity hotspot. <i>Biological Reviews</i> , 2015, 90, 762-793.	4.7	107
89	Taxonomic consequences of cryptic speciation in the Golden Whistler <i>Pachycephala pectoralis</i> complex in mainland southern Australia. <i>Zootaxa</i> , 2014, 3900, 294-300.	0.2	6
90	A new synthesis of the molecular systematics and biogeography of honeyeaters (Passeriformes: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 6 <i>Zoologica Scripta</i> , 2014, 43, 235-248.	0.7	46

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91	Molecular systematics of the world's most polytypic bird: the <i>Pachycephala pectoralis</i> / <i>melanura</i> (Aves: Pachycephalidae) species complex. <i>Zoological Journal of the Linnean Society</i> , 2014, 170, 566-588.	1.0	48
92	Are natural history collections coming to an end as timeâ€series?. <i>Frontiers in Ecology and the Environment</i> , 2014, 12, 436-438.	1.9	24
93	Australiaâ€™s arid-adapted butcherbirds experienced range expansions during Pleistocene glacial maxima. <i>Nature Communications</i> , 2014, 5, 3994.	5.8	65
94	The Maluridae: inferring avian biology and evolutionary history from DNA sequences. <i>Emu</i> , 2013, 113, 195-207.	0.2	16
95	Genetic analysis of the Australian whipbirds and wedgebills illuminates the evolution of their plumage and vocal diversity. <i>Emu</i> , 2013, 113, 359-366.	0.2	6
96	A multilocus coalescent analysis of the speciation history of the Australo-Papuan butcherbirds and their allies. <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 941-952.	1.2	25
97	Multilocus phylogeography of Australian teals (<i>Anas</i> spp.): a case study of the relationship between vagility and genetic structure. <i>Journal of Avian Biology</i> , 2013, 44, 169-178.	0.6	12
98	PERCHED AT THE MITO-NUCLEAR CROSSROADS: DIVERGENT MITOCHONDRIAL LINEAGES CORRELATE WITH ENVIRONMENT IN THE FACE OF ONGOING NUCLEAR GENE FLOW IN AN AUSTRALIAN BIRD. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 3412-3428.	1.1	97
99	Comparative phylogeography of Australo-Papuan mangrove-restricted and mangrove-associated avifaunas. <i>Biological Journal of the Linnean Society</i> , 2013, 109, 574-598.	0.7	13
100	Uncovering cryptic evolutionary diversity in extant and extinct populations of the southern Australian arid zone Western and Thick-billed Grasswrens (Passeriformes: Maluridae: Amytornis). <i>Conservation Genetics</i> , 2013, 14, 1173-1184.	0.8	27
101	<p class="HeadingRunIn">Higher classification of New World parrots (Psittaciformes; Arinae), with diagnoses of tribes</p>. <i>Zootaxa</i> , 2013, 3691, 591.	0.2	28
102	Clarification of the nomenclatural status of a recently introduced genus-group name for some honeyeaters (Meliphagidae). <i>Emu</i> , 2012, 112, 173-173.	0.2	0
103	Biogeographic models of gene flow in two waterfowl of the Australoâ€™Papuan tropics. <i>Ecology and Evolution</i> , 2012, 2, 2803-2814.	0.8	14
104	Evolution of arid zone birds in Australia: Leapfrog distribution patterns and mesic-arid connections in quail-thrush (Cinclosoma, Cinclosomatidae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 286-295.	1.2	23
105	Speciation in chestnut-shouldered fairy-wrens (<i>Malurus</i> spp.) and rapid phenotypic divergence in variegated fairy-wrens (<i>Malurus lamberti</i>): A multilocus approach. <i>Molecular Phylogenetics and Evolution</i> , 2012, 63, 668-678.	1.2	20
106	Evolution in Australasian Mangrove Forests: Multilocus Phylogenetic Analysis of the Gerygone Warblers (Aves: Acanthizidae). <i>PLoS ONE</i> , 2012, 7, e31840.	1.1	16
107	Learned Vocal Variation Is Associated with Abrupt Cryptic Genetic Change in a Parrot Species Complex. <i>PLoS ONE</i> , 2012, 7, e50484.	1.1	29
108	A revised nomenclature and classification for family-group taxa of parrots (Psittaciformes). <i>Zootaxa</i> , 2012, 3205, 26.	0.2	83

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109	A species assemblage approach to comparative phylogeography of birds in southern Australia. <i>Ecology and Evolution</i> , 2012, 2, 354-369.	0.8	52
110	A Species Tree for the Australo-Papuan Fairy-wrens and Allies (Aves: Maluridae). <i>Systematic Biology</i> , 2012, 61, 253.	2.7	55
111	Declining body size: a third universal response to warming?. <i>Trends in Ecology and Evolution</i> , 2011, 26, 285-291.	4.2	845
112	Mechanisms and consequences of changing body size: reply to Bickford et al. and McCauley and Mabry. <i>Trends in Ecology and Evolution</i> , 2011, 26, 555-556.	4.2	2
113	Systematic dismantlement of <i>Lichenostomus</i> improves the basis for understanding relationships within the honeyeaters (Meliphagidae) and the historical development of Australo-Papuan bird communities. <i>Emu</i> , 2011, 111, 202-211.	0.2	22
114	Phylogeography and taxonomy of the Little Bronze-Cuckoo (<i>Chalcites minutillus</i>) in Australia's monsoon tropics. <i>Emu</i> , 2011, 111, 113-119.	0.2	12
115	Decline of a biome: evolution, contraction, fragmentation, extinction and invasion of the Australian mesic zone biota. <i>Journal of Biogeography</i> , 2011, 38, 1635-1656.	1.4	324
116	Testing the effect of transient Plio-Pleistocene barriers in monsoonal Australo-Papua: did mangrove habitats maintain genetic connectivity in the Black Butcherbird?. <i>Molecular Ecology</i> , 2011, 20, 5042-5059.	2.0	38
117	Integrative taxonomy, or iterative taxonomy?. <i>Systematic Entomology</i> , 2011, 36, 209-217.	1.7	254
118	A cryptic and critically endangered species revealed by mitochondrial DNA analyses: the Western Ground Parrot. <i>Conservation Genetics</i> , 2011, 12, 595-600.	0.8	40
119	Molecular systematics of two enigmatic genera <i>Psittacella</i> and <i>Pezoporus</i> illuminate the ecological radiation of Australo-Papuan parrots (Aves: Psittaciformes). <i>Molecular Phylogenetics and Evolution</i> , 2011, 59, 675-684.	1.2	27
120	Museum collections in ornithology: today's record of avian biodiversity for tomorrow's world. <i>Emu</i> , 2011, 111, i-xii.	0.2	44
121	Circumscription, diagnosis and description of a subfamily of Australo-Papuan robins. <i>Zootaxa</i> , 2011, 3106, 67.	0.2	1
122	Phylogeny and evolution of the Meliphagoidea, the largest radiation of Australasian songbirds. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 1087-1102.	1.2	65
123	Multilocus analysis of honeyeaters (Aves: Meliphagidae) highlights spatio-temporal heterogeneity in the influence of biogeographic barriers in the Australian monsoonal zone. <i>Molecular Ecology</i> , 2010, 19, 2980-2994.	2.0	76
124	Biogeography of the Australian monsoon tropics. <i>Journal of Biogeography</i> , 2010, 37, 201-216.	1.4	277
125	A taxonomic framework for interpreting evolution within the <i>Amytornis textilis</i> "modestus" complex of grasswrens. <i>Emu</i> , 2010, 110, 358-363.	0.2	14
126	The impact of Pleistocene changes of climate and landscape on Australian birds: a test using the Pied Butcherbird (<i>Cracticus nigrogularis</i>). <i>Emu</i> , 2010, 110, 285-295.	0.2	30

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127	Genome 10K: A Proposal to Obtain Whole-Genome Sequence for 10,000 Vertebrate Species. <i>Journal of Heredity</i> , 2009, 100, 659-674.	1.0	504
128	Shifting latitudinal clines in avian body size correlate with global warming in Australian passerines. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2009, 276, 3845-3852.	1.2	95
129	Does the ring species concept predict vocal variation in the crimson rosella, <i>Platycercus elegans</i> , complex?. <i>Animal Behaviour</i> , 2009, 77, 581-593.	0.8	30
130	Inferring the phylogeography and evolutionary history of the splendid fairywren <i>Malurus splendens</i> from mitochondrial DNA and spectrophotometry. <i>Journal of Avian Biology</i> , 2009, 40, 7-17.	0.6	38
131	A tangled tale of two teal: population history of the grey <i>Anas gracilis</i> and chestnut teal <i>A. castanea</i> of Australia. <i>Journal of Avian Biology</i> , 2009, 40, 430-439.	0.6	23
132	Multi-locus phylogeny clarifies the systematics of the Australo-Papuan robins (Family Petroicidae). <i>Trends in Ecology & Evolution</i> , 2009, 24, 10-17.	1.2	23
133	Another piece in an Australian ornithological puzzle—a second Night Parrot is found dead in Queensland. <i>Emu</i> , 2009, 109, 198-203.	0.2	11
134	Phylogeography: its development and impact in Australo-Papuan ornithology with special reference to paraphyly in Australian birds. <i>Emu</i> , 2009, 109, 1-23.	0.2	89
135	Permanent Genetic Resources added to Molecular Ecology Resources database 1 January 2009–30 April 2009. <i>Molecular Ecology Resources</i> , 2009, 9, 1375-1379.	2.2	64
136	Birth of a biome: insights into the assembly and maintenance of the Australian arid zone biota. <i>Molecular Ecology</i> , 2008, 17, 4398-4417.	2.0	580
137	Where and when does a ring start and end? Testing the ring-species hypothesis in a species complex of Australian parrots. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2008, 275, 2431-2440.	1.2	78
138	Will Wallace's Line Save Australia from Avian Influenza?. <i>Ecology and Society</i> , 2008, 13, .	1.0	20
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