

Per G P Ericson

List of Publications by Year in descending order

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86
papers

4,759
citations

101384

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102304

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89
all docs

89
docs citations

89
times ranked

3659
citing authors

#	ARTICLE	IF	CITATIONS
1	Quantifying adaptive divergence of the snowfinches in a common landscape. <i>Diversity and Distributions</i> , 2022, 28, 2579-2592.	1.9	3
2	Utilizing museomics to trace the complex history and species boundaries in an avian-study system of conservation concern. <i>Heredity</i> , 2022, 128, 159-168.	1.2	9
3	A guide to avian museomics: Insights gained from resequencing hundreds of avian study skins. <i>Molecular Ecology Resources</i> , 2022, 22, 2672-2684.	2.2	19
4	The evolution of ancestral and species-specific adaptations in snowfinches at the Qinghai-Tibet Plateau. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	22
5	Genomic signatures of rapid adaptive divergence in a tropical montane species. <i>Biology Letters</i> , 2021, 17, 20210089.	1.0	3
6	Rapid phenotypic evolution with shallow genomic differentiation during early stages of high elevation adaptation in Eurasian Tree Sparrows. <i>National Science Review</i> , 2020, 7, 113-127.	4.6	36
7	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
8	Great journey of Great Tits (<i>Parus major</i> group): Origin, diversification and historical demographics of a broadly distributed bird lineage. <i>Journal of Biogeography</i> , 2020, 47, 1585-1598.	1.4	15
9	Genomic differentiation tracks earth-historic isolation in an Indo-Australasian archipelagic pitta (Pittidae; Aves) complex. <i>BMC Evolutionary Biology</i> , 2019, 19, 151.	3.2	14
10	No Signs of Genetic Erosion in a 19th Century Genome of the Extinct Paradise Parrot (<i>Psephotellus</i>)	0.7	11
11	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
12	Near-complete phylogeny and taxonomic revision of the world's babblers (Aves: Passeriformes). <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 346-356.	1.2	72
13	Rapid expansion and diversification into new niche space by fluvicoline flycatchers. <i>Journal of Avian Biology</i> , 2018, 49, jav-01661.	0.6	10
14	Complete taxon sampling of the avian genus <i>Pica</i> (magpies) reveals ancient relictual populations and synchronous Late-Pleistocene demographic expansion across the Northern Hemisphere. <i>Journal of Avian Biology</i> , 2018, 49, jav-01612.	0.6	20
15	A genomic perspective of the pink-headed duck <i>Rhodonessa caryophyllacea</i> suggests a long history of low effective population size. <i>Scientific Reports</i> , 2017, 7, 16853.	1.6	8
16	Phylogeny, biogeography and taxonomic consequences in a bird-of-paradise species complex, <i>Lophorina</i> - <i>Ptiloris</i> (Aves: Paradisaeidae). <i>Zoological Journal of the Linnean Society</i> , 2017, , .	1.0	0
17	Identifying Bird Remains Using Ancient DNA Barcoding. <i>Genes</i> , 2017, 8, 169.	1.0	12
18	Systematics and biodiversity research in the era of genomics. <i>Zoologica Scripta</i> , 2016, 45, 3-4.	0.7	0

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19	Sillem's Mountain Finch (<i>Leucosticte sillemi</i>) is a valid species of rosefinch (<i>Carpodacus</i>)	1.0	20
20	Shaped by uneven Pleistocene climate: mitochondrial phylogeographic pattern and population history of white wagtail (<i>Motacilla alba</i>) (Aves: Passeriformes). <i>Journal of Avian Biology</i> , 2016, 47, 263-274.	0.6	21
21	Contrasting phylogeographic signatures in two Australo-Papuan bowerbird species complexes (Aves)	0.7	21
22	Genetic responses to seasonal variation in altitudinal stress: whole-genome resequencing of great tit in eastern Himalayas. <i>Scientific Reports</i> , 2015, 5, 14256.	1.6	33
23	Dating the diversification of the major lineages of Passeriformes (Aves). <i>BMC Evolutionary Biology</i> , 2014, 14, 8.	3.2	57
24	A complete multilocus species phylogeny of the tits and chickadees (Aves: Paridae). <i>Molecular Phylogenetics and Evolution</i> , 2013, 69, 852-860.	1.2	53
25	A multi-locus phylogeny suggests an ancient hybridization event between Campephilus and melanerpine woodpeckers (Aves: Picidae). <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 578-588.	1.2	32
26	Ground tit genome reveals avian adaptation to living at high altitudes in the Tibetan plateau. <i>Nature Communications</i> , 2013, 4, 2071.	5.8	229
27	Molecular phylogeny of the manakins (Aves: Passeriformes: Pipridae), with a new classification and the description of a new genus. <i>Molecular Phylogenetics and Evolution</i> , 2013, 69, 796-804.	1.2	38
28	The spatio-temporal colonization and diversification across the Indo-Pacific by a "great speciator" (Aves, <i>Erythropitta erythrogaster</i>). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20130309.	1.2	52
29	Phylogeny and classification of the New World suboscines (Aves, Passeriformes). <i>Zootaxa</i> , 2013, 3613, 1-35.	0.2	81
30	Convergent evolution of morphological and ecological traits in the open-habitat chat complex (Aves)	0.0	10
31	Comparative phylogeography of two widespread magpies: Importance of habitat preference and breeding behavior on genetic structure in China. <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 562-572.	1.2	40
32	The phylogenetic relationships and generic limits of finches (Fringillidae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 581-596.	1.2	75
33	Molecular and morphological evidences place the extinct New Zealand endemic <i>Turnagra capensis</i> in the Oriolidae. <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 414-426.	1.2	13
34	Nuclear DNA from a 180-year-old study skin reveals the phylogenetic position of the Kinglet <i>Calyptura cristata</i> (Passeriformes: Tyrannidae). <i>Ibis</i> , 2012, 154, 533-541.	1.0	4
35	Evolution of terrestrial birds in three continents: biogeography and parallel radiations. <i>Journal of Biogeography</i> , 2012, 39, 813-824.	1.4	70
36	Phylogenetic utility and evolution of indels: A study in neognathous birds. <i>Molecular Phylogenetics and Evolution</i> , 2011, 61, 760-771.	1.2	20

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37	Neumann's Warbler (<i>Hemitesia neumanni</i>) (Sylvioidea): the sole African member of a Palaeotropical Miocene avifauna. <i>Ibis</i> , 2011, 153, 78-86.	1.0	12
38	Circumscription of a monophyletic family for the tapaculos (Aves: Rhinocryptidae): <i>Psiloramphus</i> in and <i>Melanopareia</i> out. <i>Journal of Ornithology</i> , 2010, 151, 337-345.	0.5	13
39	DNA barcoding of Scandinavian birds reveals divergent lineages in trans-Atlantic species. <i>Journal of Ornithology</i> , 2010, 151, 565-578.	0.5	129
40	The <i>Monticola</i> rock-thrushes: Phylogeny and biogeography revisited. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 901-910.	1.2	12
41	The phylogenetic position of the Black-collared Bulbul (<i>Neolestes torquatus</i>). <i>Ibis</i> , 2010, 152, 386-392.	1.0	13
42	The Cinnamon Ibon (<i>Hypocryptadius cinnamomeus</i>) is a forest canopy sparrow. <i>Ibis</i> , 2010, 152, 747-760.	1.0	19
43	A molecular phylogeny of minivets (Passeriformes: Campephagidae: <i>Pericrocotus</i>): implications for biogeography and convergent plumage evolution. <i>Zoologica Scripta</i> , 2010, 39, 1-8.	0.7	17
44	A multi-gene phylogeny disentangles the chat-flycatcher complex (Aves: Muscicapidae). <i>Zoologica Scripta</i> , 2010, 39, 213-224.	0.7	49
45	A new genus for three species of tyrant flycatchers (Passeriformes: Tyrannidae), formerly placed in <i>Myiophobus</i> . <i>Zootaxa</i> , 2009, 2290, 36-40.	0.2	5
46	An unexpectedly long history of sexual selection in birds-of-paradise. <i>BMC Evolutionary Biology</i> , 2009, 9, 235.	3.2	71
47	Convergent evolution, habitat shifts and variable diversification rates in the ovenbird-woodcreeper family (Furnariidae). <i>BMC Evolutionary Biology</i> , 2009, 9, 268.	3.2	34
48	Phylogeny of babblers (Aves, Passeriformes): major lineages, family limits and classification. <i>Zoologica Scripta</i> , 2009, 38, 225-236.	0.7	67
49	A multi-gene phylogeny reveals novel relationships for aberrant genera of Australo-Papuan core Corvoidea and polyphyly of the Pachycephalidae and Psophodidae (Aves: Passeriformes). <i>Molecular Phylogenetics and Evolution</i> , 2009, 52, 488-497.	1.2	39
50	Explosive avian radiations and multi-directional dispersal across Wallacea: Evidence from the Campephagidae and other Crown Corvidae (Aves). <i>Molecular Phylogenetics and Evolution</i> , 2008, 47, 221-236.	1.2	71
51	Molecular support for a rapid cladogenesis of the woodpecker clade Malarpicini, with further insights into the genus <i>Picus</i> (Piciformes: Picinae). <i>Molecular Phylogenetics and Evolution</i> , 2008, 48, 34-46.	1.2	25
52	The systematic affinity of the enigmatic <i>Lamprolia victoriae</i> (Aves: Passeriformes) – An example of avian dispersal between New Guinea and Fiji over Miocene intermittent land bridges?. <i>Molecular Phylogenetics and Evolution</i> , 2008, 48, 1218-1222.	1.2	39
53	Tyrant flycatchers coming out in the open: phylogeny and ecological radiation of Tyrannidae (Aves). <i>Trends in Ecology and Evolution</i> , 2008, 23, 87-94.	0.7	87
54	Phylogenetic relationships among Palearctic-Oriental starlings and mynas (genera <i>Sturnus</i>). <i>Trends in Ecology and Evolution</i> , 2008, 23, 10-17.	0.7	24

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55	Current perspectives on the evolution of birds. <i>Contributions To Zoology</i> , 2008, 77, 109-116.	0.2	7
56	Flight Speeds among Bird Species: Allometric and Phylogenetic Effects. <i>PLoS Biology</i> , 2007, 5, e197.	2.6	220
57	Hangin' on to our rocks 'n clocks: a reply to Brown et al.. <i>Biology Letters</i> , 2007, 3, 260-261.	1.0	9
58	Systematic placement of an enigmatic Southeast Asian taxon <i>Eupetes macrocerus</i> and implications for the biogeography of a main songbird radiation, the Passerida. <i>Biology Letters</i> , 2007, 3, 323-326.	1.0	32
59	A molecular phylogeny of the cotingas (Aves: Cotingidae). <i>Molecular Phylogenetics and Evolution</i> , 2007, 42, 25-37.	1.2	28
60	Phylogeny of the ovenbird genus <i>Upucerthia</i> : a case of independent adaptations for terrestrial life. <i>Zoologica Scripta</i> , 2007, 36, 133-141.	0.7	17
61	Evolution of the ovenbird-woodcreeper assemblage (Aves: Furnariidae) - major shifts in nest architecture and adaptive radiation. <i>Journal of Avian Biology</i> , 2006, 37, 260-272.	0.6	55
62	Diversification of Neoaves: integration of molecular sequence data and fossils. <i>Biology Letters</i> , 2006, 2, 543-547.	1.0	617
63	Nuclear DNA from old collections of avian study skins reveals the evolutionary history of the Old World suboscines (Aves, Passeriformes). <i>Zoologica Scripta</i> , 2006, 35, 567-580.	0.7	129
64	Phylogeny and classification of the avian superfamily Sylvioidea. <i>Molecular Phylogenetics and Evolution</i> , 2006, 38, 381-397.	1.2	143
65	Systematic affinities of two enigmatic New Zealand passerines of high conservation priority, the hihi or stitchbird <i>Notiomystis cincta</i> and the kokako <i>Callaeas cinerea</i> . <i>Molecular Phylogenetics and Evolution</i> , 2006, 40, 281-284.	1.2	25
66	Molecular phylogenetic relationship of snow finch complex (genera <i>Montifringilla</i> , <i>Pyrgilauda</i> , and) <i>Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50</i>	1.2	30
67	Higher-level phylogeny and morphological evolution of tyrant flycatchers, cotingas, manakins, and their allies (Aves: Tyrannida). <i>Molecular Phylogenetics and Evolution</i> , 2006, 40, 471-483.	1.2	69
68	Nuclear and mitochondrial sequence data reveal the major lineages of starlings, mynas and related taxa. <i>Molecular Phylogenetics and Evolution</i> , 2006, 41, 333-344.	1.2	36
69	Out of Gondwanaland; the evolutionary history of cooperative breeding and social behaviour among crows, magpies, jays and allies. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006, 273, 1117-1125.	1.2	53
70	Inter-generic relationships of the crows, jays, magpies and allied groups (Aves: Corvidae) based on nucleotide sequence data. <i>Journal of Avian Biology</i> , 2005, 36, 222-234.	0.6	79
71	Molecular data reveal some major adaptational shifts in the early evolution of the most diverse avian family, the Furnariidae. <i>Journal Fur Ornithologie</i> , 2005, 146, 1-13.	1.2	44
72	Phylogenetic relationships of woodcreepers (Aves: Dendrocolaptinae) - incongruence between molecular and morphological data. <i>Journal of Avian Biology</i> , 2004, 35, 280-288.	0.6	39

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73	Phylogenetic relationships of typical antbirds (Thamnophilidae) and test of incongruence based on Bayes factors. <i>BMC Evolutionary Biology</i> , 2004, 4, 23.	3.2	57
74	Inter-familial relationships of the shorebirds (Aves: Charadriiformes) based on nuclear DNA sequence data. <i>BMC Evolutionary Biology</i> , 2003, 3, 16.	3.2	70
75	Phylogeny of Passerida (Aves: Passeriformes) based on nuclear and mitochondrial sequence data. <i>Molecular Phylogenetics and Evolution</i> , 2003, 29, 126-138.	1.2	135
76	Molecular support for a sister group relationship between Pici and Galbulae (Piciformes sensu) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622	0.6	40
77	Evolution, biogeography, and patterns of diversification in passerine birds. <i>Journal of Avian Biology</i> , 2003, 34, 3-15.	0.6	134
78	<i>Pseudopodoces humilis</i> , a misclassified terrestrial tit (Paridae) of the Tibetan Plateau: evolutionary consequences of shifting adaptive zones. <i>Ibis</i> , 2003, 145, 185-202.	1.0	65
79	<i>Sapayoa aenigma</i> : a New World representative of 'Old World suboscines'. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2003, 270, S238-41.	1.2	84
80	BASAL PHYLOGENY OF THE TYRANNOIDEA BASED ON COMPARISONS OF CYTOCHROME b AND EXONS OF NUCLEAR c-myc AND RAG-1 GENES. <i>Auk</i> , 2002, 119, 984.	0.7	33
81	A MIDDLE EOCENE SHOREBIRD FROM CHINA. <i>Condor</i> , 2002, 104, 896.	0.7	10
82	A Gondwanan origin of passerine birds supported by DNA sequences of the endemic New Zealand wrens. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2002, 269, 235-241.	1.2	305
83	A Middle Eocene Shorebird from China. <i>Condor</i> , 2002, 104, 896-899.	0.7	15
84	Systematic relationships and biogeography of the tracheophone suboscines (Aves: Passeriformes). <i>Molecular Phylogenetics and Evolution</i> , 2002, 23, 499-512.	1.2	125
85	Systematic affinities of the lyrebirds (Passeriformes: Menura), with a novel classification of the major groups of passerine birds. <i>Molecular Phylogenetics and Evolution</i> , 2002, 25, 53-62.	1.2	78
86	Phylogeny of major lineages of suboscines (Passeriformes) analysed by nuclear DNA sequence data. <i>Journal of Avian Biology</i> , 2001, 32, 15-25.	0.6	84