

Allan J Baker

List of Publications by Year in descending order

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Version: 2024-02-01

64
papers

3,930
citations

109321

35
h-index

123424

61
g-index

65
all docs

65
docs citations

65
times ranked

3962
citing authors

#	ARTICLE	IF	CITATIONS
1	Rapid population decline in red knots: fitness consequences of decreased refuelling rates and late arrival in Delaware Bay. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2004, 271, 875-882.	2.6	373
2	A Mitogenomic Timescale for Birds Detects Variable Phylogenetic Rates of Molecular Evolution and Refutes the Standard Molecular Clock. <i>Molecular Biology and Evolution</i> , 2006, 23, 1731-1740.	8.9	222
3	Convergent regulatory evolution and loss of flight in paleognathous birds. <i>Science</i> , 2019, 364, 74-78.	12.6	189
4	Phylogenetic relationships and divergence times of Charadriiformes genera: multigene evidence for the Cretaceous origin of at least 14 clades of shorebirds. <i>Biology Letters</i> , 2007, 3, 205-210.	2.3	173
5	Single mitochondrial gene barcodes reliably identify sister-species in diverse clades of birds. <i>BMC Evolutionary Biology</i> , 2008, 8, 81.	3.2	170
6	RAG-1 sequences resolve phylogenetic relationships within Charadriiform birds. <i>Molecular Phylogenetics and Evolution</i> , 2003, 29, 268-278.	2.7	145
7	Complete mitochondrial DNA genome sequences show that modern birds are not descended from transitional shorebirds. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2002, 269, 839-846.	2.6	119
8	Multiple gene evidence for expansion of extant penguins out of Antarctica due to global cooling. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006, 273, 11-17.	2.6	118
9	HISTORICAL DEMOGRAPHY AND PRESENT DAY POPULATION STRUCTURE OF THE GREENFINCH, <i>CARDUEUS CHLORIS</i> -AN ANALYSIS OF mtDNA CONTROL-REGION SEQUENCES. <i>Evolution; International Journal of Organic Evolution</i> , 1997, 51, 946-956.	2.3	111
10	Mitochondrial and Nuclear DNA Sequences Support a Cretaceous Origin of Columbiformes and a Dispersal-Driven Radiation in the Paleogene. <i>Systematic Biology</i> , 2007, 56, 656-672.	5.6	110
11	Phylogenetic Relationships and Historical Biogeography of Neotropical Parrots (Psittaciformes: Tj ETQq1 1 0.784314 rgBT /Overlock 10 55, 454-470.	5.6	108
12	Natural selection shaped the rise and fall of passenger pigeon genomic diversity. <i>Science</i> , 2017, 358, 951-954.	12.6	105
13	A molecular timescale for galliform birds accounting for uncertainty in time estimates and heterogeneity of rates of DNA substitutions across lineages and sites. <i>Molecular Phylogenetics and Evolution</i> , 2006, 38, 499-509.	2.7	103
14	Rates of mass gain and energy deposition in red knot on their final spring staging site is both time- and condition-dependent. <i>Journal of Applied Ecology</i> , 2007, 44, 885-895.	4.0	89
15	Whole-Genome Analyses Resolve the Phylogeny of Flightless Birds (Palaeognathae) in the Presence of an Empirical Anomaly Zone. <i>Systematic Biology</i> , 2019, 68, 937-955.	5.6	88
16	Multiple nuclear genes and retroposons support vicariance and dispersal of the palaeognaths, and an Early Cretaceous origin of modern birds. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 4617-4625.	2.6	86
17	A phylogenetic framework for the terns (Sternini) inferred from mtDNA sequences: implications for taxonomy and plumage evolution. <i>Molecular Phylogenetics and Evolution</i> , 2005, 35, 459-469.	2.7	82
18	Reconstructing the tempo and mode of evolution in an extinct clade of birds with ancient DNA: The giant moas of New Zealand. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 8257-8262.	7.1	82

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19	Genomic Support for a Moa–Tinamou Clade and Adaptive Morphological Convergence in Flightless Ratites. <i>Molecular Biology and Evolution</i> , 2014, 31, 1686-1696.	8.9	80
20	Explosive ice age diversification of kiwi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E5580-7.	7.1	78
21	Combined Nuclear and Mitochondrial DNA Sequences Resolve Generic Relationships within the Cracidae (Galliformes, Aves). <i>Systematic Biology</i> , 2002, 51, 946-958.	5.6	75
22	Countering criticisms of single mitochondrial DNA gene barcoding in birds. <i>Molecular Ecology Resources</i> , 2009, 9, 257-268.	4.8	75
23	Association between mitochondrial DNA and morphological evolution in Canada geese. <i>Journal of Molecular Evolution</i> , 1990, 31, 373-382.	1.8	69
24	Sequences from 14 mitochondrial genes provide a well-supported phylogeny of the Charadriiform birds congruent with the nuclear RAG-1 tree. <i>Molecular Phylogenetics and Evolution</i> , 2006, 39, 657-667.	2.7	69
25	Title is missing!. <i>Conservation Genetics</i> , 2003, 4, 167-177.	1.5	67
26	A POPULATION MEMETICS APPROACH TO CULTURAL EVOLUTION IN CHAFFINCH SONG: DIFFERENTIATION AMONG POPULATIONS. <i>Evolution; International Journal of Organic Evolution</i> , 1994, 48, 351-359.	2.3	65
27	Unravelling the migration and moult strategies of a long-distance migrant using stable isotopes: Red Knot <i>Calidris canutus</i> movements in the Americas. <i>Ibis</i> , 2005, 147, 738-749.	1.9	63
28	DNA Barcode Detects High Genetic Structure within Neotropical Bird Species. <i>PLoS ONE</i> , 2011, 6, e28543.	2.5	63
29	VICARIANT SPECIATION OF CURASSOWS (AVES, CRACIDAE): A HYPOTHESIS BASED ON MITOCHONDRIAL DNA PHYLOGENY. <i>Auk</i> , 2004, 121, 682.	1.4	58
30	Feather Development Genes and Associated Regulatory Innovation Predate the Origin of Dinosauria. <i>Molecular Biology and Evolution</i> , 2015, 32, 23-28.	8.9	57
31	Vicariant Speciation of Curassows (Aves, Cracidae): A Hypothesis Based on Mitochondrial DNA Phylogeny. <i>Auk</i> , 2004, 121, 682-694.	1.4	55
32	DNA evidence for a Paleocene origin of the Alcidae (Aves: Charadriiformes) in the Pacific and multiple dispersals across northern oceans. <i>Molecular Phylogenetics and Evolution</i> , 2008, 46, 430-445.	2.7	47
33	POPULATION DIVERGENCE TIMES AND HISTORICAL DEMOGRAPHY IN RED KNOTS AND DUNLINS. <i>Condor</i> , 2005, 107, 497.	1.6	46
34	Conserved Nonexonic Elements: A Novel Class of Marker for Phylogenomics. <i>Systematic Biology</i> , 2017, 66, 1028-1044.	5.6	46
35	Population Divergence Times and Historical Demography in red Knots and Dunlins. <i>Condor</i> , 2005, 107, 497-513.	1.6	44
36	Gastro-intestinal microbiota of two migratory shorebird species during spring migration staging in Delaware Bay, USA. <i>Journal of Ornithology</i> , 2014, 155, 969-977.	1.1	42

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37	Mechanisms of song differentiation in introduced populations of Chaffinches (<i>Fringilla coelebs</i>) in New Zealand. <i>Ibis</i> , 1984, 126, 510-524.	1.9	36
38	A novel mitochondrial gene order in shorebirds (Scolopacidae, Charadriiformes). <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 411-416.	2.7	32
39	MULTIPLE GENE EVIDENCE FOR PARALLEL EVOLUTION AND RETENTION OF ANCESTRAL MORPHOLOGICAL STATES IN THE SHANKS (CHARADRIIFORMES: SCOLOPACIDAE). <i>Condor</i> , 2005, 107, 514.	1.6	29
40	Contrasting Phylogeographic Patterns in Mitochondrial DNA and Microsatellites: Evidence of Female Philopatry and Male-biased Gene Flow among Regional Populations of the Blue-and-yellow Macaw (<i>Psittaciformes:Ara ararauna</i>) in Brazil. <i>Auk</i> , 2009, 126, 359-370.	1.4	28
41	Linking intronic polymorphism on the <i>CHD1</i> gene with fitness correlates in Black-tailed Godwits (<i>Limosa l. limosa</i>). <i>Ibis</i> , 2010, 152, 368-377.	1.9	23
42	Multiple Gene Evidence for Parallel Evolution and Retention of Ancestral Morphological States in the Shanks (Charadriiformes: Scolopacidae). <i>Condor</i> , 2005, 107, 514-526.	1.6	20
43	Multigene phylogeny and DNA barcoding indicate that the Sandwich tern complex (<i>Thalasseus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 52, 263-267.	2.7	20
44	Characterization of MHC class I in a long-distance migrant shorebird suggests multiple transcribed genes and intergenic recombination. <i>Immunogenetics</i> , 2013, 65, 211-225.	2.4	19
45	Phylogenetic and coalescent analysis of three loci suggest that the Water Rail is divisible into two species, <i>Rallus aquaticus</i> and <i>R. indicus</i> . <i>BMC Evolutionary Biology</i> , 2010, 10, 226.	3.2	18
46	Criteria for aging and sexing New Zealand oystercatchers. <i>New Zealand Journal of Marine and Freshwater Research</i> , 1974, 8, 211-221.	2.0	16
47	Eight independent nuclear genes support monophyly of the plovers: The role of mutational variance in gene trees. <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 631-641.	2.7	15
48	Molecular Evidence for Recent Radiation in Southern Hemisphere Masked Gulls. <i>Auk</i> , 2005, 122, 268-279.	1.4	13
49	MORPHOMETRIC VARIABILITY IN CONTINENTAL AND ATLANTIC ISLAND POPULATIONS OF CHAFFINCHES (<i>FRINGILLA COELEBS</i>). <i>Evolution; International Journal of Organic Evolution</i> , 1991, 45, 29-39.	2.3	12
50	MOLECULAR EVIDENCE FOR RECENT RADIATION IN SOUTHERN HEMISPHERE MASKED GULLS. <i>Auk</i> , 2005, 122, 268.	1.4	12
51	High genetic diversity in the blue-listed British Columbia population of the purple martin maintained by multiple sources of immigrants. <i>Conservation Genetics</i> , 2008, 9, 495-505.	1.5	10
52	A rare case of <i>Plasmodium</i> (<i>Haemamoeba</i>) <i>relictum</i> infection in a free-living Red Knot (<i>Calidris canutus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 1.1	1.1	10
53	Islands in the sky: the impact of Pleistocene climate cycles on biodiversity. <i>Journal of Biology</i> , 2008, 7, 32.	2.7	7
54	One hundred new universal exonic markers for birds developed from a genomic pipeline. <i>Journal of Ornithology</i> , 2014, 155, 561-569.	1.1	7

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55	Global flyway evolution in red knots <i>Calidris canutus</i> and genetic evidence for a Nearctic refugium. <i>Molecular Ecology</i> , 2022, 31, 2124-2139.	3.9	7
56	Species limits and population differentiation in New Zealand snipes (Scolopacidae: Coenocorypha). <i>Conservation Genetics</i> , 2010, 11, 1363-1374.	1.5	6
57	The enigmatic monotypic crab plover <i>Dromas ardeola</i> is closely related to pratincoles and coursers (Aves, Charadriiformes, Glareolidae). <i>Genetics and Molecular Biology</i> , 2010, 33, 583-586.	1.3	6
58	Lipid levels in the South Island pied oystercatcher (<i>Haematopus ostralegus finschi</i>). <i>New Zealand Journal of Zoology</i> , 1975, 2, 425-434.	1.1	3
59	Mitochondrial-DNA evidence shows the Australian Painted Snipe is a full species, <i>Rostratula australis</i> . <i>Emu</i> , 2007, 107, 185-189.	0.6	3
60	Novel and cross-species microsatellite markers for parentage analysis in Sanderling <i>Calidris alba</i> . <i>Journal of Ornithology</i> , 2011, 152, 807-810.	1.1	2
61	Relationships of gulls – A reply to Bourne. <i>Auk</i> , 2006, 123, 906-907.	1.4	1
62	Speciation in Birds. <i>Condor</i> , 2008, 110, 396-398.	1.6	1
63	Molecular Advances in the Study of Geographic Variation and Speciation in Birds. <i>Ornithological Monographs</i> , 2007, , 18-29.	1.3	0
64	Molecular evidence for introgressive hybridization in New Zealand masked gulls. <i>Ibis</i> , 2023, 165, 248-269.	1.9	0