

# Martin Irestedt

## List of Publications by Year in descending order

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Version: 2024-02-01

73  
papers

2,943  
citations

186265

28  
h-index

189892

50  
g-index

80  
all docs

80  
docs citations

80  
times ranked

2372  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | 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.                       | 2.6 | 305       |
| 2  | Ecological and evolutionary determinants for the adaptive radiation of the Madagascan vangas. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 6620-6625.  | 7.1 | 151       |
| 3  | Evolution, biogeography, and patterns of diversification in passerine birds. <i>Journal of Avian Biology</i> , 2003, 34, 3-15.  | 1.2 | 134       |
| 4  | 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.                       | 1.7 | 129       |
| 5  | Systematic relationships and biogeography of the tracheophone suboscines (Aves: Passeriformes). <i>Molecular Phylogenetics and Evolution</i> , 2002, 23, 499-512.   | 2.7 | 125       |
| 6  | Identifying the causes and consequences of assembly gaps using a multiplatform genome assembly of a bird-of-paradise. <i>Molecular Ecology Resources</i> , 2021, 21, 263-286.                                 | 4.8 | 103       |
| 7  | Phylogeny of major lineages of suboscines (Passeriformes) analysed by nuclear DNA sequence data. <i>Journal of Avian Biology</i> , 2001, 32, 15-25.   | 1.2 | 84        |
| 8  | Sapayoa aenigma : a New World representative of 'Old World suboscines'. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2003, 270, S238-41.  | 2.6 | 84        |
| 9  | Phylogeny and classification of the New World suboscines (Aves, Passeriformes). <i>Zootaxa</i> , 2013, 3613, 1-35.  | 0.5 | 81        |
| 10 | 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.           | 2.7 | 78        |
| 11 | Supermatrix phylogeny and biogeography of the Australasian Meliphagidae radiation (Aves: Tj ETQq1 1 0.784314 rBT /Overlock 10 TFS   | 2.7 | 77        |
| 12 | Near-complete phylogeny and taxonomic revision of the world's babblers (Aves: Passeriformes). <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 346-356.  | 2.7 | 72        |
| 13 | Explosive avian radiations and multi-directional dispersal across Wallacea: Evidence from the Campephagidae and other Crown Corvida (Aves). <i>Molecular Phylogenetics and Evolution</i> , 2008, 47, 221-236. | 2.7 | 71        |
| 14 | An unexpectedly long history of sexual selection in birds-of-paradise. <i>BMC Evolutionary Biology</i> , 2009, 9, 235.  | 3.2 | 71        |
| 15 | 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        |
| 16 | Dynamic evolutionary history and gene content of sex chromosomes across diverse songbirds. <i>Nature Ecology and Evolution</i> , 2019, 3, 834-844.  | 7.8 | 68        |
| 17 | Evidence of taxon cycles in an Indo-Pacific passerine bird radiation (Aves: Pachycephala ). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20131727.                             | 2.6 | 67        |
| 18 | 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        |

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|----|--|-----|-----------|
| 19 | Dating the diversification of the major lineages of Passeriformes (Aves). <i>BMC Evolutionary Biology</i> , 2014, 14, 8.   | 3.2 | 57        |
| 20 | 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.  | 1.2 | 55        |
| 21 | 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.                              | 2.6 | 52        |
| 22 | The avian W chromosome is a refugium for endogenous retroviruses with likely effects on female-biased mutational load and genetic incompatibilities. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2021, 376, 20200186.          | 4.0 | 46        |
| 23 | 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        |
| 24 | Phylogenetic relationships of woodcreepers (Aves: Dendrocolaptinae) - incongruence between molecular and morphological data. <i>Journal of Avian Biology</i> , 2004, 35, 280-288.  | 1.2 | 39        |
| 25 | 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. | 2.7 | 39        |
| 26 | The division of the major songbird radiation into Passerida and "core Corvoidea" (Aves: Passeriformes) - the species tree vs. gene trees. <i>Zoologica Scripta</i> , 2008, 37, 305-313.  | 1.7 | 37        |
| 27 | 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        |
| 28 | Phylogeny and historical biogeography of gnateaters (Passeriformes, Conopophagidae) in the South America forests. <i>Molecular Phylogenetics and Evolution</i> , 2014, 79, 422-432.  | 2.7 | 33        |
| 29 | 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.   | 2.3 | 32        |
| 30 | 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.  | 3.0 | 30        |
| 31 | Dynamic colonization exchanges between continents and islands drive diversification in paradise-flycatchers ( <i>Terpsiphone</i> , Monarchidae). <i>Journal of Biogeography</i> , 2012, 39, 1900-1918.   | 3.0 | 29        |
| 32 | Dramatic niche shifts and morphological change in two insular bird species. <i>Royal Society Open Science</i> , 2015, 2, 140364.   | 2.4 | 29        |
| 33 | Reconciling supertramps, great speciators and relict species with the taxon cycle stages of a large island radiation (Aves: Campephagidae). <i>Journal of Biogeography</i> , 2019, 46, 1214-1225.  | 3.0 | 26        |
| 34 | Mitochondrial and nuclear DNA phylogenies reveal a complex evolutionary history in the Australasian robins (Passeriformes: Petroicidae). <i>Molecular Phylogenetics and Evolution</i> , 2011, 61, 726-738.   | 2.7 | 25        |
| 35 | Molecular systematics and evolution of the <i>Synallaxis ruficapilla</i> complex (Aves: Furnariidae) in the Atlantic Forest. <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 86-94.   | 2.7 | 24        |
| 36 | Comparative analyses identify genomic features potentially involved in the evolution of birds-of-paradise. <i>GigaScience</i> , 2019, 8, .   | 6.4 | 22        |

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|----|--|------|-----------|
| 37 | Contrasting phylogeographic signatures in two Australo-Papuan bowerbird species complexes (Aves: Tj ETQq1 1,0,784314,rgBT /Ome   | 1.7  | 21        |
| 38 | 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. | 1.2  | 20        |
| 39 | The Cinnamon Ibon <i>Hypocryptadius cinnamomeus</i> is a forest canopy sparrow. <i>Ibis</i> , 2010, 152, 747-760.  | 1.9  | 19        |
| 40 | A guide to avian museomics: Insights gained from resequencing hundreds of avian study skins. <i>Molecular Ecology Resources</i> , 2022, 22, 2672-2684.   | 4.8  | 19        |
| 41 | Complete subspecies-level phylogeny of the Oriolidae (Aves: Passeriformes): Out of Australasia and return. <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 200-209.  | 2.7  | 18        |
| 42 | Satellite DNA evolution in Corvoidea inferred from short and long reads. <i>Molecular Ecology</i> , 2023, 32, 1288-1305.   | 3.9  | 18        |
| 43 | Phylogeny of the ovenbird genus <i>Upucerthia</i> : a case of independent adaptations for terrestrial life. <i>Zoologica Scripta</i> , 2007, 36, 133-141.  | 1.7  | 17        |
| 44 | 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.  | 1.7  | 17        |
| 45 | Systematics and biogeography of Indo-Pacific ground-doves. <i>Molecular Phylogenetics and Evolution</i> , 2011, 59, 538-543.   | 2.7  | 17        |
| 46 | Phylogenomics of white-eyes, a "great speciator", reveals Indonesian archipelago as the center of lineage diversity. <i>ELife</i> , 2020, 9, .   | 6.0  | 17        |
| 47 | Molecular phylogenetics and species limits in a cryptically coloured radiation of Australo-Papuan passerine birds (Pachycephalidae: Colluricincla). <i>Molecular Phylogenetics and Evolution</i> , 2018, 124, 100-105.                               | 2.7  | 16        |
| 48 | Parallel Evolution of Bower-Building Behavior in Two Groups of Bowerbirds Suggested by Phylogenomics. <i>Systematic Biology</i> , 2020, 69, 820-829.   | 5.6  | 15        |
| 49 | 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.  | 3.0  | 15        |
| 50 | Systematic revision of the avian family Cisticolidae based on a multi-locus phylogeny of all genera. <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 790-799.   | 2.7  | 14        |
| 51 | Densely sampled phylogenetic analyses of the Lesser Short-toed Lark ( <i>Alaudala rufescens</i> ) " Sand Lark ( <i>A. Araytal</i> ) species complex (Aves, Passeriformes) reveal cryptic diversity. <i>Zoologica Scripta</i> , 2020, 49, 427-439.    | 1.7  | 14        |
| 52 | Complex population structure of the Atlantic puffin revealed by whole genome analyses. <i>Communications Biology</i> , 2021, 4, 922.   | 4.4  | 14        |
| 53 | The formation of avian montane diversity across barriers and along elevational gradients. <i>Nature Communications</i> , 2022, 13, 268.  | 12.8 | 14        |
| 54 | 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.   | 1.1  | 13        |

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|----|--|-----|-----------|
| 55 | Fine-scale barriers to connectivity across a fragmented South-East Asian landscape in six songbird species. <i>Evolutionary Applications</i> , 2020, 13, 1026-1036.  | 3.1 | 13        |
| 56 | Neumann's Warbler ( <i>Hemitesia neumanni</i> ) (Sylvioidea): the sole African member of a Palaeotropical Miocene avifauna. <i>Ibis</i> , 2011, 153, 78-86.  | 1.9 | 12        |
| 57 | Identifying Bird Remains Using Ancient DNA Barcoding. <i>Genes</i> , 2017, 8, 169.   | 2.4 | 12        |
| 58 | Novel genome and genome-wide SNPs reveal early fragmentation effects in an edge-tolerant songbird population across an urbanized tropical metropolis. <i>Scientific Reports</i> , 2018, 8, 12804.                  | 3.3 | 12        |
| 59 | No Signs of Genetic Erosion in a 19th Century Genome of the Extinct Paradise Parrot ( <i>Psephotellus</i> )  | 1.7 | 11        |
| 60 | Rapid expansion and diversification into new niche space by fluvicoline flycatchers. <i>Journal of Avian Biology</i> , 2018, 49, jav-01661.  | 1.2 | 10        |
| 61 | Unrecognised (species) diversity in New Guinean passerine birds. <i>Emu</i> , 2019, 119, 233-241.  | 0.6 | 10        |
| 62 | 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.  | 2.6 | 9         |
| 63 | Molecular phylogeny of the Indian Ocean Terpsiphone paradise flycatchers: Undetected evolutionary diversity revealed amongst island populations. <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 336-347. | 2.7 | 8         |
| 64 | 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.                               | 3.3 | 8         |
| 65 | Relicts of the lost arc: High-throughput sequencing of the <i>Eutrichomyias rowleyi</i> (Aves)   | 2.7 | 8         |
| 66 | Diversification and community assembly of the world's largest tropical island. <i>Global Ecology and Biogeography</i> , 2022, 31, 1078-1089.   | 5.8 | 8         |
| 67 | Multiple species within the Striated Prinia ( <i>Prinia crinigera</i> ) Brown Prinia ( <i>P. polychroa</i> ) complex revealed through an integrative taxonomic approach. <i>Ibis</i> , 2020, 162, 936-967.         | 1.9 | 7         |
| 68 | The New Zealand Thrush: An Extinct Oriole. <i>PLoS ONE</i> , 2011, 6, e24317.  | 2.5 | 7         |
| 69 | Sequence Transpositions Restore Genes on the Highly Degenerated W Chromosomes of Songbirds. <i>Genes</i> , 2020, 11, 1267.   | 2.4 | 5         |
| 70 | Nuclear DNA from a 180-year-old study skin reveals the phylogenetic position of the Kinglet <i>Calyptura cristata</i> (Passeriformes: Tyrannides). <i>Ibis</i> , 2012, 154, 533-541.                               | 1.9 | 4         |
| 71 | 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, 984-995.   | 1.4 | 2         |
| 72 | Speciation and population divergence in a mutualistic seed dispersing bird. <i>Communications Biology</i> , 2022, 5, 429.  | 4.4 | 1         |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 73 | Phylogeny, biogeography and taxonomic consequences in a bird-of-paradise species complex, Lophorinaâ€Ptiloris (Aves: Paradisaeidae). Zoological Journal of the Linnean Society, 2017, , . | 2.3 | 0         |