# Rebecca T Kimball

## List of Publications by Citations



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| 125 papers | 5,299 citations | 35 h-index | 71 g-index |
| 143 ext. papers | 6,287 ext. citations | 3.6 avg, IF | 5.63 L-index |

| # | Paper | IF | Citations |
|---|---|---|---|
| 125 | A phylogenomic study of birds reveals their evolutionary history. *Science*, **2008**, 320, 1763-8 | 33.3 | 1433 |
| 124 | Phylogenomic evidence for multiple losses of flight in ratite birds. *Proceedings of the National Academy of Sciences of the United States of America*, **2008**, 105, 13462-7 | 11.5 | 157 |
| 123 | A well-tested set of primers to amplify regions spread across the avian genome. *Molecular Phylogenetics and Evolution*, **2009**, 50, 654-60 | 4.1 | 156 |
| 122 | Why Do Phylogenomic Data Sets Yield Conflicting Trees? Data Type Influences the Avian Tree of Life more than Taxon Sampling. *Systematic Biology*, **2017**, 66, 857-879 | 8.4 | 151 |
| 121 | Mitochondrial genomes and avian phylogeny: complex characters and resolvability without explosive radiations. *Molecular Biology and Evolution*, **2007**, 24, 269-80 | 8.3 | 151 |
| 120 | Avoiding Missing Data Biases in Phylogenomic Inference: An Empirical Study in the Landfowl (Aves: Galliformes). *Molecular Biology and Evolution*, **2016**, 33, 1110-25 | 8.3 | 145 |
| 119 | A molecular phylogeny of the pheasants and partridges suggests that these lineages are not monophyletic. *Molecular Phylogenetics and Evolution*, **1999**, 11, 38-54 | 4.1 | 135 |
| 118 | Dispersers shape fruit diversity in Ficus (Moraceae). *Proceedings of the National Academy of Sciences of the United States of America*, **2010**, 107, 14668-72 | 11.5 | 130 |
| 117 | Evolution of Avian Plumage Dichromatism from a Proximate Perspective. *American Naturalist*, **1999** , 154, 182-193 | 3.7 | 128 |
| 116 | Earth history and the passerine superradiation. *Proceedings of the National Academy of Sciences of the United States of America*, **2019**, 116, 7916-7925 | 11.5 | 121 |
| 115 | Biogeography of Discontinuously Distributed Hydrophytes: A Molecular Appraisal of Intercontinental Disjunctions. *International Journal of Plant Sciences*, **2003**, 164, 917-932 | 2.6 | 106 |
| 114 | Co-phylogeography and comparative population genetics of the threatened Galapagos hawk and three ectoparasite species: ecology shapes population histories within parasite communities. *Molecular Ecology*, **2007**, 16, 4759-73 | 5.7 | 102 |
| 113 | Analysis of a Rapid Evolutionary Radiation Using Ultraconserved Elements: Evidence for a Bias in Some Multispecies Coalescent Methods. *Systematic Biology*, **2016**, 65, 612-27 | 8.4 | 100 |
| 112 | Building the avian tree of life using a large-scale, sparse supermatrix. *Molecular Phylogenetics and Evolution*, **2015**, 84, 53-63 | 4.1 | 90 |
| 111 | Parsimony and model-based analyses of indels in avian nuclear genes reveal congruent and incongruent phylogenetic signals. *Biology*, **2013**, 2, 419-44 | 4.9 | 76 |
| 110 | Examining Basal avian divergences with mitochondrial sequences: model complexity, taxon sampling, and sequence length. *Systematic Biology*, **2002**, 51, 614-25 | 8.4 | 69 |
| 109 | Identifying localized biases in large datasets: a case study using the avian tree of life. *Molecular Phylogenetics and Evolution*, **2013**, 69, 1021-32 | 4.1 | 67 |

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| 108 | Introns outperform exons in analyses of basal avian phylogeny using clathrin heavy chain genes. *Gene*, **2008**, 410, 89-96 | 3.8 | 64 |
| 107 | Sexual ornamentation, condition, and immune defence in the house sparrow Passer domesticus. *Behavioral Ecology and Sociobiology*, **1996**, 39, 317-322 | 2.5 | 63 |
| 106 | Incongruence among different mitochondrial regions: a case study using complete mitogenomes. *Molecular Phylogenetics and Evolution*, **2014**, 78, 314-23 | 4.1 | 60 |
| 105 | Ratite nonmonophyly: independent evidence from 40 novel Loci. *Systematic Biology*, **2013**, 62, 35-49 | 8.4 | 59 |
| 104 | Assessing phylogenetic relationships among galliformes: a multigene phylogeny with expanded taxon sampling in Phasianidae. *PLoS ONE*, **2013**, 8, e64312 | 3.7 | 57 |
| 103 | Testing hypotheses about the sister group of the passeriformes using an independent 30-locus data set. *Molecular Biology and Evolution*, **2012**, 29, 737-50 | 8.3 | 56 |
| 102 | Phylogeography of the Galþagos hawk (Buteo galapagoensis): a recent arrival to the Galþagos Islands. *Molecular Phylogenetics and Evolution*, **2006**, 39, 237-47 | 4.1 | 56 |
| 101 | Phylogenetic Utility of Avian Ovomucoid Intron G: A Comparison of Nuclear and Mitochondrial Phylogenies in Galliformes. *Auk*, **2001**, 118, 799-804 | 2.1 | 56 |
| 100 | Correlated evolution of fig size and color supports the dispersal syndromes hypothesis. *Oecologia*, **2008**, 156, 783-96 | 2.9 | 55 |
| 99 | Are transposable element insertions homoplasy free?: an examination using the avian tree of life. *Systematic Biology*, **2011**, 60, 375-86 | 8.4 | 51 |
| 98 | Mate choice by female red junglefowl: the issues of multiple ornaments and fluctuating asymmetry. *Animal Behaviour*, **1998**, 55, 41-50 | 2.8 | 50 |
| 97 | A Phylogenomic Supertree of Birds. *Diversity*, **2019**, 11, 109 | 2.5 | 47 |
| 96 | Evolution of the mitochondrial DNA control region and cytochrome b genes and the inference of phylogenetic relationships in the avian genus Lophura (Galliformes). *Molecular Phylogenetics and Evolution*, **2001**, 19, 187-201 | 4.1 | 46 |
| 95 | Geographic variation in the acoustic traits of greater horseshoe bats: testing the importance of drift and ecological selection in evolutionary processes. *PLoS ONE*, **2013**, 8, e70368 | 3.7 | 44 |
| 94 | Phylogeny of Coreopsideae (Asteraceae) using ITS sequences suggests lability in reproductive characters. *Molecular Phylogenetics and Evolution*, **2004**, 33, 127-39 | 4.1 | 39 |
| 93 | The evolution of peafowl and other taxa with ocelli (eyespots): a phylogenomic approach. *Proceedings of the Royal Society B: Biological Sciences*, **2014**, 281, | 4.4 | 37 |
| 92 | Duplication of accelerated evolution and growth hormone gene in passerine birds. *Molecular Biology and Evolution*, **2008**, 25, 352-61 | 8.3 | 37 |
| 91 | Patterns of vertebrate isochore evolution revealed by comparison of expressed mammalian, avian, and crocodilian genes. *Journal of Molecular Evolution*, **2007**, 65, 259-66 | 3.1 | 35 |

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| 90 | A macroevolutionary perspective on multiple sexual traits in the phasianidae (galliformes). *International Journal of Evolutionary Biology*, **2011**, 2011, 423938 | | 33 |
| 89 | Phylogenetic Position of the New World Quail (Odontophoridae): Eight Nuclear Loci and Three Mitochondrial Regions Contradict Morphology and the Sibley-Ahlquist Tapestry. *Auk*, **2007**, 124, 71-84 | 2.1 | 31 |
| 88 | A molecular phylogeny of the peacock-pheasants (Galliformes: Polyplectron spp.) indicates loss and reduction of ornamental traits and display behaviours. *Biological Journal of the Linnean Society*, **2001**, 73, 187-198 | 1.9 | 31 |
| 87 | Does more sequence data improve estimates of galliform phylogeny? Analyses of a rapid radiation using a complete data matrix. *PeerJ*, **2014**, 2, e361 | 3.1 | 30 |
| 86 | Ancestral range reconstruction of Galliformes: the effects of topology and taxon sampling. *Journal of Biogeography*, **2017**, 44, 122-135 | 4.1 | 29 |
| 85 | Comparative molecular evolution and phylogenetic utility of 3'-UTRs and introns in Galliformes. *Molecular Phylogenetics and Evolution*, **2010**, 56, 536-42 | 4.1 | 29 |
| 84 | Molecular and morphological reassessment of relationships within the Vittadinia group of Astereae (Asteraceae). *American Journal of Botany*, **2001**, 88, 1279-1289 | 2.7 | 29 |
| 83 | Rapid morphological change of a top predator with the invasion of a novel prey. *Nature Ecology and Evolution*, **2018**, 2, 108-115 | 12.3 | 29 |
| 82 | Land connectivity changes and global cooling shaped the colonization history and diversification of New World quail (Aves: Galliformes: Odontophoridae). *Journal of Biogeography*, **2015**, 42, 1883-1895 | 4.1 | 28 |
| 81 | Polytomies, the power of phylogenetic inference, and the stochastic nature of molecular evolution: a comment on Walsh et al. (1999). *Evolution; International Journal of Organic Evolution*, **2001**, 55, 1261-3; discussion 1264-6 | 3.8 | 28 |
| 80 | Insight from an ultraconserved element bait set designed for hemipteran phylogenetics integrated with genomic resources. *Molecular Phylogenetics and Evolution*, **2019**, 130, 297-303 | 4.1 | 28 |
| 79 | Homoplastic microinversions and the avian tree of life. *BMC Evolutionary Biology*, **2011**, 11, 141 | 3 | 27 |
| 78 | Testicular Asymmetry and Secondary Sexual Characters in Red Junglefowl. *Auk*, **1997**, 114, 221-228 | 2.1 | 27 |
| 77 | Resolution of the phylogenetic position of the Congo peafowl, Afropavo congensis: a biogeographic and evolutionary enigma. *Proceedings of the Royal Society B: Biological Sciences*, **1997**, 264, 1517-23 | 4.4 | 27 |
| 76 | Sorting out relationships among the grouse and ptarmigan using intron, mitochondrial, and ultra-conserved element sequences. *Molecular Phylogenetics and Evolution*, **2016**, 98, 123-32 | 4.1 | 27 |
| 75 | Tempo and Pattern of Avian Brain Size Evolution. *Current Biology*, **2020**, 30, 2026-2036.e3 | 6.3 | 26 |
| 74 | How do seemingly non-vagile clades accomplish trans-marine dispersal? Trait and dispersal evolution in the landfowl (Aves: Galliformes). *Proceedings of the Royal Society B: Biological Sciences*, **2017**, 284, | 4.4 | 25 |
| 73 | Phylogenetic Position of the New World Quail (Odontophoridae): Eight Nuclear Loci and Three Mitochondrial Regions Contradict Morphology and the Sibley-Ahlquist Tapestry. *Auk*, **2007**, 124, 71 | 2.1 | 24 |

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| 72 | Rapid and recent diversification of curassows, guans, and chachalacas (Galliformes: Cracidae) out of Mesoamerica: Phylogeny inferred from mitochondrial, intron, and ultraconserved element sequences. *Molecular Phylogenetics and Evolution*, **2016**, 102, 320-30 | 4.1 | 21 |
| 71 | Fluctuating asymmetry in red junglefowl. *Journal of Evolutionary Biology*, **1997**, 10, 441 | 2.3 | 21 |
| 70 | What are the roles of taxon sampling and model fit in tests of cyto-nuclear discordance using avian mitogenomic data?. *Molecular Phylogenetics and Evolution*, **2019**, 130, 132-142 | 4.1 | 21 |
| 69 | Female Choice for Male Morphological Traits in House Sparrows, Passer domesticus. *Ethology*, **2010** , 102, 639-648 | 1.7 | 20 |
| 68 | Historical relationships of three enigmatic phasianid genera (Aves: Galliformes) inferred using phylogenomic and mitogenomic data. *Molecular Phylogenetics and Evolution*, **2017**, 109, 217-225 | 4.1 | 18 |
| 67 | Evolutionary processes in the genus Coreocarpus: insights from molecular phylogenetics. *Evolution; International Journal of Organic Evolution*, **2003**, 57, 52-61 | 3.8 | 18 |
| 66 | Mitochondrial genome diversity and population structure of two western honey bee subspecies in the Republic of South Africa. *Scientific Reports*, **2018**, 8, 1333 | 4.9 | 17 |
| 65 | Do pups recognize maternal calls in pomona leaf-nosed bats, Hipposideros pomona?. *Animal Behaviour*, **2015**, 100, 200-207 | 2.8 | 16 |
| 64 | OCCURRENCE AND EVOLUTION OF COOPERATIVE BREEDING AMONG THE DIURNAL RAPTORS (ACCIPITRIDAE AND FALCONIDAE). *Auk*, **2003**, 120, 717 | 2.1 | 16 |
| 63 | The complete mitochondrial genome of the Cape honey bee, Esch. (Insecta: hymenoptera: apidae). *Mitochondrial DNA Part B: Resources*, **2016**, 1, 817-819 | 0.5 | 16 |
| 62 | Data Types and the Phylogeny of Neoaves. *Birds*, **2021**, 2, 1-22 | 1 | 16 |
| 61 | The complete mitochondrial genome of (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2017**, 2, 268-269 | 0.5 | 15 |
| 60 | Phylogenomic analysis suggests Coreidae and Alydidae (Hemiptera: Heteroptera) are not monophyletic. *Zoologica Scripta*, **2019**, 48, 520-534 | 2.5 | 14 |
| 59 | Fine-Scale Spatial Genetic Structure in the Cooperatively Breeding Brown-Headed Nuthatch (Sitta pusilla). *Southeastern Naturalist*, **2010**, 9, 743-756 | 0.4 | 13 |
| 58 | Out of Africa: molecular phylogenetics and biogeography of Wolffiella (Lemnaceae). *Biological Journal of the Linnean Society*, **2003**, 79, 565-576 | 1.9 | 13 |
| 57 | Comparative Genomics Reveals a Burst of Homoplasy-Free Numt Insertions. *Molecular Biology and Evolution*, **2018**, 35, 2060-2064 | 8.3 | 13 |
| 56 | The complete mitochondrial genome of the Egyptian honey bee, (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2017**, 2, 270-272 | 0.5 | 12 |
| 55 | Interspecific brood parasitism in galliform birds. *Ibis*, **2009**, 151, 373-381 | 1.9 | 12 |

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| 54 | On the origin of the Galapagos hawk: an examination of phenotypic differentiation and mitochondrial paraphyly. *Biological Journal of the Linnean Society*, **2008**, 95, 779-789 | 1.9 | 12 |
| 53 | The complex evolutionary history of big-eared horseshoe bats (Rhinolophus macrotis complex): insights from genetic, morphological and acoustic data. *Scientific Reports*, **2016**, 6, 35417 | 4.9 | 11 |
| 52 | Occurrence and Evolution of Cooperative Breeding among the Diurnal Raptors (Accipitridae and Falconidae). *Auk*, **2003**, 120, 717-729 | 2.1 | 11 |
| 51 | A molecular phylogeny of the peacock-pheasants (Galliformes Polyplectron spp.) indicates loss and reduction of ornamental traits and display behaviours. *Biological Journal of the Linnean Society*, **2001**, 73, 187-198 | 1.9 | 11 |
| 50 | The evolution of autotomy in leaf-footed bugs. *Evolution; International Journal of Organic Evolution*, **2020**, 74, 897-910 | 3.8 | 11 |
| 49 | Whole genome phylogeny of Gallus: introgression and data-type effects. *Avian Research*, **2020**, 11, | 2 | 10 |
| 48 | The mitochondrial genome of (Hymenoptera: Apidae), an Ethiopian honey bee. *Mitochondrial DNA Part B: Resources*, **2019**, 5, 9-10 | 0.5 | 10 |
| 47 | Allozyme variation within and divergence between Lemna gibba and L. disperma: Systematic and biogeographic implications. *Aquatic Botany*, **2005**, 83, 119-128 | 1.8 | 9 |
| 46 | Are Rooster Crows Honest Signals of Fighting Ability?. *Auk*, **1998**, 115, 763-766 | 2.1 | 9 |
| 45 | Pleistocene glaciation explains the disjunct distribution of the Chestnut-vented Nuthatch (Aves, Sittidae). *Zoologica Scripta*, **2019**, 48, 33-45 | 2.5 | 9 |
| 44 | Phylogenomics of manakins (Aves: Pipridae) using alternative locus filtering strategies based on informativeness. *Molecular Phylogenetics and Evolution*, **2021**, 155, 107013 | 4.1 | 9 |
| 43 | A simple strategy for recovering ultraconserved elements, exons, and introns from low coverage shotgun sequencing of museum specimens: Placement of the partridge genus Tropicoperdix within the galliformes. *Molecular Phylogenetics and Evolution*, **2018**, 129, 304-314 | 4.1 | 9 |
| 42 | The complete mitochondrial genome of an east African honey bee, Smith (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2017**, 2, 589-590 | 0.5 | 8 |
| 41 | The complete mitochondrial genome and phylogenetic placement of Smith (Insecta: Hymenoptera: Apidae), an Asian, cavity-nesting honey bee. *Mitochondrial DNA Part B: Resources*, **2017**, 2, 249-250 | 0.5 | 8 |
| 40 | Allozyme studies in Lemnaceae: variation and relationships in Lemna sections Alatae and Biformes. *Taxon*, **2001**, 50, 987-999 | 0.8 | 8 |
| 39 | The complete mitochondrial genome of (Insecta: Hymenoptera: Apidae), the Malagasy honey bee. *Mitochondrial DNA Part B: Resources*, **2019**, 4, 3286-3287 | 0.5 | 7 |
| 38 | Inter-simple sequence repeat (ISSR) diversity within Monarda fistulosa var. brevis (Lamiaceae) and divergence between var. brevis and var. fistulosa in West Virginia. *Brittonia*, **2001**, 53, 511-518 | 0.5 | 7 |
| 37 | The complete mitochondrial genome of (Insecta: Hymenoptera: Apidae), the Arabian honey bee. *Mitochondrial DNA Part B: Resources*, **2020**, 5, 875-876 | 0.5 | 6 |

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| 36 | Nestmate killing by obligate brood parasitic chicks: is this linked to obligate siblicidal behavior?. *Journal of Ornithology*, **2012**, 153, 825-831 | 1.5 | 6 |
| 35 | Isolation and characterization of polymorphic microsatellite markers for the brown-headed nuthatch (Sitta pusilla). *Conservation Genetics*, **2009**, 10, 1393-1395 | 2.6 | 6 |
| 34 | Molecular and morphological divergence in a pair of bird species and their ectoparasites. *Journal of Parasitology*, **2009**, 95, 1372-82 | 0.9 | 6 |
| 33 | The mitochondrial genome of the Spanish honey bee, (Insecta: Hymenoptera: Apidae), from Portugal. *Mitochondrial DNA Part B: Resources*, **2019**, 5, 17-18 | 0.5 | 5 |
| 32 | The complete mitochondrial genome of the hybrid honey bee, , from South Africa. *Mitochondrial DNA Part B: Resources*, **2016**, 1, 856-857 | 0.5 | 5 |
| 31 | Uncommon Levels of Relatedness and Parentage in a Cooperatively Breeding Bird, the Brown-Headed Nuthatch (Sitta pusilla). *Wilson Journal of Ornithology*, **2015**, 127, 593-600 | 0.4 | 5 |
| 30 | A multigene phylogeny of Galliformes supports a single origin of erectile ability in non-feathered facial traits. *Journal of Avian Biology*, **2008**, 39, ???-??? | 1.9 | 5 |
| 29 | The generic placement of a morphologically enigmatic species in Asteraceae: evidence from ITS sequences. *Plant Systematics and Evolution*, **2001**, 228, 63-69 | 1.3 | 5 |
| 28 | Does the use of a multi-trait, multi-test approach to´measure animal personality yield different behavioural´syndrome results?. *Behaviour*, **2018**, 155, 115-150 | 1.4 | 5 |
| 27 | The complete mitochondrial genome of the West African honey bee (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2019**, 5, 11-12 | 0.5 | 4 |
| 26 | Phylogeny and diversification of the gallopheasants (Aves: Galliformes): Testing roles of sexual selection and environmental niche divergence. *Zoologica Scripta*, **2020**, 49, 549-562 | 2.5 | 4 |
| 25 | A phylogenomic supermatrix of Galliformes (Landfowl) reveals biased branch lengths. *Molecular Phylogenetics and Evolution*, **2021**, 158, 107091 | 4.1 | 4 |
| 24 | The mitochondrial genome of the Maltese honey bee, (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2020**, 5, 877-878 | 0.5 | 3 |
| 23 | Testing hypotheses driving genetic structure in the cooperatively breeding Brown-headed Nuthatch Sitta pusilla. *Ibis*, **2019**, 161, 387-400 | 1.9 | 3 |
| 22 | The mitochondrial genome of the Carniolan honey bee, (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2019**, 4, 3288-3290 | 0.5 | 3 |
| 21 | Genetic divergence among Snail Kite subspecies: implications for the conservation of the endangered Florida Snail Kite Rostrhamus sociabilis. *Ibis*, **2009**, 151, 181-185 | 1.9 | 3 |
| 20 | Divergence time estimation of Galliformes based on the best gene shopping scheme of ultraconserved elements. *Bmc Ecology and Evolution*, **2021**, 21, 209 | 21 | 3 |
| 19 | Phylogenomics of the Leaf-Footed Bug Subfamily Coreinae (Hemiptera: Coreidae). *Insect Systematics and Diversity*, **2020**, 4, | 1.8 | 3 |

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| 18 | Kin-biased conspecific brood parasitism in a native Mandarin duck population. *Journal of Ornithology*, **2016**, 157, 1063-1072 | 1.5 | 3 |
| 17 | When good mitochondria go bad: Cyto-nuclear discordance in landfowl (Aves: Galliformes). *Gene*, **2021**, 801, 145841 | 3.8 | 3 |
| 16 | Phylogenetic definitions for 25 higher-level clade names of birds. *Avian Research*, **2022**, 100027 | 2 | 3 |
| 15 | Coreopsis sect. Pseudoagarista (Asteraceae: Coreopsideae): Molecular phylogeny, chromosome numbers, and comments on taxonomy and distribution. *Taxon*, **2014**, 63, 1092-1102 | 0.8 | 2 |
| 14 | Protein Structure, Models of Sequence Evolution, and Data Type Effects in Phylogenetic Analyses of Mitochondrial Data: A Case Study in Birds. *Diversity*, **2021**, 13, 555 | 2.5 | 2 |
| 13 | Habitat urbanization and stress response are primary predictors of personality variation in northern cardinals (Cardinalis cardinalis). *Journal of Urban Ecology*, **2020**, 6, | 2 | 2 |
| 12 | Strong phenotypic divergence in spite of low genetic structure in the endemic Mangrove Warbler subspecies () of Costa Rica. *Ecology and Evolution*, **2019**, 9, 13902-13918 | 2.8 | 2 |
| 11 | Ecological niche differentiation in Chiroxiphia and Antilophia manakins (Aves: Pipridae). *PLoS ONE*, **2021**, 16, e0243760 | 3.7 | 2 |
| 10 | Re-evaluating the distribution of cooperative breeding in birds: is it tightly linked with altriciality?. *Journal of Avian Biology*, **2016**, 47, 724-730 | 1.9 | 1 |
| 9 | Gene tree quality affects empirical coalescent branch length estimation. *Zoologica Scripta*, | 2.5 | 1 |
| 8 | Extracting "legacy loci" from an invertebrate sequence capture data set. *Zoologica Scripta*, | 2.5 | 1 |
| 7 | Phylogenomic analysis with improved taxon sampling corroborates an Alydidae + Hydarinae + Pseudophloeinae clade (Heteroptera: Coreoidea: Alydidae, Coreidae). *Organisms Diversity and Evolution*,1 | 1.7 | 1 |
| 6 | Dancing drives evolution of sexual size dimorphism in manakins.. *Proceedings of the Royal Society B: Biological Sciences*, **2022**, 289, 20212540 | 4.4 | 1 |
| 5 | Dynamic Patterns of Sex Chromosome Evolution in Neognath Birds: Many Independent Barriers to Recombination at the ATP5F1A Locus. *Birds*, **2022**, 3, 51-70 | 1 | 0 |
| 4 | Occurrence and Evolution of Cooperative Breeding Among the Diurnal Raptors (Accipitridae and Falconidae). *Auk*, **2003**, 120, 717-729 | 2.1 | 0 |
| 3 | The complete mitochondrial genome of Tingek, an Asian honey bee (Insecta: Hymenoptera: Apidae). *Mitochondrial DNA Part B: Resources*, **2017**, 2, 552-553 | 0.5 | |
| 2 | Phylogenetic Utility of Avian Ovomucoid Intron G: A Comparison of Nuclear and Mitochondrial Phylogenies in Galliformes. *Auk*, **2001**, 118, 799-804 | 2.1 | |
| 1 | Genome-wide assessment of population structure in Florida's coastal seaside sparrows. *Conservation Genetics*,1 | 2.6 | |