Hong-Yi Liu

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| # | Paper | IF | Citations |
|----|--|-----|-----------|
| 34 | Five new mitogenomes of Phylloscopus (Passeriformes, Phylloscopidae): Sequence, structure, and phylogenetic analyses. <i>International Journal of Biological Macromolecules</i> , 2020 , 146, 638-647 | 7.9 | 16 |
| 33 | Analysis of the Gut Microbiome of Wild and Captive PEe David's Deer. <i>Frontiers in Microbiology</i> , 2019 , 10, 2331 | 5.7 | 12 |
| 32 | Potential natural exposure of endangered red-crowned crane (Grus japonensis) to mycotoxins aflatoxin B1, deoxynivalenol, zearalenone, T-2 toxin, and ochratoxin A. <i>Journal of Zhejiang University: Science B</i> , 2016 , 17, 158-68 | 4.5 | 9 |
| 31 | Comparative analysis of the gut microbiota of hornbill and toucan in captivity. <i>MicrobiologyOpen</i> , 2018 , 8, e786 | 3.4 | 9 |
| 30 | Mitogenome of the little owl Athene noctua and phylogenetic analysis of Strigidae. <i>International Journal of Biological Macromolecules</i> , 2020 , 151, 924-931 | 7.9 | 8 |
| 29 | Characterization and comparison of gut microbiomes in nine species of parrots in captivity. <i>Symbiosis</i> , 2019 , 78, 241-250 | 3 | 8 |
| 28 | The complete mitochondrial genome of (Oligochaeta, Lumbricidae) and its phylogenetic analysis. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2677-2678 | 0.5 | 6 |
| 27 | Mitochondrial Genome Structures and Phylogenetic Analyses of Two Tropical Characidae Fishes. <i>Frontiers in Genetics</i> , 2021 , 12, 627402 | 4.5 | 6 |
| 26 | Limited polymorphism of the functional MHC class II B gene in the black-spotted frog () identified by locus-specific genotyping. <i>Ecology and Evolution</i> , 2017 , 7, 9860-9868 | 2.8 | 4 |
| 25 | Factors affecting genotyping success in giant panda fecal samples. <i>PeerJ</i> , 2017 , 5, e3358 | 3.1 | 4 |
| 24 | Characterization of the complete mitochondrial genome of Drawida gisti (Metagynophora, Moniligastridae) and comparison with other Metagynophora species. <i>Genomics</i> , 2020 , 112, 3056-3064 | 4.3 | 4 |
| 23 | Genetic diversity and structure of the round-tailed paradise fish (Macropodus ocellatus): Implications for population management. <i>Global Ecology and Conservation</i> , 2020 , 21, e00876 | 2.8 | 4 |
| 22 | Mitochondrial genomes of four American characins and phylogenetic relationships within the family Characidae (Teleostei: Characiformes). <i>Gene</i> , 2020 , 762, 145041 | 3.8 | 3 |
| 21 | Development and characterization of microsatellite markers in the earthworm Drawida gisti Michaelsen, 1931 and cross-amplification in two other congeners. <i>Molecular Biology Reports</i> , 2020 , 47, 8265-8269 | 2.8 | 3 |
| 20 | Genetic diversity of wild wintering red-crowned crane () by microsatellite markers and mitochondrial gene sequence in the Yancheng reserve. <i>Animal Biotechnology</i> , 2021 , 32, 531-536 | 1.4 | 2 |
| 19 | Complete mitogenomic and phylogenetic characteristics of the speckled wood-pigeon (Columba hodgsonii). <i>Molecular Biology Reports</i> , 2020 , 47, 3567-3576 | 2.8 | 2 |
| 18 | Characterization of MHC class II A genes in Hainan EldE deer (Cervus eldi hainanus). <i>Science Bulletin</i> , 2013 , 58, 2148-2153 | | 2 |

LIST OF PUBLICATIONS

| 17 | MHC class II Igenes in the endangered Hainan Eld's deer (Cervus eldi hainanus). <i>Journal of Heredity</i> , 2013 , 104, 874-80 | 2.4 | 2 |
|----|---|-----|---|
| 16 | The complete mitochondrial genome sequence of Yarkand hare (). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3727-3728 | 0.5 | 2 |
| 15 | The complete mitogenome of Hong Kong paradise fish (), an endemic freshwater fish in South China. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2849-2850 | 0.5 | 1 |
| 14 | The complete mitogenome of red-collared lorikeet () and its phylogenetic analysis. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3116-3117 | 0.5 | 1 |
| 13 | Characterization of Five New Earthworm Mitogenomes (Oligochaeta: Lumbricidae): Mitochondrial Phylogeny of Lumbricidae. <i>Diversity</i> , 2021 , 13, 580 | 2.5 | 1 |
| 12 | Effects of Montmorillonite on Growth Performance, Serum Biochemistry and Oxidative Stress of Red-Crowned Crane () Fed Mycotoxin-Contaminated Feed. <i>Current Drug Metabolism</i> , 2020 , 21, 626-632 | 3.5 | 1 |
| 11 | Development and characterization of 33 SNP markers for the red-crowned crane (Grus japonensis) using a genotyping-by-sequencing approach. <i>Conservation Genetics Resources</i> , 2020 , 12, 385-387 | 0.8 | 1 |
| 10 | The mitochondrial genome and phylogenetic characteristics of the Thick-billed Green-Pigeon, : the first sequence for the genus. <i>ZooKeys</i> , 2021 , 1041, 167-182 | 1.2 | 1 |
| 9 | Genetic Diversity and Population Structure of Based on the Mitochondrial COI Gene and Microsatellites. <i>Frontiers in Genetics</i> , 2021 , 12, 686246 | 4.5 | 1 |
| 8 | Polymorphism and varying selection within the MHC class I of four Anas species. <i>Immunogenetics</i> , 2021 , 73, 395-404 | 3.2 | 1 |
| 7 | Comparative mitogenomes of three species in Moenkhausia: Rare irregular gene rearrangement within Characidae. <i>International Journal of Biological Macromolecules</i> , 2021 , 183, 1079-1086 | 7.9 | 1 |
| 6 | Mitochondrial genomes of five tetras and their phylogenetic implications. <i>Ecology and Evolution</i> , 2021 , 11, 12754-12764 | 2.8 | 1 |
| 5 | Historical gene flow and profound spatial genetic structure among golden pheasant populations suggested by multi-locus analysis. <i>Molecular Phylogenetics and Evolution</i> , 2017 , 110, 93-103 | 4.1 | O |
| 4 | MHC-associated load informs the giant panda reintroduction program. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2020 , 12, 113-120 | 2.6 | О |
| 3 | Comprehensive transcriptome characterization of Grus japonensis using PacBio SMRT and Illumina sequencing <i>Scientific Reports</i> , 2021 , 11, 23927 | 4.9 | 0 |
| 2 | Two complete mitochondrial genomes in Scolopendra and a comparative analysis of tRNA rearrangements in centipedes <i>Molecular Biology Reports</i> , 2022 , 1 | 2.8 | 0 |
| 1 | Development and characterization of 12 microsatellite markers for the roundtail paradisefish (Macropodus ocellatus) using RAD sequencing. <i>Journal of Applied Ichthyology</i> , 2020 , 36, 243-245 | 0.9 | |