

Hong-Yi Liu

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

34
papers

133
citations

7
h-index

9
g-index

44
ext. papers

236
ext. citations

2.9
avg, IF

3.42
L-index

#	Paper	IF	Citations
34	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
33	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
32	Characterization of Five New Earthworm Mitogenomes (Oligochaeta: Lumbricidae): Mitochondrial Phylogeny of Lumbricidae. <i>Diversity</i> , 2021 , 13, 580	2.5	1
31	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
30	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
29	Polymorphism and varying selection within the MHC class I of four Anas species. <i>Immunogenetics</i> , 2021 , 73, 395-404	3.2	1
28	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
27	Mitochondrial Genome Structures and Phylogenetic Analyses of Two Tropical Characidae Fishes. <i>Frontiers in Genetics</i> , 2021 , 12, 627402	4.5	6
26	Mitochondrial genomes of five tetras and their phylogenetic implications. <i>Ecology and Evolution</i> , 2021 , 11, 12754-12764	2.8	1
25	Comprehensive transcriptome characterization of Grus japonensis using PacBio SMRT and Illumina sequencing.. <i>Scientific Reports</i> , 2021 , 11, 23927	4.9	0
24	MHC-associated load informs the giant panda reintroduction program. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2020 , 12, 113-120	2.6	0
23	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	
22	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
21	Complete mitogenomic and phylogenetic characteristics of the speckled wood-pigeon (Columba hodgsonii). <i>Molecular Biology Reports</i> , 2020 , 47, 3567-3576	2.8	2
20	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
19	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
18	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

17	Genetic diversity and structure of the round-tailed paradise fish (<i>Macropodus ocellatus</i>): Implications for population management. <i>Global Ecology and Conservation</i> , 2020 , 21, e00876	2.8	4
16	Five new mitogenomes of <i>Phylloscopus</i> (Passeriformes, Phylloscopidae): Sequence, structure, and phylogenetic analyses. <i>International Journal of Biological Macromolecules</i> , 2020 , 146, 638-647	7.9	16
15	Mitochondrial genomes of four American characins and phylogenetic relationships within the family Characidae (Teleostei: Characiformes). <i>Gene</i> , 2020 , 762, 145041	3.8	3
14	Development and characterization of microsatellite markers in the earthworm <i>Drawida gisti</i> Michaelsen, 1931 and cross-amplification in two other congeners. <i>Molecular Biology Reports</i> , 2020 , 47, 8265-8269	2.8	3
13	The complete mitochondrial genome of (<i>Oligochaeta</i> , Lumbricidae) and its phylogenetic analysis. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2677-2678	0.5	6
12	The complete mitogenome of Hong Kong paradise fish (<i></i>), an endemic freshwater fish in South China. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 2849-2850	0.5	1
11	Characterization and comparison of gut microbiomes in nine species of parrots in captivity. <i>Symbiosis</i> , 2019 , 78, 241-250	3	8
10	Analysis of the Gut Microbiome of Wild and Captive Phe David's Deer. <i>Frontiers in Microbiology</i> , 2019 , 10, 2331	5.7	12
9	The complete mitogenome of red-collared lorikeet (<i></i>) and its phylogenetic analysis. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3116-3117	0.5	1
8	The complete mitochondrial genome sequence of Yarkand hare (<i></i>). <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 3727-3728	0.5	2
7	Comparative analysis of the gut microbiota of hornbill and toucan in captivity. <i>MicrobiologyOpen</i> , 2018 , 8, e786	3.4	9
6	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	0
5	Limited polymorphism of the functional MHC class II B gene in the black-spotted frog (<i></i>) identified by locus-specific genotyping. <i>Ecology and Evolution</i> , 2017 , 7, 9860-9868	2.8	4
4	Factors affecting genotyping success in giant panda fecal samples. <i>PeerJ</i> , 2017 , 5, e3358	3.1	4
3	Potential natural exposure of endangered red-crowned crane (<i>Grus japonensis</i>) 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
2	Characterization of MHC class II A genes in Hainan Eld's deer (<i>Cervus eldi hainanus</i>). <i>Science Bulletin</i> , 2013 , 58, 2148-2153		2
1	MHC class II genes in the endangered Hainan Eld's deer (<i>Cervus eldi hainanus</i>). <i>Journal of Heredity</i> , 2013 , 104, 874-80	2.4	2