

De Chen

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

179
citations

1163117

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1199594

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docs citations

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times ranked

241
citing authors

#	ARTICLE	IF	CITATIONS
1	A pipeline for effectively developing highly polymorphic simple sequence repeats markers based on multi-sample genomic data. <i>Ecology and Evolution</i> , 2022, 12, e8705.	1.9	5
2	Disentangling the relative roles of geographical and ecological factors in driving genomic variations of a widely distributed bird across a longitudinal gradient. <i>Journal of Avian Biology</i> , 2022, 2022, .	1.2	0
3	Genomic Consequences of Long-Term Population Decline in Brown Eared Pheasant. <i>Molecular Biology and Evolution</i> , 2021, 38, 263-273.	8.9	36
4	Population genomic, climatic and anthropogenic evidence suggest the role of human forces in endangerment of green peafowl (<i>Pavo muticus</i>). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20210073.	2.6	26
5	Comparative analysis of mite genomes reveals positive selection for diet adaptation. <i>Communications Biology</i> , 2021, 4, 668.	4.4	6
6	Divergence time estimation of Galliformes based on the best gene shopping scheme of ultraconserved elements. <i>Bmc Ecology and Evolution</i> , 2021, 21, 209.	1.6	17
7	Disentangling the evolutionary history and biogeography of hill partridges (Phasianidae, Arborophila) from low coverage shotgun sequences. <i>Molecular Phylogenetics and Evolution</i> , 2020, 151, 106895.	2.7	2
8	Genome assembly of the common pheasant <i>Phasianus colchicus</i> , a model for speciation and ecological genomics. <i>Genome Biology and Evolution</i> , 2019, 11, 3326-3331.	2.5	6
9	Demographic Histories and Genome-Wide Patterns of Divergence in Incipient Species of Shorebirds. <i>Frontiers in Genetics</i> , 2019, 10, 919.	2.3	14
10	A simple strategy for recovering ultraconserved elements, exons, and introns from low coverage shotgun sequencing of museum specimens: Placement of the partridge genus <i>Tropicoperdix</i> within the galliformes. <i>Molecular Phylogenetics and Evolution</i> , 2018, 129, 304-314.	2.7	13
11	Characterization of novel microsatellite markers of the Emei Shan <i>Liocichla</i> using restriction site-associated DNA sequencing. <i>Avian Research</i> , 2017, 8, .	1.2	5
12	Revival of the genus <i>Tropicoperdix</i> Blyth 1859 (Phasianidae, Aves) using multilocus sequence data. <i>Zoological Journal of the Linnean Society</i> , 2015, 175, 429-438.	2.3	10
13	Was the exposed continental shelf a long-distance colonization route in the ice age? The Southeast Asia origin of Hainan and Taiwan partridges. <i>Molecular Phylogenetics and Evolution</i> , 2015, 83, 167-173.	2.7	18
14	Species delimitation in the Chinese bamboo partridge <i>Bambusicola thoracica</i> (Phasianidae; Phasianinae). <i>Zoologica Scripta</i> , 2014, 43, 562-575.	1.7	15
15	Molecular demographic history of the Hainan Peacock Pheasant (<i>Polyplectron katsumatae</i>) and its conservation implications. <i>Science Bulletin</i> , 2013, 58, 2185-2190.	1.7	6