

Yinchang Hu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5141326/publications.pdf>

Version: 2024-02-01

16
papers

237
citations

1163117

8
h-index

996975

15
g-index

18
all docs

18
docs citations

18
times ranked

558
citing authors

#	ARTICLE	IF	CITATIONS
1	A global analysis of enemy release and its variation with latitude. <i>Global Ecology and Biogeography</i> , 2021, 30, 277-288.	5.8	15
2	The complete mitochondrial genome of <i>Moolgarda perusii</i> (Teleostei: Mugilidae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 93-94.	0.4	0
3	Do non-native ornamental fishes pose a similar level of invasion risk in neighbouring regions of similar current and future climate? Implications for conservation and management. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2021, 31, 2041-2057.	2.0	11
4	The complete mitochondrial genome of longhorn cowfish, <i>Lactoria cornuta</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 3749-3750.	0.4	0
5	Tissue expression and bioinformatics analysis of the vitellogenin gene of Asian arowana (<i>Scleropages formosus</i>). <i>Journal of Herpetology</i> , 2021, 55, 1-10.	0.7	4
6	The influence of warming on the biogeographic and phylogenetic dependence of herbivore-plant interactions. <i>Ecology and Evolution</i> , 2019, 9, 2231-2241.	1.9	4
7	Current status and potential risks of established alien fish species in China. <i>Aquatic Ecosystem Health and Management</i> , 2019, 22, 371-384.	0.6	5
8	China is initiating the Aquatic 10-100-1,000 Genomics Program. <i>Science China Life Sciences</i> , 2017, 60, 329-332.	4.9	6
9	A chromosome-level genome assembly of the Asian arowana, <i>Scleropages formosus</i> . <i>Scientific Data</i> , 2016, 3, 160105.	5.3	13
10	The Asian arowana (<i>Scleropages formosus</i>) genome provides new insights into the evolution of an early lineage of teleosts. <i>Scientific Reports</i> , 2016, 6, 24501.	3.3	89
11	Warming mediates the relationship between plant nutritional properties and herbivore functional responses. <i>Ecology and Evolution</i> , 2016, 6, 8777-8784.	1.9	11
12	Spatial variation in adult sex ratio across multiple scales in the invasive golden apple snail, <i>Pomacea canaliculata</i> . <i>Ecology and Evolution</i> , 2016, 6, 2308-2317.	1.9	12
13	Comparative Functional Responses Predict the Invasiveness and Ecological Impacts of Alien Herbivorous Snails. <i>PLoS ONE</i> , 2016, 11, e0147017.	2.5	26
14	Characterization of the <i>Macropodus opercularis</i> complete mitochondrial genome and family Channidae taxonomy using Illumina-based de novo transcriptome sequencing. <i>Gene</i> , 2015, 559, 189-195.	2.2	24
15	An unusual mitochondrial genome structure of the tonguefish, <i>Cynoglossus trigrammus</i> : Control region translocation and a long additional non-coding region inversion. <i>Gene</i> , 2015, 573, 216-224.	2.2	12
16	Characterization of the mitochondrial genome and phylogeny of the black arowana (<i>Osteoglossum</i>). <i>Journal of Herpetology</i> , 2015, 49, 1-10.	0.5	1