

Dengqiang Wang

List of Publications by Year in descending order

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papers

176
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1478505

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#	ARTICLE	IF	CITATIONS
1	Using environmental DNA to detect <i>Hypophthalmichthys molitrix</i> during the spawning period in the Yangtze River. <i>Conservation Genetics Resources</i> , 2020, 12, 37-39.	0.8	1
2	Population genetics and sympatric divergence of the freshwater gudgeon, <i>Gobiobotia filifer</i> , in the Yangtze River inferred from mitochondrial DNA. <i>Ecology and Evolution</i> , 2020, 10, 50-58.	1.9	1
3	The mitochondrial genome of <i>Culter oxycephaloides</i> (Cypriniformes, Cyprinidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2392-2393.	0.4	1
4	Development of 28 EST-SSR markers based on transcriptome sequences of <i>Gobiobotia filifer</i> and cross-species amplification. <i>Journal of Applied Ichthyology</i> , 2019, 35, 1295-1299.	0.7	1
5	Monitoring the genetic effects of broodstock enhancement of silver carp (<i>Hypophthalmichthys</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 62 Td <i>Freshwater Ecology</i> , 2019, 34, 323-332.	1.2	3
6	Hatchery-reared enhancement program for silver carp (<i>Hypophthalmichthys molitrix</i>) in the middle Yangtze River: monitoring the effectiveness based on parentage analysis. <i>PeerJ</i> , 2019, 7, e6836.	2.0	2
7	Genetic structure of two sympatric gudgeon fishes (<i>Xenophysogobio boulengeri</i> and <i>X.</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 62 Td	2.0	3
8	Species diversity of drifting fish eggs in the Yangtze River using molecular identification. <i>PeerJ</i> , 2018, 6, e5807.	2.0	18
9	The complete mitochondrial genome of <i>Triplophysa robusta</i> (Teleostei, Cypriniformes) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 62 Td	0.6	2
10	The complete mitochondrial genome of <i>Xenophysogobio nudicorpa</i> (Teleostei, Cypriniformes,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 Td	0.6	2
11	The complete mitochondrial genome of <i>Phoxinus lagowskii</i> (Teleostei, Cypriniformes) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 62 Td	0.6	6
12	DNA barcoding and evaluation of genetic diversity in Cyprinidae fish in the midstream of the Yangtze River. <i>Ecology and Evolution</i> , 2016, 6, 2702-2713.	1.9	60
13	The mitochondrial genome of <i>Pseudolaubuca sinensis</i> (Cypriniformes, Cyprinidae). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016, 27, 3356-3357.	0.7	1
14	The mitochondrial genome of <i>Gnathopogon imberbis</i> (Cypriniformes, Cyprinidae). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016, 27, 2543-2544.	0.7	3
15	Characterization and cross-species amplification of 19 polymorphic microsatellite loci of <i>Hemibagrus macropterus</i> (Bleeker) in the Yangtze River. <i>Conservation Genetics Resources</i> , 2015, 7, 5-8.	0.8	0
16	Development of 27 novel cross-species microsatellite markers for the endangered <i>Hucho bleekeri</i> using next-generation sequencing technology. <i>Conservation Genetics Resources</i> , 2015, 7, 263-267.	0.8	12
17	Isolation and characterization of twelve microsatellite loci of <i>Leiocassis albomarginatus</i> (Rendhal) in the Yangtze River. <i>Conservation Genetics Resources</i> , 2015, 7, 315-317.	0.8	1
18	Isolation and characterization of 23 microsatellite loci in the Chinese sucker (<i>Myxocyprinus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 Td	0.8	4

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19	Characterization and cross-species amplification of 14 polymorphic microsatellite loci in <i>Xenophysogobio boulengeri</i> . <i>Conservation Genetics Resources</i> , 2012, 4, 1015-1017.	0.8	1
20	Genetic diversity in the bronze gudgeon, <i>Coreius heterodon</i> , from the Yangtze River system based on mtDNA sequences of the control region. <i>Environmental Biology of Fishes</i> , 2008, 82, 35-40.	1.0	13
21	Cloning and expression analysis of a prion protein encoding gene in guppy (<i>Poecilia reticulata</i>). <i>Journal of Ocean University of China</i> , 2008, 7, 425-431.	1.2	0
22	Development of microsatellite DNA markers of silver carp (<i>Hypophthalmichthys molitrix</i>) and their cross-species application in bighead carp (<i>Aristichthys nobilis</i>). <i>Molecular Ecology Notes</i> , 2006, 7, 95-99.	1.7	18
23	Molecular phylogenetic systematics of twelve species of Acipenseriformes based on mtDNA ND4 gene sequence analysis. <i>Science in China Series C: Life Sciences</i> , 2000, 43, 129-137.	1.3	23