

# Dengqiang Wang

## List of Publications by Year in descending order

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23  
papers

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citations

1478505

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times ranked

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citing authors

#	ARTICLE	IF	CITATIONS
1	<sc>DNA</sc> barcoding and evaluation of genetic diversity in Cyprinidae fish in the midstream of the Yangtze River. Ecology and Evolution, 2016, 6, 2702-2713.	1.9	60
2	Molecular phylogenetic systematics of twelve species of Acipenseriformes based on mtDNA ND4 gene sequence analysis. Science in China Series C: Life Sciences, 2000, 43, 129-137.	1.3	23
3	Development of microsatellite DNA markers of silver carp ( <i>Hypophthalmichthys molitrix</i> ) and their cross-species application in bighead carp ( <i>Aristichthys nobilis</i> ). Molecular Ecology Notes, 2006, 7, 95-99.	1.7	18
4	Species diversity of drifting fish eggs in the Yangtze River using molecular identification. PeerJ, 2018, 6, e5807.	2.0	18
5	Genetic diversity in the bronze gudgeon, <i>Coreius heterodon</i> , from the Yangtze River system based on mtDNA sequences of the control region. Environmental Biology of Fishes, 2008, 82, 35-40.	1.0	13
6	Development of 27 novel cross-species microsatellite markers for the endangered <i>Hucho bleekeri</i> using next-generation sequencing technology. Conservation Genetics Resources, 2015, 7, 263-267.	0.8	12
7	The complete mitochondrial genome of <i>Phoxinus lagowskii</i> (Teleostei, Cypriniformes). Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 462 T	0.6	6
8	Isolation and characterization of 23 microsatellite loci in the Chinese sucker ( <i>Myxocyprinus</i> ). Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 462 T	0.8	4
9	The mitochondrial genome of <i>Gnathopogon imberbis</i> (Cypriniformes, Cyprinidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 2543-2544.	0.7	3
10	Monitoring the genetic effects of broodstock enhancement of silver carp ( <i>Hypophthalmichthys</i> ). Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 462 T Freshwater Ecology, 2019, 34, 323-332.	1.2	3
11	Genetic structure of two sympatric gudgeon fishes ( <i>Xenophysogobio boulengeri</i> and <i>X.</i> ). Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 462 T	2.0	3
12	The complete mitochondrial genome of <i>Triplophysa robusta</i> (Teleostei, Cypriniformes). Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 302 T	0.6	2
13	The complete mitochondrial genome of <i>Xenophysogobio nudicorpa</i> (Teleostei, Cypriniformes). Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 462 T	0.6	2
14	Hatchery-reared enhancement program for silver carp ( <i>Hypophthalmichthys molitrix</i> ) in the middle Yangtze River: monitoring the effectiveness based on parentage analysis. PeerJ, 2019, 7, e6836.	2.0	2
15	Characterization and cross-species amplification of 14 polymorphic microsatellite loci in <i>Xenophysogobio boulengeri</i> . Conservation Genetics Resources, 2012, 4, 1015-1017.	0.8	1
16	Isolation and characterization of twelve microsatellite loci of <i>Leiocassis albomarginatus</i> (Rendhal) in the Yangtze River. Conservation Genetics Resources, 2015, 7, 315-317.	0.8	1
17	The mitochondrial genome of <i>Pseudolaubuca sinensis</i> (Cypriniformes, Cyprinidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 3356-3357.	0.7	1
18	The mitochondrial genome of <i>Culter oxycephaloides</i> (Cypriniformes, Cyprinidae). Mitochondrial DNA Part B: Resources, 2019, 4, 2392-2393.	0.4	1

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19	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
20	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
21	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
22	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
23	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