Yong Zhang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	WEGO: a web tool for plotting GO annotations. Nucleic Acids Research, 2006, 34, W293-W297.	14.5	2,529
2	SOAPnuke: a MapReduce acceleration-supported software for integrated quality control and preprocessing of high-throughput sequencing data. GigaScience, 2018, 7, 1-6.	6.4	1,265
3	The draft genome of the grass carp (Ctenopharyngodon idellus) provides insights into its evolution and vegetarian adaptation. Nature Genetics, 2015, 47, 625-631.	21.4	352
4	The kiss/kissr Systems Are Dispensable for Zebrafish Reproduction: Evidence From Gene Knockout Studies. Endocrinology, 2015, 156, 589-599.	2.8	153
5	Structural diversity of the gnih/gnih receptor system in teleost: Its involvement in early development and the negative control of LH release. Peptides, 2010, 31, 1034-1043.	2.4	145
6	Mudskipper genomes provide insights into the terrestrial adaptation of amphibious fishes. Nature Communications, 2014, 5, 5594.	12.8	135
7	Molecular Identification of the Kiss2/Kiss1ra System and Its Potential Function During 17Alpha-Methyltestosterone-Induced Sex Reversal in the Orange-Spotted Grouper, Epinephelus coioides1. Biology of Reproduction, 2010, 83, 63-74.	2.7	96
8	Transcriptome analysis reveals the molecular mechanisms underlying growth superiority in a novel grouper hybrid (Epinephelus fuscogutatus♀ × E. lanceolatusâ™,). BMC Genetics, 2016, 17, 24.	2.7	94
9	Molecular cloning, characterization and expression profiles of multiple leptin genes and a leptin receptor gene in orange-spotted grouper (Epinephelus coioides). General and Comparative Endocrinology, 2013, 181, 295-305.	1.8	88
10	Signatures of selection in tilapia revealed by whole genome resequencing. Scientific Reports, 2015, 5, 14168.	3.3	86
11	A novel neuropeptide in suppressing luteinizing hormone release in goldfish, Carassius auratus. Molecular and Cellular Endocrinology, 2013, 374, 65-72.	3.2	83
12	Evidences for the regulation of GnRH and GTH expression by GnIH in the goldfish, Carassius auratus. Molecular and Cellular Endocrinology, 2013, 366, 9-20.	3.2	83
13	Orange-spotted grouper (Epinephelus coioides) toll-like receptor 22: Molecular characterization, expression pattern and pertinent signaling pathways. Fish and Shellfish Immunology, 2012, 33, 494-503.	3.6	78
14	Molecular identification of GnIH/GnIHR signal and its reproductive function in protogynous hermaphroditic orange-spotted grouper (Epinephelus coioides). General and Comparative Endocrinology, 2015, 216, 9-23.	1.8	64
15	Two distinct cytochrome P450 aromatases in the orange-spotted grouper (Epinephelus coioides): cDNA cloning and differential mRNA expression. Journal of Steroid Biochemistry and Molecular Biology, 2004, 92, 39-50.	2.5	62
16	Interleukin-1β gene in orange-spotted grouper, Epinephelus coioides: Molecular cloning, expression, biological activities and signal transduction. Molecular Immunology, 2008, 45, 857-867.	2.2	62
17	Genetic Evidence for Multifactorial Control of the Reproductive Axis in Zebrafish. Endocrinology, 2017, 158, 604-611.	2.8	62
18	Spexin Suppress Food Intake in Zebrafish: Evidence from Gene Knockout Study. Scientific Reports, 2017, 7, 14643.	3.3	61

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19	The mRNA expression of P450 aromatase, gonadotropin βâ€subunits and FTZâ€F1 in the orangeâ€spotted grouper (<i>Epinephelus Coioides</i>) during 17αâ€methyltestosteroneâ€induced precocious sex change. Molecular Reproduction and Development, 2007, 74, 665-673.	2.0	59
20	Molecular cloning and functional characterization of spexin in orange-spotted grouper (Epinephelus) Tj ETQqO 196-197, 85-91.	0 0 rgBT /0 1.6	Overlock 10 Tf 50
21	Gene knockout of nuclear progesterone receptor provides insights into the regulation of ovulation by LH signaling in zebrafish. Scientific Reports, 2016, 6, 28545.	3.3	49
22	De novo assembly of a chromosomeâ€level reference genome of redâ€spotted grouper (<i>Epinephelus) Tj ETQ</i>	q0 0 0 rgE 4.8	3T /Qyerlock 10
23	The evolution of tachykinin/tachykinin receptor (TAC/TACR) in vertebrates and molecular identification of the TAC3/TACR3 system in zebrafish (Danio rerio). Molecular and Cellular Endocrinology, 2012, 361, 202-212.	3.2	44
24	Molecular regulation of sex change induced by methyltestosterone -feeding and methyltestosterone -feeding withdrawal in the protogynous orange-spotted grouperâ€. Biology of Reproduction, 2017, 97, 324-333.	2.7	40
25	Construction of high-density genetic linkage maps for orange-spotted grouper Epinephelus coioidesusing multiplexed shotgun genotyping. BMC Genetics, 2013, 14, 113.	2.7	39
26	Sexual Dimorphism of Steroidogenesis Regulated by GnIH in the Goldfish, Carassius auratus1. Biology of Reproduction, 2013, 88, 89.	2.7	39
27	A chromosomeâ€level genome assembly of the giant grouper (<i>Epinephelus lanceolatus</i>) provides insights into its innate immunity and rapid growth. Molecular Ecology Resources, 2019, 19, 1322-1332.	4.8	39
28	Comparative Transcriptomic Study of Muscle Provides New Insights into the Growth Superiority of a Novel Grouper Hybrid. PLoS ONE, 2016, 11, e0168802.	2.5	38
29	Expression profiles of dmrts and foxls during gonadal development and sex reversal induced by 171±-methyltestosterone in the orange-spotted grouper. General and Comparative Endocrinology, 2019, 274, 26-36.	1.8	37
30	Genetic Diversity and Differentiation of the Orange-Spotted Grouper (Epinephelus coioides) Between and Within Cultured Stocks and Wild Populations Inferred from Microsatellite DNA Analysis. International Journal of Molecular Sciences, 2011, 12, 4378-4394.	4.1	34
31	Molecular cloning, characterization and expression profiles of three estrogen receptors in protogynous hermaphroditic orange-spotted grouper (Epinephelus coioides). General and Comparative Endocrinology, 2011, 172, 371-381.	1.8	33
32	Intracellular TLR22 acts as an inflammation equalizer via suppression of NF-κB and selective activation of MAPK pathway in fish. Fish and Shellfish Immunology, 2018, 72, 646-657.	3.6	32
33	Discovery of four estrogen receptors and their expression profiles during testis recrudescence in male Spinibarbus denticulatus. General and Comparative Endocrinology, 2008, 156, 265-276.	1.8	31
34	Goldfish neurokinin B: Cloning, tissue distribution, and potential role in regulating reproduction. General and Comparative Endocrinology, 2015, 221, 267-277.	1.8	31
35	Genome-Wide Mapping of Growth-Related Quantitative Trait Loci in Orange-Spotted Grouper (Epinephelus coioides) Using Double Digest Restriction-Site Associated DNA Sequencing (ddRADseq). International Journal of Molecular Sciences, 2016, 17, 501.	4.1	31
36	Screening and characterization of sex-specific markers developed by a simple NGS method in mandarin fish (Siniperca chuatsi). Aquaculture, 2020, 527, 735495.	3.5	31

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37	Day-night and reproductive cycle profiles of melatonin receptor, kiss , and gnrh expression in orange-spotted grouper (Epinephelus coioides). Molecular Reproduction and Development, 2013, 80, 535-548.	2.0	30
38	The complete mitochondrial genome of the <i>Trachinotus ovatus</i> (Teleostei, Carangidae). Mitochondrial DNA, 2015, 26, 644-646.	0.6	30
39	Identification and functional characterization of Toll-like receptor 13 from orange-spotted grouper (Epinephelus coioides). Fish and Shellfish Immunology, 2018, 74, 309-317.	3.6	30
40	Gonadal development, aromatase activity and P450 aromatase gene expression during sex inversion of protogynous red-spotted grouper Epinephelus akaara (Temminck and Schlegel) after implantation of the aromatase inhibitor, fadrozole. Aquaculture Research, 2006, 37, 484-491.	1.8	29
41	Female-to-male sex reversal in orange-spotted grouper (<i>Epinephelus coioides</i>) caused by overexpressing of Amh in vivoâ€. Biology of Reproduction, 2018, 99, 1205-1215.	2.7	29
42	Vibrio parahaemolyticus flagellin induces cytokines expression via toll-like receptor 5 pathway in orange-spotted grouper, Epinephelus coioides. Fish and Shellfish Immunology, 2019, 87, 573-581.	3.6	29
43	Two distinct interferon-Î ³ genes in Tetraodon nigroviridis : Functional analysis during Vibrio parahaemolyticus infection. Molecular Immunology, 2016, 70, 34-46.	2.2	28
44	Phoenixin participated in regulation of food intake and growth in spotted scat, Scatophagus argus. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2018, 226, 36-44.	1.6	25
45	Whole Genome Sequencing of the Giant Grouper (Epinephelus lanceolatus) and High-Throughput Screening of Putative Antimicrobial Peptide Genes. Marine Drugs, 2019, 17, 503.	4.6	25
46	Molecular cloning of the insulin-like growth factor 3 and difference in the expression of igf genes in orange-spotted grouper (Epinephelus coioides). Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2015, 186, 68-75.	1.6	23
47	Wnt4 in protogynous hermaphroditic orange-spotted grouper (Epinephelus coioides): Identification and expression. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2015, 183, 67-74.	1.6	23
48	Characterization of triploid hybrid groupers from interspecies hybridization (<i>Epinephelus) Tj ETQq0 0 0 rgBT /(</i>	Dverlock 1 1.8	.0 <u>Tf</u> 50 302
49	Neurokinin B signaling in hermaphroditic species, a study of the orange-spotted grouper (Epinephelus) Tj ETQq1	1 0.78431 1.8	4 rgBT /Ove
50	Probiotics Improve Eating Disorders in Mandarin Fish (Siniperca chuatsi) Induced by a Pellet Feed Diet via Stimulating Immunity and Regulating Gut Microbiota. Microorganisms, 2021, 9, 1288.	3.6	23
51	Transcriptome analysis of the spleen provides insight into the immunoregulation of Mastacembelus armatus under Aeromonas veronii infection. Fish and Shellfish Immunology, 2019, 88, 272-283.	3.6	22
52	Socially controlled maleâ€ŧoâ€ŧemale sex reversal in the protogynous orangeâ€spotted grouper, <scp><i>Epinephelus coioides</i></scp> . Journal of Fish Biology, 2019, 94, 414-421.	1.6	20
53	Tetraodon nigroviridis : A model of Vibrio parahaemolyticus infection. Fish and Shellfish Immunology, 2016, 56, 388-396.	3.6	18
54	Identification and characterization of a motilin-like peptide and its receptor in teleost. General and Comparative Endocrinology, 2013, 186, 85-93.	1.8	17

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55	Two IFNGR1 homologues in Tetraodon nigroviridis: Origin, expression analysis and ligand-binding preference. Developmental and Comparative Immunology, 2014, 44, 270-279.	2.3	17
56	MT-Feeding-Induced Impermanent Sex Reversal in the Orange-Spotted Grouper during Sex Differentiation. International Journal of Molecular Sciences, 2018, 19, 2828.	4.1	17
57	Expression profiles of gonadotropins and their receptors during 17αâ€methyltestosterone implantationâ€induced sex change in the orangeâ€spotted grouper (<i>Epinephelus coioides</i>). Molecular Reproduction and Development, 2011, 78, 376-390.	2.0	15
58	Polymorphisms of Leptin-b Gene Associated with Growth Traits in Orange-Spotted Grouper (Epinephelus coioides). International Journal of Molecular Sciences, 2014, 15, 11996-12006.	4.1	15
59	Two Distinct Interferon-γ in the Orange-Spotted Grouper (Epinephelus coioides): Molecular Cloning, Functional Characterization, and Regulation in Toll-Like Receptor Pathway by Induction of miR-146a. Frontiers in Endocrinology, 2018, 9, 41.	3.5	15
60	An estradiol-17β/miRNA-26a/cyp19a1a regulatory feedback loop in the protogynous hermaphroditic fish, Epinephelus coioides. Molecular and Cellular Endocrinology, 2020, 504, 110689.	3.2	15
61	Beta-Hydroxysteroid Dehydrogenase Genes in Orange-Spotted Grouper (Epinephelus coioides): Genome-Wide Identification and Expression Analysis During Sex Reversal. Frontiers in Genetics, 2020, 11, 161.	2.3	15
62	Production of neoâ€male mandarin fish Siniperca chuatsi by masculinization with orally administered 17α-methyltestosterone. Aquaculture, 2021, 530, 735904.	3.5	15
63	Promotion of pellet-feed feeding in mandarin fish (Siniperca chuatsi) by Bdellovibrio bacteriovorus is influenced by immune and intestinal flora. Aquaculture, 2021, 542, 736864.	3.5	15
64	MicroRNA-182-3p negatively regulates cytokines expression by targeting TLR5M in orange-spotted grouper, Epinephelus coioides. Fish and Shellfish Immunology, 2019, 93, 589-596.	3.6	13
65	Two alternatively spliced GPR39 transcripts in seabream: molecular cloning, genomic organization, and regulation of gene expression by metabolic signals. Journal of Endocrinology, 2008, 199, 457-470.	2.6	12
66	The complete mitochondrial genome of theHemibagrus wyckioides(Siluriformes, Bagridae). Mitochondrial DNA, 2016, 27, 766-768.	0.6	12
67	The complete mitochondrial genome of theHemibarbus medius(Cypriniformes, Cyprinidae). Mitochondrial DNA, 2016, 27, 1070-1072.	0.6	12
68	Copy Number Variations in Tilapia Genomes. Marine Biotechnology, 2017, 19, 11-21.	2.4	12
69	Leptin Stimulates Prolactin mRNA Expression in the Goldfish Pituitary through a Combination of the PI3K/Akt/mTOR, MKK3/6/p38MAPK and MEK1/2/ERK1/2 Signalling Pathways. International Journal of Molecular Sciences, 2017, 18, 2781.	4.1	12
70	Characterization, evolution, and expression analysis of TLR7 gene subfamily members in Mastacembelus armatus (Synbranchiformes: Mastacembelidae). Developmental and Comparative Immunology, 2019, 95, 77-88.	2.3	12
71	Single Nucleotide Polymorphisms in the Leptin-a Gene and Associations with Growth Traits in the Orange-Spotted Grouper (Epinephelus coioides). International Journal of Molecular Sciences, 2013, 14, 8625-8637.	4.1	11
72	The complete mitochondrial genome of the hybrid grouper (<i>Cromileptes altivelis♀</i> ×) Tj ETQq0 0 0 rg	BT /Overlo 0.4	ck 10 Tf 50 6 11

Resources, 2017, 2, 171-172.

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			Chinana
73	Identification and functional characterization of two Secretogranin II genes in orange-spotted grouper (Epinephelus coioides). General and Comparative Endocrinology, 2018, 261, 115-126.	1.8	11
74	Molecular identification of the Dyn/Kor system and its potential role in the reproductive axis of goldfish. General and Comparative Endocrinology, 2018, 257, 29-37.	1.8	11
75	Transcriptomic Analysis Revealed the Regulatory Mechanisms of Oocyte Maturation and Hydration in Orange-Spotted Grouper (Epinephelus coioides). Marine Biotechnology, 2019, 21, 537-549.	2.4	11
76	NKB/NK3 system negatively regulates the reproductive axis in sexually immature goldfish (Carassius) Tj ETQqO O (0 rgBT /Ov F.8	erlock 10 Tf
77	Natural sex change in mature protogynous orangeâ€spotted grouper (<i>Epinephelus coioides</i>):	1.6	11

78	The Administration of Cortisol Induces Female-to-Male Sex Change in the Protogynous Orange-Spotted Grouper, Epinephelus coioides. Frontiers in Endocrinology, 2020, 11, 12.	3.5	11
79	Efficient RNA Virus Targeting via CRISPR/CasRx in Fish. Journal of Virology, 2021, 95, e0046121.	3.4	11
80	Characterization of dmrts and their potential role in gonadal development of mandarin fish (Siniperca chuatsi). Aquaculture Reports, 2021, 21, 100802.	1.7	11
81	An SNP-Based Genetic Map and QTL Mapping for Growth Traits in the Red-Spotted Grouper (Epinephelus) Tj ETQc	1 _{2.4} 1,0.784	•314 rgBT 10
82	New Insights Into the Role of Follicle-Stimulating Hormone in Sex Differentiation of the Protogynous Orange-Spotted Grouper, Epinephelus coioides. Frontiers in Endocrinology, 2019, 10, 304.	3.5	10
83	Comparative transcriptome analysis of diploid and triploid hybrid groupers (Epinephelus coioides♀â€ ⁻ ×â€ ⁻ E.) 2019, 111, 251-259.	Tj ETQq1 2.9	1 0.78431 10
84	Induction of oocyte maturation and changes in the biochemical composition, physiology and molecular biology of oocytes during maturation and hydration in the orange-spotted grouper (Epinephelus coioides). Aquaculture, 2020, 522, 735115.	3.5	9
85	Identification of potential sex-related genes in Siniperca chuatsi. Journal of Oceanology and Limnology, 2021, 39, 1500-1512.	1.3	9
86	The complete mitochondrial genome of the <i>Rhabdosargus sarba</i> (Perciformes: Sparidae). Mitochondrial DNA, 2016, 27, 1-2.	0.6	8
87	Cloning, expression and functional characterization of a novel luteinizing hormone receptor in the orange-spotted grouper, Epinephelus coioides. General and Comparative Endocrinology, 2018, 267, 90-97.	1.8	8
88	Estradiol-17β regulates the expression of insulin-like growth factors 1 and 2 via estradiol receptors in spotted scat (Scatophagus argus). Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2019, 237, 110328.	1.6	8
89	The complete mitochondrial genome of the hybrid grouper <i>Epinephelus coioides</i> ♀ <i> × Epinephelus akaara</i> â™, with phylogenetic consideration. Mitocho Resources, 2017, 2, 31-32.	n o kial DN/	\₿art B:
90	23S rRNA from Vibrio parahaemolyticus regulates the innate immune response via recognition by TLR13 in orange-spotted grouper (Epinephelus coioides). Developmental and Comparative Immunology, 2021, 114, 103837.	2.3	7

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91	Physical interactions facilitate sex change in the protogynous orangeâ€spotted grouper, Epinephelus coioides. Journal of Fish Biology, 2021, 98, 1308-1320.	1.6	7
92	A simple PCR-based genetic sex identification method in the blotched snakehead (Channa maculata) developed by high-throughput sequencing. Aquaculture, 2021, 538, 736579.	3.5	7
93	Comparative Metabolomics and Proteomics Reveal Vibrio parahaemolyticus Targets Hypoxia-Related Signaling Pathways of Takifugu obscurus. Frontiers in Immunology, 2021, 12, 825358.	4.8	7
94	Development and gene expression analysis of gonad during 17α-methyltestosterone-induced sex reversal in mandarin fish (Siniperca chuatsi). Aquaculture Reports, 2022, 23, 101049.	1.7	7
95	The complete mitochondrial genome of the orange-spotted grouper Epinephelus coioides (Perciformes, Serranidae). Mitochondrial DNA, 2014, 27, 1-3.	0.6	6
96	Knockout of tac3 genes in zebrafish shows no impairment of reproduction. General and Comparative Endocrinology, 2021, 311, 113839.	1.8	6
97	The flagellin of Vibrio parahaemolyticus induces the inflammatory response of Tetraodon nigroviridis through TLR5M. Fish and Shellfish Immunology, 2021, 120, 102-110.	3.6	6
98	Comparison of Gonadal Development in Diploid and Triploid Hybrid Groupers, <i>Epinephelus coioides</i> ♀ × <i>Epinephelus lanceolatus</i> â™,. Journal of the World Aquaculture Society, 2018, 49, 328-337.	2.4	5
99	Molecular characterization and functional analysis of IKK $\hat{I}\pm$ in orange-spotted grouper (Epinephelus) Tj ETQq1 1	0.784314	rgBT /Overloc
100	The complete mitochondrial genome of the <i>Platax teira</i> (Osteichthyes: Ephippidae). Mitochondrial DNA, 2016, 27, 796-797.	0.6	4
101	Retinoic acid and androgen influence germ cells development and meiotic initiation in juvenile orange-spotted grouper, Epinephelus coioides. General and Comparative Endocrinology, 2020, 289, 113379.	1.8	4
102	MicroRNA-29b modulates the innate immune response by suppressing IFNγs production in orange-spotted grouper (Epinephelus coioides). Fish and Shellfish Immunology, 2020, 104, 537-544.	3.6	4
103	Chromosome-Level Genome Assembly and Transcriptome Comparison Analysis of Cephalopholis sonnerati and Its Related Grouper Species. Biology, 2022, 11, 1053.	2.8	4
104	The complete mitochondrial genome of the hybrid grouper <i>Epinehelus_moara ♀</i> × <i>Epinephelus_lanceolatus â™,</i> with phylogenetic consideration. Mitochondrial DNA Part B: Resources, 2016, 1, 584-585.	0.4	3
105	Microsatellite analysis of the genetic relationships between wild and cultivated giant grouper in the South China Sea. Journal of Genetics, 2016, 95, 369-376.	0.7	3
106	Formation of diploid and triploid hybrid groupers (hybridization of Epinephelus coioides ♀ × Epine	ephelus) Tj 2 .7	ETQq0 0 0 rg
107	The complete mitochondrial genome of thePampus nozawae(Perciformes: Stromateidae). Mitochondrial DNA, 2016, 27, 988-989.	0.6	3

108 The complete mitochondrial genome of the hybrid grouper Epinephelus coioides♀ × Epinephelus 0.7 3 lanceolatusâ™,. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 4181-4182.

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109	The next-generation sequencing reveals the complete mitochondrial genome of <i>Alosa sapidissima</i> (Perciformes: Clupeidae) with phylogenetic consideration. Mitochondrial DNA Part B: Resources, 2017, 2, 304-306.	0.4	3
110	A PCR-based genetic sex identification method in spotted mandarin fish (Siniperca scherzeri) and big eye mandarin fish (Siniperca kneri). Aquaculture Reports, 2020, 18, 100552.	1.7	3
111	Mutation of <i>spexin2</i> promotes feeding, somatic growth, adiposity, and insulin resistance in zebrafish. American Journal of Physiology - Regulatory Integrative and Comparative Physiology, 2022, 322, R454-R465.	1.8	3
112	The complete mitochondrial genome of theEpinephelus lanceolatus(Perciformes: Serranidae). Mitochondrial DNA, 2014, 27, 1-2.	0.6	2
113	The complete mitochondrial genome of theEpinephelus akaara(Perciformes: Serranidae). Mitochondrial DNA, 2014, 27, 1-2.	0.6	2
114	The complete mitochondrial genome of the Epinephelus fuscoguttatus (Perciformes: Serranidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 4110-4111.	0.7	2
115	Molecular cloning, expression patterns and functional characterization of Gpr3 in the orange-spotted grouper (Epinephelus coioides). Aquaculture Reports, 2022, 23, 101050.	1.7	2
116	Vibrio parahaemolyticus flagellin F (FlaF) induces the inflammatory response of the Tetraodon nigroviridis through the TLR5M. Aquaculture, 2022, 555, 738140.	3.5	2
117	Estrogen receptor-related receptors in mandarin fish (Siniperca chuatsi): Molecular cloning, characterization, and estrogen responsiveness. Aquaculture Reports, 2022, 24, 101137.	1.7	2
118	The complete mitochondrial genome of theEpinephelus moara(Osteichthyes: Ephippidae). Mitochondrial DNA, 2014, 27, 1-2.	0.6	1
119	The complete mitochondrial genome of Epinephelus awoara (Perciformes: Epinephelus) with phylogenetic consideration. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 4286-4287.	0.7	1
120	The complete mitochondrial genome of the Epinephelus corallicola (Perciformes: Serranidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 3971-3972.	0.7	1
121	Molecular cloning and characterization of estrogen and androgen receptors in Mandarin fish, Siniperca chuatsi. Aquaculture Reports, 2021, 21, 100834.	1.7	1
122	The complete mitochondrial genome of the <i>Drepane punctata</i> (Perciformes: Drepanidae). Mitochondrial DNA, 2016, 27, 1-2.	0.6	0
123	The complete mitochondrial genome of the <i>Siganus canaliculatus</i> (Perciformes: Siganidae). Mitochondrial DNA, 2016, 27, 1111-1112.	0.6	0
124	Cloning, pattern of gonadal soma-derived factor mRNA in the orange-spotted grouper, Epinephelus coioides. Aquaculture Reports, 2021, 20, 100754.	1.7	0