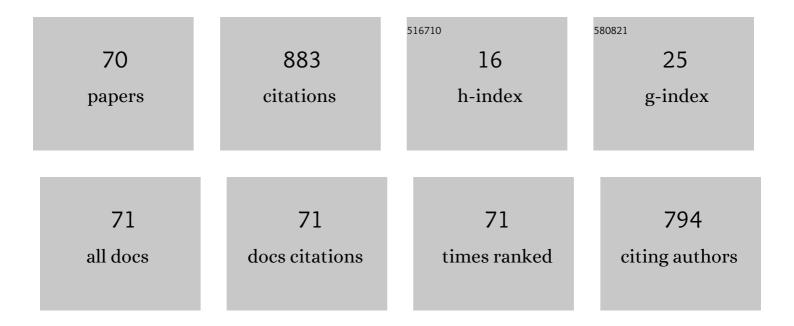
## Xinxin Du

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptome profiling based on protein–protein interaction networks provides a core set of genes for understanding blood immune response mechanisms against Edwardsiella tarda infection in Japanese flounder ( Paralichthys olivaceus ). Developmental and Comparative Immunology, 2018, 78, 100-113.	2.3	56
2	Transcriptome profiling provides gene resources for understanding gill immune responses in Japanese flounder (Paralichthys olivaceus) challenged with Edwardsiella tarda. Fish and Shellfish Immunology, 2018, 72, 593-603.	3.6	55
3	Comparative transcriptome analysis of ovary and testis reveals potential sex-related genes and pathways in spotted knifejaw Oplegnathus punctatus. Gene, 2017, 637, 203-210.	2.2	51
4	A chromosomeâ€level genome of black rockfish, <i>Sebastes schlegelii</i> , provides insights into the evolution of live birth. Molecular Ecology Resources, 2019, 19, 1309-1321.	4.8	44
5	Identification and characterization of a nanog homolog in Japanese flounder (Paralichthys) Tj ETQq1 1 0.78431	4 rgBT /Ov 2.2	erlock 10 Tf 5
6	Sequencing-based network analysis provides a core set of gene resource for understanding kidney immune response against Edwardsiella tarda infection in Japanese flounder. Fish and Shellfish Immunology, 2017, 67, 643-654.	3.6	38
7	Transcriptome Profiling Insights the Feature of Sex Reversal Induced by High Temperature in Tongue Sole Cynoglossus semilaevis. Frontiers in Genetics, 2019, 10, 522.	2.3	34
8	Transcriptome analysis of the gonads of olive flounder (Paralichthys olivaceus). Fish Physiology and Biochemistry, 2016, 42, 1581-1594.	2.3	33
9	Detection of Alternative Splice and Gene Duplication by RNA Sequencing in Japanese Flounder, <i>Paralichthys olivaceus</i> . G3: Genes, Genomes, Genetics, 2014, 4, 2419-2424.	1.8	27
10	Identification and characterization of a Sox2 homolog in the Japanese flounder Paralichthys olivaceus. Gene, 2014, 544, 165-176.	2.2	21
11	Isolation and characterization of 64 novel microsatellite markers from a fosmid library of female halfâ€smooth tongue sole ( <i>Cynoglossus semilaevis</i> ). Molecular Ecology Resources, 2008, 8, 1303-1306.	4.8	20
12	Fosmid Library Construction and Initial Analysis of End Sequences in Female Half-smooth Tongue Sole (Cynoglossus semilaevis). Marine Biotechnology, 2009, 11, 236-242.	2.4	20
13	Sexually dimorphic expression in developing and adult gonads shows an important role of gonadal soma-derived factor during sex differentiation in olive flounder ( Paralichthys olivaceus ). Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2017, 210, 1-8.	1.6	19
14	Piwil1 gene is regulated by hypothalamic-pituitary-gonadal axis in turbot ( Scophthalmus maximus ): A different effect in ovaries and testes. Gene, 2018, 658, 86-95.	2.2	19
15	Spotted knifejaw (Oplegnathus punctatus) MyD88: Intracellular localization, signal transduction function and immune responses to bacterial infection. Fish and Shellfish Immunology, 2019, 89, 719-726.	3.6	19
16	Molecular Cloning, Promoter Analysis and Expression Profiles of the sox3 Gene in Japanese Flounder, Paralichthys olivaceus. International Journal of Molecular Sciences, 2015, 16, 27931-27944.	4.1	18
17	Edwardsiella tarda-induced miR-7a functions as a suppressor in PI3K/AKT/GSK3β signaling pathway by targeting insulin receptor substrate-2 (IRS2a and IRS2b) in Paralichthys olivaceus. Fish and Shellfish Immunology, 2019, 89, 477-485.	3.6	17
18	Identification, evolution and expression analyses of mapk gene family in Japanese flounder (Paralichthys olivaceus) provide insight into its divergent functions on biotic and abiotic stresses response. Aquatic Toxicology, 2021, 241, 106005.	4.0	17

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19	Molecular characterization and functional analysis of the GATA4 in tongue sole (Cynoglossus) Tj ETQq1 1 0.7843		
	193, 1-8.	1.6	16
20	Identification and expression of piwil2 in turbot Scophthalmus maximus , with implications of the involvement in embryonic and gonadal development. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2017, 208-209, 84-93.	1.6	16
21	Genome-wide identification and transcriptome-based expression analysis of sox gene family in the Japanese flounder Paralichthys olivaceus. Journal of Oceanology and Limnology, 2018, 36, 1731-1745.	1.3	16
22	A novel C-type lectin from spotted knifejaw, Oplegnathus punctatus possesses antibacterial and anti-inflammatory activity. Fish and Shellfish Immunology, 2019, 92, 11-20.	3.6	16
23	Long-term exposure to environmental relevant triclosan induces reproductive toxicity on adult zebrafish and its potential mechanism. Science of the Total Environment, 2022, 826, 154026.	8.0	16
24	Molecular characterization and expression profiles of GATA6 in tongue sole (Cynoglossus) Tj ETQq0 0 0 rgBT /Ove 198, 19-26.	erlock 10 <sup>-</sup> 1.6	Tf 50 547 Td 15
25	Roles of Two Sox9 Genes during Gonadal Development in Japanese Flounder: Sex Differentiation, Spermatogenesis and Gonadal Function Maintenance. International Journal of Molecular Sciences, 2018, 19, 512.	4.1	14
26	Evolutionary Conservation of pou5f3 Genomic Organization and Its Dynamic Distribution during Embryogenesis and in Adult Gonads in Japanese Flounder Paralichthys olivaceus. International Journal of Molecular Sciences, 2017, 18, 231.	4.1	13
27	Kunitz-type serine protease inhibitor is a novel participator in anti-bacterial and anti-inflammatory responses in Japanese flounder ( Paralichthys olivaceus ). Fish and Shellfish Immunology, 2018, 80, 22-30.	3.6	12
28	Genetic variation assessed with microsatellites in mass selection lines of the Pacific oyster (Crassostrea gigas) in China. Journal of Ocean University of China, 2016, 15, 1039-1045.	1.2	11
29	Roles of piwil1 gene in gonad development and gametogenesis in Japanese flounder, Paralichthys olivaceus. Gene, 2019, 701, 104-112.	2.2	11
30	Expression and functional analysis of receptor-interacting serine/threonine kinase 2 (RIP2) in Japanese flounder (Paralichthys olivaceus). Fish and Shellfish Immunology, 2018, 75, 327-335.	3.6	10
31	Identification, molecular characterization and gene expression analysis of sox1a and sox1b genes in Japanese flounder, Paralichthys olivaceus. Gene, 2015, 574, 225-234.	2.2	9
32	Rapid evolution of piRNA pathway and its transposon targets in Japanese flounder (Paralichthys) Tj ETQq0 0 0 rgB 100609.	T /Overloo 1.0	ck 10 Tf 50 2 9
33	Molecular Mechanism Based on Histopathology, Antioxidant System and Transcriptomic Profiles in Heat Stress Response in the Gills of Japanese Flounder. International Journal of Molecular Sciences, 2022, 23, 3286.	4.1	9
34	Sex determination mechanisms in fish. Journal of Ocean University of China, 2009, 8, 155-160.	1.2	8
35	Identification of two major histocompatibility (MH) class II A genes and their association to Vibrio anguillarum infection in half-smooth tongue sole (Cynoglossus semilaevis). Journal of Ocean University of China, 2012, 11, 32-44.	1.2	8
36	Sequences analyses and expression profiles in tissues and embryos of Japanese flounder (Paralichthys) Tj ETQqO C	0 rgBT /C	Vygrlock 10 T

#	Article	IF	CITATIONS
37	Roles of two <i>cyp11</i> genes in sex hormone biosynthesis in Japanese flounder ( <i>Paralichthys) Tj ETQq1</i>	1 0.784314 2.0	rg&T /Overloo
38	Weighted Correlation Network Analysis (WGCNA) of Japanese Flounder (Paralichthys olivaceus) Embryo Transcriptome Provides Crucial Gene Sets for Understanding Haploid Syndrome and Rescue by Diploidization. Journal of Ocean University of China, 2018, 17, 1441-1450.	1.2	7
39	Comparative studies on duplicated tdrd7 paralogs in teleosts: Molecular evolution caused neo-functionalization. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2019, 30, 347-357.	1.0	7
40	Two cyp17 genes perform different functions in the sex hormone biosynthesis and gonadal differentiation in Japanese flounder (Paralichthys olivaceus). Gene, 2019, 702, 17-26.	2.2	7
41	Isolation and characterization of 21 novel microsatellite markers from spotted halibut (Verasper) Tj ETQq1 1 0.	784314 rgB 1.8	T /Overlock ]
42	miR-1388 regulates the expression of nectin2l in Paralichthys olivaceus. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2018, 28, 9-16.	1.0	6
43	Evolutionary significance and regulated expression of Tdrd family genes in gynogenetic Japanese flounder (Paralichthys olivaceus). Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2019, 31, 100593.	1.0	6
44	Genomeâ€wide identification of nonvisual opsin family reveals amplification of RPEâ€retinal G protein receptor gene ( <i>RGR</i> ) and offers novel insights into functions of <i>RGR</i> (s) in <i>Paralichthys olivaceus</i> (Paralichthyidae, Teleostei). Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2020, 334, 25-36.	1.3	6
45	Comparative Evolution of Duplicated <i>Ddx3</i> Genes in Teleosts: Insights from Japanese Flounder, <i>Paralichthys olivaceus</i> . G3: Genes, Genomes, Genetics, 2015, 5, 1765-1773.	1.8	5
46	The evolution and possible role of two Sox8 genes during sex differentiation in Japanese flounder () Tj ETQq0 0	0 rgBT /Ove	erlock 10 Tf 5
47	Molecular characterization, expression and functional analysis of cystatin C in Japanese flounder (Paralichthys olivaceus). Fish and Shellfish Immunology, 2019, 86, 695-701.	3.6	5
48	Evolution history of duplicated <i>smad3</i> genes in teleost: insights from Japanese flounder, <i>Paralichthys olivaceus</i> . PeerJ, 2016, 4, e2500.	2.0	5
49	Molecular cloning and expression of PoIR2, a novel gene involved in immune response in Japanese flounder (Paralichthys olivaceus). Journal of Ocean University of China, 2010, 9, 59-64.	1.2	4
50	Identification and Characterization of a PRDM14 Homolog in Japanese Flounder (Paralichthys) Tj ETQq0 0 0 rgI	3T /Overlock 4.1	10 Tf 50 222
51	Locus Number Estimation of MHC Class II B in Stone Flounder and Japanese Flounder. International Journal of Molecular Sciences, 2015, 16, 6000-6017.	4.1	4
52	Characterization and functional analysis of the Paralichthys olivaceus prdm1 gene promoter. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2017, 212, 32-40.	1.6	3
53	Klf4 could be activated by Pou5f3/Stat3 to induce the expression of Nanog in Japanese flounder. Gene, 2019, 687, 56-63.	2.2	3

54	Establishment and Characterization of Four Long-Term Cultures of Neural Stem/Progenitor Cell Lines from the Japanese Flounder Paralichthys olivaceus. Journal of Ocean University of China, 2020, 19, 1153-1162.	1	.2	3
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#	Article	IF	CITATIONS
55	Hypothalamic-Pituitary-Gonadal (HPG) Axis and Transcriptional Regulatory Elements Regulate piwil2 Gene Expression During Gametogenesis and Gonadal Development in Japanese Flounder (Paralichthys) Tj ETQq1 I	1 <b>0.2</b> 8431	4 <b>s</b> gBT /Over
56	Genome-Wide Identification of heat shock protein 10/60 Genes in Japanese Flounder (Paralichthys) Tj ETQq0 0 0 c China, 2021, 20, 886-896.	rgBT /Over 1.2	rlock 10 Tf 5 3
57	Cloning and stage-specific expression of CK-M1 gene during metamorphosis of Japanese flounder, Paralichthys olivaceus. Chinese Journal of Oceanology and Limnology, 2010, 28, 558-564.	0.7	2
58	Sequence polymorphism of two major histocompatibility (MH) class II B genes and their association with Vibrio anguillarum infection in half-smooth tongue sole (Cynoglossus semilaevis). Chinese Journal of Oceanology and Limnology, 2011, 29, 1275-1286.	0.7	2
59	Molecular cloning, expression profiles and promoter analysis of insulin-like growth factor binding protein-4 (IGFBP-4) in Japanese flounder (Paralichthys olivaceus). Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2014, 175, 41-52.	1.6	2
60	Novel Functions of Interleukin-1 Beta in Spotted Halibut (Verasper variegatus) at Its Embryonic and Metamorphic Stages. Journal of Ocean University of China, 2018, 17, 1197-1208.	1.2	2
61	Discovery and functional characterization of microRNAs and their potential roles for gonadal development in spotted knifejaw, Oplegnathus punctatus. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2018, 28, 1-8.	1.0	2
62	Screening of eye-position related genes with DD-RT-PCR and RDA in the hybrids between Japanese flounder Paralichthys olivaceus and stone flounder Kareius bicoloratus. Chinese Journal of Oceanology and Limnology, 2009, 27, 92-99.	0.7	1
63	Molecular cloning, expression pattern, and 3D structural prediction of the cold inducible RNA-binding protein (CIRP) in Japanese flounder (Paralichthys olivaceus). Journal of Ocean University of China, 2015, 14, 161-170.	1.2	1
64	Functional characterization of the Japanese flounder (Paralichthys olivaceus) Sox2 gene promoter. Fish Physiology and Biochemistry, 2016, 42, 1275-1285.	2.3	1
65	Roles of GATA6 during Gonadal Development in Japanese Flounder: Gonadogenesis, Regulation of Gender-Related Genes, Estrogen Formation and Gonadal Function Maintenance. International Journal of Molecular Sciences, 2017, 18, 160.	4.1	1
66	Analysis of new microsatellite markers developed from reported sequences of Japanese flounder Paralichthys olivaceus. Journal of Ocean University of China, 2010, 9, 365-370.	1.2	0
67	Isolation and characterization of eighteen polymorphic microsatellite markers from the frog flounder, Pleuronichthys cornutus. Conservation Genetics Resources, 2013, 5, 335-337.	0.8	0
68	Isolation and characterization of twenty novel microsatellite markers of Pointhead flounder (Cleisthenes herzensteini). Conservation Genetics Resources, 2013, 5, 137-139.	0.8	0
69	Tumor necrosis factor-alpha (TNF-α) in spotted halibut Verasper variegatus at the embryonic and metamorphic stages. Journal of Oceanology and Limnology, 2020, 38, 454-466.	1.3	0
70	GATA4 Is a Transcriptional Regulator of SOX2 in Japanese Flounder (Paralichthys olivaceus). Journal of Ocean University of China, 2022, 21, 163-170.	1.2	0