Ji-Fang Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3410109/publications.pdf

Version: 2024-02-01

	687363	642732
583	13	23 g-index
citations	h-index	g-index
21	21	474
31	31	474
docs citations	times ranked	citing authors
	citations 31	583 13 citations h-index 31 31

#	Article	IF	CITATIONS
1	DNA methylation in promoter region of immune related genes STAT3 and VEGFA and biochemical parameters change in muscle of Japanese flounder under acute hypoxia. Developmental and Comparative Immunology, 2022, 129, 104295.	2.3	8
2	Cyclooxygenases of ovoviviparous black rockfish (Sebastes schlegelii): Cloning, tissue distribution and potential role in mating and parturition. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2022, 257, 110677.	1.6	11
3	Molecular characterization and expression patterns of glucocorticoid receptors in the viviparous black rockfish Sebastes schlegelii. General and Comparative Endocrinology, 2022, 316, 113947.	1.8	2
4	Identification and characterization of mkk genes and their expression profiles in rainbow trout (Oncorhynchus mykiss) symptomatically or asymptomatically infected with Vibrio anguillarum. Fish and Shellfish Immunology, 2022, 121, 1-11.	3.6	4
5	Transcriptional Signatures of Immune, Neural, and Endocrine Functions in the Brain and Kidney of Rainbow Trout (Oncorhynchus mykiss) in Response to Aeromonas salmonicida Infection. International Journal of Molecular Sciences, 2022, 23, 1340.	4.1	6
6	Characterization of CYP11A1 and its potential role in sex asynchronous gonadal development of viviparous black rockfish Sebastes schlegelii (Sebastidae). General and Comparative Endocrinology, 2021, 302, 113689.	1.8	15
7	Transcriptional Profiles of Genes Related to Stress and Immune Response in Rainbow Trout (Oncorhynchus mykiss) Symptomatically or Asymptomatically Infected With Vibrio anguillarum. Frontiers in Immunology, 2021, 12, 639489.	4.8	9
8	Identification and characterization of caspases genes in rainbow trout (Oncorhynchus mykiss) and their expression profiles after Aeromonas salmonicida and Vibrio anguillarum infection. Developmental and Comparative Immunology, 2021, 118, 103987.	2.3	19
9	Acute hypoxia effects on Keap1/Nrf2 (Mafs)-GST pathway related oxidative metabolism in muscle of Japanese flounder (Paralichthys olivaceus). Science of the Total Environment, 2021, 795, 148646.	8.0	19
10	HSP90 and HSP70 Families in Lateolabrax maculatus: Genome-Wide Identification, Molecular Characterization, and Expression Profiles in Response to Various Environmental Stressors. Frontiers in Physiology, 2021, 12, 784803.	2.8	10
11	Environmental hypoxia causes growth retardation, osteoclast differentiation and calcium dyshomeostasis in juvenile rainbow trout (Oncorhynchus mykiss). Science of the Total Environment, 2020, 705, 135272.	8.0	32
12	Identification and Characterization of IncRNAs Related to the Muscle Growth and Development of Japanese Flounder (Paralichthys olivaceus). Frontiers in Genetics, 2020, 11, 1034.	2.3	11
13	GHRH-SST-GH-IGF axis regulates crosstalk between growth and immunity in rainbow trout (Oncorhynchus mykiss) infected with Vibrio anguillarum. Fish and Shellfish Immunology, 2020, 106, 887-897.	3.6	9
14	First High-Density Linkage Map and QTL Fine Mapping for Growth-Related Traits of Spotted Sea bass (Lateolabrax maculatus). Marine Biotechnology, 2020, 22, 526-538.	2.4	18
15	Alternative splicing (AS) mechanism plays important roles in response to different salinity environments in spotted sea bass. International Journal of Biological Macromolecules, 2020, 155, 50-60.	7.5	18
16	Genome-wide identification and characterization of toll-like receptor genes in spotted sea bass (Lateolabrax maculatus) and their involvement in the host immune response to Vibrio harveyi infection. Fish and Shellfish Immunology, 2019, 92, 782-791.	3.6	34
17	Effects of long-term crowding stress on neuro-endocrine-immune network of rainbow trout (Oncorhynchus mykiss). Fish and Shellfish Immunology, 2019, 95, 180-189.	3.6	9
18	Analysis of apolipoprotein multigene family in spotted sea bass (Lateolabrax maculatus) and their expression profiles in response to Vibrio harveyi infection. Fish and Shellfish Immunology, 2019, 92, 111-118.	3.6	22

#	Article	IF	CITATIONS
19	The impact of acute thermal stress on the metabolome of the black rockfish (Sebastes schlegelii). PLoS ONE, 2019, 14, e0217133.	2.5	39
20	14-3-3 gene family in spotted sea bass (Lateolabrax maculatus): Genome-wide identification, phylogenetic analysis and expression profiles after salinity stress. Comparative Biochemistry and Physiology Part A, Molecular & Entry integrative Physiology, 2019, 235, 1-11.	1.8	10
21	Characterization of Full-Length Transcriptome Sequences and Splice Variants of Lateolabrax maculatus by Single-Molecule Long-Read Sequencing and Their Involvement in Salinity Regulation. Frontiers in Genetics, 2019, 10, 1126.	2.3	29
22	Stocking density affects the growth performance and metabolism of Amur sturgeon by regulating expression of genes in the GH/IGF axis. Journal of Oceanology and Limnology, 2018, 36, 956-972.	1.3	12
23	Deep Transcriptomic Analysis of Black Rockfish (Sebastes schlegelii) Provides New Insights on Responses to Acute Temperature Stress. Scientific Reports, 2018, 8, 9113.	3.3	53
24	Liver transcriptome analysis reveals extensive transcriptional plasticity during acclimation to low salinity in Cynoglossus semilaevis. BMC Genomics, 2018, 19, 464.	2.8	48
25	Methylation Status of the Follistatin Gene at Different Development Stages of Japanese Flounder (Paralichthys olivaceus). Journal of Ocean University of China, 2018, 17, 1243-1252.	1.2	2
26	DNA methylation levels and expression patterns of Smyd1a and Smyd1b genes during Metamorphosis of the Japanese Flounder (Paralichthys olivaceus). Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2018, 223, 16-22.	1.6	12
27	Effects of stocking density on lipid deposition and expression of lipid-related genes in Amur sturgeon (Acipenser schrenckii). Fish Physiology and Biochemistry, 2017, 43, 1707-1720.	2.3	11
28	Low salinity affects cellularity, DNA methylation, and mRNA expression of igf1 in the liver of half smooth tongue sole (Cynoglossus semilaevis). Fish Physiology and Biochemistry, 2017, 43, 1587-1602.	2.3	30
29	Analysis of DNA methylation level by methylation-sensitive amplification polymorphism in half smooth tongue sole (Cynoglossus semilaevis) subjected to salinity stress. Journal of Ocean University of China, 2017, 16, 269-278.	1.2	14
30	Genetic polymorphisms and DNA methylation in exon 1 CpG-rich regions of PACAP gene and its effect on mRNA expression and growth traits in half smooth tongue sole (Cynoglossus semilaevis). Fish Physiology and Biochemistry, 2016, 42, 407-421.	2.3	11
31	DNA methylation level of cyp19a1a and Foxl2 gene related to their expression patterns and reproduction traits during ovary development stages of Japanese flounder (Paralichthys olivaceus). Gene, 2016, 575, 321-330.	2.2	56