Guangjun Wang

List of Publications by Year in descending order

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933447 888059 26 344 10 17 citations g-index h-index papers 27 27 27 283 docs citations times ranked citing authors all docs

| # | Article | IF | CITATIONS |
|----|---|------------------|---------------------|
| 1 | Impact of microecological agents on water environment restoration and microbial community structures of trench system in a Baiyangdian wetland ecosystem. Journal of Applied Microbiology, 2022, 132, 2450-2463. | 3.1 | 7 |
| 2 | MicroRNA-dependent regulation of targeted mRNAs for improved muscle texture in crisp grass carp fed with broad bean. Food Research International, 2022, 155, 111071. | 6.2 | 6 |
| 3 | Effectiveness of agricultural waste in the enhancement of biological denitrification of aquaculture wastewater. PeerJ, 2022, 10, e13339. | 2.0 | 7 |
| 4 | Effects of <scp>BBR</scp> on growth performance, serum and hepatic biochemistry parameters, hepatic morphology and gene expression levels related to glucose metabolism in largemouth bass, <i>Micropterus salmoides</i> . Aquaculture Research, 2022, 53, 3807-3817. | 1.8 | 9 |
| 5 | Both TGF- \hat{l}^21 and Smad4 regulate type I collagen expression in the muscle of grass carp, Ctenopharyngodon idella. Fish Physiology and Biochemistry, 2021, 47, 907-917. | 2.3 | 8 |
| 6 | Effect of the Aerobic Denitrifying Bacterium Pseudomonas furukawaii ZS1 on Microbiota Compositions in Grass Carp Culture Water. Water (Switzerland), 2021, 13, 1329. | 2.7 | 11 |
| 7 | Influence mechanism of filling ratio on solid-phase denitrification with polycaprolactone as biofilm carrier. Bioresource Technology, 2021, 337, 125401. | 9.6 | 25 |
| 8 | Quantitative phosphoproteomic analysis of soft and firm grass carp muscle. Food Chemistry, 2020, 303, 125367. | 8.2 | 33 |
| 9 | Comparative analysis of microbial community structure in the ponds with different aquaculture model and fish by high-throughput sequencing. Microbial Pathogenesis, 2020, 142, 104101. | 2.9 | 32 |
| 10 | Water Treatment Effect, Microbial Community Structure, and Metabolic Characteristics in a Field-Scale Aquaculture Wastewater Treatment System. Frontiers in Microbiology, 2020, 11, 930. | 3.5 | 18 |
| 11 | Proteomic and metabolomic basis for improved textural quality in crisp grass carp (Ctenopharyngodon idellus C.et V) fed with a natural dietary pro-oxidant. Food Chemistry, 2020, 325, 126906. | 8.2 | 53 |
| 12 | Identification and expression analysis of miRNA in hybrid snakehead by deep sequencing approach and their targets prediction. Genomics, 2019, 111, 1315-1324. | 2.9 | 2 |
| 13 | Epizootic ulcerative syndrome causes cutaneous dysbacteriosis in hybrid snakehead (<i>Channa) Tj ETQq1 1 0.7</i> | 84314 rgB 2.0 | BT /Overlock 1 |
| 14 | Artificial substrata increase pond farming density of grass carp (<i>Ctenopharyngodon idella</i>) by increasing the bacteria that participate in nitrogen and phosphorus cycles in pond water. PeerJ, 2019, 7, e7906. | 2.0 | 5 |
| 15 | Denitrification potential evaluation of a newly indigenous aerobic denitrifier isolated from largemouth bass Micropterus salmoides culture pond. Journal of Oceanology and Limnology, 2018, 36, 913-925. | 1.3 | 2 |
| 16 | Identification and comparative analysis of the miRNA expression profiles from four tissues of Micropterus salmoides using deep sequencing. Genomics, 2018, 110, 414-422. | 2.9 | 8 |
| 17 | Broad Bean (Vicia faba L.) Induces Intestinal Inflammation in Grass Carp (Ctenopharyngodon idellus C.) Tj ETQq1 in Microbiology, 2018, 9, 1913. | 1 0.78433 3.5 | 14 rgBT /Over 17 |
| 18 | The complete mitochondrial genome of <i>Carassius auratus var.</i> high back crucian carp (Cypriniformes: Cyprinidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2017, 28, 91-92. | 0.7 | 0 |

| # | Article | IF | CITATIONS |
|----|--|---------------------|------------|
| 19 | Genome-wide identification and characterization of conserved and novel microRNAs in grass carp () Tj ETQq $1\ 1\ C$ |).7 <u>84</u> 314 r | gBT/Overlo |
| 20 | Microbial succession in biofilms growing on artificial substratum in subtropical freshwater aquaculture ponds. FEMS Microbiology Letters, 2017, 364, fnx017. | 1.8 | 25 |
| 21 | Genome-wide identification of novel microRNAs from genome sequences using computational approach in the mudskipper (Boleophthalmus pectinirostris). Russian Journal of Bioorganic Chemistry, 2017, 43, 397-408. | 1.0 | 0 |
| 22 | The complete mitochondrial genome of <i>Shizothorax grahami</i> (Cypriniformes: Cyprinidae). Mitochondrial DNA Part B: Resources, 2016, 1, 775-776. | 0.4 | 2 |
| 23 | The complete mitochondrial genome of <i>Sinilabeo rendahli</i> (Cypriniformes: Cyprinidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 4603-4604. | 0.7 | 0 |
| 24 | An outbreak of granulomatous inflammation associated with Francisella noatunensis subsp. orientalis in farmed tilapia (Oreochromis niloticus \tilde{A} — O. aureus) in China. Chinese Journal of Oceanology and Limnology, 2016, 34, 460-466. | 0.7 | 19 |
| 25 | Gene Expression Profiling of Grass Carp (<i>Ctenopharyngodon idellus</i>) and Crisp Grass Carp. International Journal of Genomics, 2014, 2014, 1-15. | 1.6 | 26 |
| 26 | Laboratory evaluation of entomopathogenic fungi against the white grubs, <i>Holotrichia oblita </i> Anomala corpulenta (Coleoptera: Scarabaeidae) from the field of peanut, <i>Arachis hypogaea </i> . Biocontrol Science and Technology, 2011, 21, 593-603. | 1.3 | 13 |