

# Zhi-Jian Huang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7947709/publications.pdf>

Version: 2024-02-01

25  
papers

1,107  
citations

567281

15  
h-index

610901

24  
g-index

26  
all docs

26  
docs citations

26  
times ranked

993  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Efficient assembly of nanopore reads via highly accurate and intact error correction. <i>Nature Communications</i> , 2021, 12, 60.   | 12.8 | 166       |
| 2  | Environmental Factors Shape Water Microbial Community Structure and Function in Shrimp Cultural Enclosure Ecosystems. <i>Frontiers in Microbiology</i> , 2017, 8, 2359.  | 3.5  | 137       |
| 3  | Microecological Koch's postulates reveal that intestinal microbiota dysbiosis contributes to shrimp white feces syndrome. <i>Microbiome</i> , 2020, 8, 32.   | 11.1 | 126       |
| 4  | Intestinal bacterial signatures of white feces syndrome in shrimp. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 3701-3709.   | 3.6  | 118       |
| 5  | Composition, diversity and function of intestinal microbiota in pacific white shrimp ( <i>Litopenaeus</i> ) Tj ETQq1 1 0.784314 rgBT / Over  | 2.0  | 108       |
| 6  | Comparative analysis of the bacterial community compositions of the shrimp intestine, surrounding water and sediment. <i>Journal of Applied Microbiology</i> , 2018, 125, 792-799.   | 3.1  | 72        |
| 7  | Host development overwhelms environmental dispersal in governing the ecological succession of zebrafish gut microbiota. <i>Npj Biofilms and Microbiomes</i> , 2021, 7, 5.  | 6.4  | 64        |
| 8  | Potential biosorbent based on sugarcane bagasse modified with tetraethylenepentamine for removal of eosin Y. <i>International Journal of Biological Macromolecules</i> , 2012, 50, 707-712.  | 7.5  | 38        |
| 9  | Occurrence of human pathogenic bacteria carrying antibiotic resistance genes revealed by metagenomic approach: A case study from an aquatic environment. <i>Journal of Environmental Sciences</i> , 2019, 80, 248-256.   | 6.1  | 31        |
| 10 | Antibiotic supplement in feed can perturb the intestinal microbial composition and function in Pacific white shrimp. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 3111-3122.   | 3.6  | 28        |
| 11 | Immunological evaluation of <i>Vibrio alginolyticus</i> , <i>Vibrio harveyi</i> , <i>Vibrio vulnificus</i> and infectious spleen and kidney necrosis virus (ISKNV) combined-vaccine efficacy in <i>Epinephelus coioides</i> . <i>Veterinary Immunology and Immunopathology</i> , 2012, 150, 61-68. | 1.2  | 27        |
| 12 | Dissimilarity of microbial diversity of pond water, shrimp intestine and sediment in Aquamimicry system. <i>AMB Express</i> , 2020, 10, 180.   | 3.0  | 23        |
| 13 | Identification of Multigene Biomarker for Shrimp White Feces Syndrome by Full-Length Transcriptome Sequencing. <i>Frontiers in Genetics</i> , 2020, 11, 71.  | 2.3  | 22        |
| 14 | Intestine Bacterial Community Composition of Shrimp Varies Under Low- and High-Salinity Culture Conditions. <i>Frontiers in Microbiology</i> , 2020, 11, 589164.   | 3.5  | 20        |
| 15 | Stochastic processes shape the bacterial community assembly in shrimp cultural pond sediments. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 5013-5022.   | 3.6  | 20        |
| 16 | Community diversity and abundance of ammonia-oxidizing archaea and bacteria in shrimp pond sediment at different culture stages. <i>Journal of Applied Microbiology</i> , 2021, 130, 1442-1455.  | 3.1  | 18        |
| 17 | Distinct bacterial communities in the environmental water, sediment and intestine between two crayfish-plant coculture ecosystems. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 5087-5101.   | 3.6  | 17        |
| 18 | Sediment microbiota in polyculture of shrimp and fish pattern is distinctive from those in monoculture intensive shrimp or fish ponds. <i>Science of the Total Environment</i> , 2021, 787, 147594.  | 8.0  | 16        |

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|----|--|-----|-----------|
| 19 | Shrimp TAB1 interacts with TAK1 and p38 and activates the host innate immune response to bacterial infection. <i>Molecular Immunology</i> , 2017, 88, 10-19.   | 2.2 | 15        |
| 20 | Temporal variation of antibiotic resistance genes carried by culturable bacteria in the shrimp hepatopancreas and shrimp culture pond water. <i>Ecotoxicology and Environmental Safety</i> , 2020, 199, 110738.      | 6.0 | 15        |
| 21 | Interactions and Stability of Gut Microbiota in Zebrafish Increase with Host Development. <i>Microbiology Spectrum</i> , 2022, 10, e0169621.   | 3.0 | 11        |
| 22 | Environmental Water and Sediment Microbial Communities Shape Intestine Microbiota for Host Health: The Central Dogma in an Anthropogenic Aquaculture Ecosystem. <i>Frontiers in Microbiology</i> , 2021, 12, 772149. | 3.5 | 8         |
| 23 | Bacterial and eukaryotic community interactions might contribute to shrimp culture pond soil ecosystem at different culture stages. <i>Soil Ecology Letters</i> , 0, , 1.  | 4.5 | 2         |
| 24 | Sedimentary Nitrogen and Sulfur Reduction Functional-Couplings Interplay With the Microbial Community of Anthropogenic Shrimp Culture Pond Ecosystem. <i>Frontiers in Microbiology</i> , 2022, 13, 830777.           | 3.5 | 2         |
| 25 | Abundant and Rare Microbial Communities Respectively Contribute to an Aquaculture Pond Ecosystem. <i>Frontiers in Marine Science</i> , 2022, 9, .  | 2.5 | 2         |