

# Baozhong Liu

## List of Publications by Year in descending order

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103  
papers

2,732  
citations

201385

27  
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223531

46  
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106  
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106  
docs citations

106  
times ranked

2388  
citing authors

#	ARTICLE	IF	CITATIONS
1	Early mesodermal development in the patellogastropod <i>Lottia goshimai</i> . <i>Evolutionary Applications</i> , 2023, 16, 250-261.	1.5	2
2	Molluscan Dorsal-Ventral Patterning Relying on BMP2/4 and Chordin Provides Insights into Spiralian Development and Evolution. <i>Molecular Biology and Evolution</i> , 2022, 39, .	3.5	16
3	Changes in gluconeogenesis pathways and key genes associated with mass mortality in the clam <i>Meretrix petechialis</i> upon <i>Vibrio</i> infection. <i>Aquaculture</i> , 2022, 548, 737691.	1.7	2
4	Proteomics reveals the changes in energy metabolism associated with reproduction in the clam <i>Meretrix petechialis</i> . <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2022, 41, 100954.	0.4	2
5	Nonmuscle Myosin II is Required for Larval Shell Formation in a Patellogastropod. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 813741.	1.8	1
6	Transcriptomic analysis reveals the immune changes associated with reproduction in the clam <i>Meretrix petechialis</i> . <i>Fish and Shellfish Immunology</i> , 2021, 108, 24-31.	1.6	3
7	Microbial community changes in the digestive tract of the clam <i>Meretrix petechialis</i> in response to <i>Vibrio parahaemolyticus</i> challenge. <i>Journal of Oceanology and Limnology</i> , 2021, 39, 329-339.	0.6	5
8	Lipid metabolism changes in clam <i>Meretrix petechialis</i> in response to <i>Vibrio</i> infection and the identification of <i>Vibrio</i> -resistance markers. <i>Aquaculture</i> , 2021, 539, 736611.	1.7	11
9	CRISPR/Cas9-mediated mutagenesis reveals the roles of calaxin in gastropod larval cilia. <i>Gene</i> , 2021, 787, 145640.	1.0	5
10	Dorsoventral decoupling of Hox gene expression underpins the diversification of molluscs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 503-512.	3.3	37
11	Heritability of resistance-related gene expression traits and their correlation with body size of clam <i>Meretrix petechialis</i> . <i>Journal of Oceanology and Limnology</i> , 2020, 38, 571-578.	0.6	2
12	TAF5L functions as transcriptional coactivator of MITF involved in the immune response of the clam <i>Meretrix petechialis</i> . <i>Fish and Shellfish Immunology</i> , 2020, 98, 1017-1023.	1.6	7
13	Early shell field morphogenesis of a patellogastropod mollusk predominantly relies on cell movement and F-actin dynamics. <i>BMC Developmental Biology</i> , 2020, 20, 18.	2.1	6
14	Identification of three cell populations from the shell gland of a bivalve mollusc. <i>Development Genes and Evolution</i> , 2020, 230, 39-45.	0.4	8
15	MITF Regulates Downstream Genes in Response to <i>Vibrio parahaemolyticus</i> Infection in the Clam <i>Meretrix Petechialis</i> . <i>Frontiers in Immunology</i> , 2019, 10, 1547.	2.2	12
16	Dynamic immune and metabolism response of clam <i>Meretrix petechialis</i> to <i>Vibrio</i> challenge revealed by a time series of transcriptome analysis. <i>Fish and Shellfish Immunology</i> , 2019, 94, 17-26.	1.6	20
17	p38 MAPK is involved in the immune response to pathogenic <i>Vibrio</i> in the clam <i>Meretrix petechialis</i> . <i>Fish and Shellfish Immunology</i> , 2019, 95, 456-463.	1.6	6
18	Gill symbionts of the cold-seep mussel <i>Bathymodiolus platifrons</i> : Composition, environmental dependency and immune control. <i>Fish and Shellfish Immunology</i> , 2019, 86, 246-252.	1.6	18

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19	An investigation of oyster TGF- $\beta$ 2 receptor genes and their potential roles in early molluscan development. <i>Gene</i> , 2018, 663, 65-71.	1.0	13
20	Integrated transcriptomic and proteomic analyses reveal potential mechanisms linking thermal stress and depressed disease resistance in the turbot <i>Scophthalmus maximus</i> . <i>Scientific Reports</i> , 2018, 8, 1896.	1.6	17
21	Integrating the <i>Vibrio</i> -resistance phenotype and gene expression data for discovery of markers used for resistance evaluation in the clam <i>Meretrix petechialis</i> . <i>Aquaculture</i> , 2018, 482, 130-136.	1.7	8
22	Identification of a gene encoding microphthalmia-associated transcription factor and its association with shell color in the clam <i>Meretrix petechialis</i> . <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2018, 225, 75-83.	0.7	25
23	Transcriptome profiles of the clam <i>Meretrix petechialis</i> hepatopancreas in response to <i>Vibrio</i> infection. <i>Fish and Shellfish Immunology</i> , 2017, 62, 175-183.	1.6	32
24	Population genetics of the Manila clam ( <i>Ruditapes philippinarum</i> ) introduced in North America and Europe. <i>Scientific Reports</i> , 2017, 7, 39745.	1.6	62
25	Mining and identification of SNP markers associated with growth traits in the clam <i>Meretrix meretrix</i> . <i>Aquaculture International</i> , 2017, 25, 1185-1196.	1.1	0
26	Scallop genome provides insights into evolution of bilaterian karyotype and development. <i>Nature Ecology and Evolution</i> , 2017, 1, 120.	3.4	353
27	A <i>SoxC</i> gene related to larval shell development and co-expression analysis of different shell formation genes in early larvae of oyster. <i>Development Genes and Evolution</i> , 2017, 227, 181-188.	0.4	10
28	Expression patterns indicate that BMP2/4 and Chordin, not BMP5-8 and Gremlin, mediate dorsal-ventral patterning in the mollusk <i>Crassostrea gigas</i> . <i>Development Genes and Evolution</i> , 2017, 227, 75-84.	0.4	28
29	Identification of an <i>MITF</i> gene and its polymorphisms associated with the <i>Vibrio</i> resistance trait in the clam <i>Meretrix petechialis</i> . <i>Fish and Shellfish Immunology</i> , 2017, 68, 466-473.	1.6	21
30	A comparative proteomic analysis reveals important proteins for the fertilization and early embryonic development of the oyster <i>Crassostrea gigas</i> . <i>Proteomics</i> , 2017, 17, 1600251.	1.3	1
31	Genetic variation in <i>vibrio</i> resistance in the clam <i>Meretrix petechialis</i> under the challenge of <i>Vibrio parahaemolyticus</i> . <i>Aquaculture</i> , 2017, 468, 458-463.	1.7	31
32	The polymorphisms of a <i>MIF</i> gene and their association with <i>Vibrio</i> resistance in the clam <i>Meretrix meretrix</i> . <i>Developmental and Comparative Immunology</i> , 2016, 62, 116-126.	1.0	11
33	Influence of parental sample sizes on the estimating genetic parameters in cultured clam <i>Meretrix meretrix</i> based on factorial mating designs. <i>Acta Oceanologica Sinica</i> , 2016, 35, 42-49.	0.4	2
34	Characterization and expression of a novel caspase gene: Evidence of the expansion of caspases in <i>Crassostrea gigas</i> . <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2016, 201, 37-45.	0.7	10
35	Assessment of housekeeping genes as internal references in quantitative expression analysis during early development of oyster. <i>Genes and Genetic Systems</i> , 2016, 91, 257-265.	0.2	20
36	Identification of the <i>MmeHairy</i> gene and expression analysis affected by two SNPs in the 3'-untranslated region in the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2016, 51, 46-52.	1.6	1

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37	The impact of selection on population genetic structure in the clam <i>Meretrix petechialis</i> revealed by microsatellite markers. <i>Genetica</i> , 2016, 144, 1-8.	0.5	19
38	Characterization of a long-chain fatty acid-CoA ligase 1 gene and association between its SNPs and growth traits in the clam <i>Meretrix meretrix</i> . <i>Gene</i> , 2015, 566, 194-200.	1.0	10
39	The sperm proteome of the Pacific oyster <i>Crassostrea gigas</i> and immunolocalization of heat shock proteins. <i>Invertebrate Reproduction and Development</i> , 2015, 59, 111-118.	0.3	4
40	Development of <i>Vibrio</i> spp. infection resistance related SNP markers using multiplex SNaPshot genotyping method in the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2015, 43, 469-476.	1.6	25
41	Transcriptome Analysis of Shell Color-Related Genes in the Clam <i>Meretrix meretrix</i> . <i>Marine Biotechnology</i> , 2015, 17, 364-374.	1.1	61
42	Identification of a Serum amyloid A gene and the association of SNPs with <i>Vibrio</i> -resistance and growth traits in the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2015, 43, 301-309.	1.6	18
43	A calaxin Gene in the Pacific Oyster <i>Crassostrea gigas</i> and Its Potential Roles in Cilia. <i>Zoological Science</i> , 2015, 32, 419.	0.3	3
44	A GATA2/3 gene potentially involved in larval shell formation of the Pacific oyster <i>Crassostrea gigas</i> . <i>Development Genes and Evolution</i> , 2015, 225, 253-257.	0.4	15
45	The role of Cu/Zn-SOD and Mn-SOD in the immune response to oxidative stress and pathogen challenge in the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2015, 42, 58-65.	1.6	102
46	A Label-Free Proteomic Analysis on Competent Larvae and Juveniles of the Pacific Oyster <i>Crassostrea gigas</i> . <i>PLoS ONE</i> , 2015, 10, e0135008.	1.1	20
47	Combining ability and heterosis analysis over two environments in a diallel cross of three families of the clam <i>Meretrix meretrix</i> . <i>Acta Oceanologica Sinica</i> , 2014, 33, 37-42.	0.4	9
48	The involvement of cysteine-rich intestinal protein in early development and innate immunity of Asiatic hard clam, <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2014, 40, 435-440.	1.6	2
49	Assessment of parental contribution and effective population size from a 3 $\times$ 3 diallel cross of clam <i>Meretrix meretrix</i> . <i>Chinese Journal of Oceanology and Limnology</i> , 2014, 32, 306-315.	0.7	1
50	Cloning and expression patterns of two Smad genes during embryonic development and shell formation of the Pacific oyster <i>Crassostrea gigas</i> . <i>Chinese Journal of Oceanology and Limnology</i> , 2014, 32, 1224-1231.	0.7	9
51	An EGFR gene of the Pacific oyster <i>Crassostrea gigas</i> functions in wound healing and promotes cell proliferation. <i>Molecular Biology Reports</i> , 2014, 41, 2757-2765.	1.0	11
52	Multiple ferritin subunit genes of the Pacific oyster <i>Crassostrea gigas</i> and their distinct expression patterns during early development. <i>Gene</i> , 2014, 546, 80-88.	1.0	20
53	Identification of a tyrosinase gene potentially involved in early larval shell biogenesis of the Pacific oyster <i>Crassostrea gigas</i> . <i>Development Genes and Evolution</i> , 2013, 223, 389-394.	0.4	47
54	Transcriptional response of lysozyme, metallothionein, and superoxide dismutase to combined exposure to heavy metals and bacteria in <i>Macrura veneriformis</i> . <i>Comparative Biochemistry and Physiology Part - C: Toxicology and Pharmacology</i> , 2013, 157, 54-62.	1.3	18

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55	Growth performance of the clam, <i>Meretrix meretrix</i> , breeding-selection populations cultured in different conditions. <i>Acta Oceanologica Sinica</i> , 2013, 32, 82-87.	0.4	1
56	The role of catalase in the immune response to oxidative stress and pathogen challenge in the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2013, 34, 91-99.	1.6	59
57	Expression patterns of an i-type lysozyme in the clam <i>Meretrix meretrix</i> along with larval development. <i>Developmental and Comparative Immunology</i> , 2013, 41, 27-32.	1.0	6
58	Three vibrio-resistance related EST-SSR markers revealed by selective genotyping in the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2013, 35, 421-428.	1.6	17
59	Three EST-SSR Markers Associated with QTL for the Growth of the Clam <i>Meretrix meretrix</i> Revealed by Selective Genotyping. <i>Marine Biotechnology</i> , 2013, 15, 16-25.	1.1	27
60	Single nucleotide polymorphisms in i-type lysozyme gene and their correlation with vibrio-resistance and growth of clam <i>Meretrix meretrix</i> based on the selected resistance stocks. <i>Fish and Shellfish Immunology</i> , 2012, 33, 559-568.	1.6	33
61	Hatching, growth and hatchling dietary preference in <i>Hemifusus ternatanus</i> (Gmelin, 1791). <i>Aquaculture</i> , 2012, 326-329, 141-147.	1.7	7
62	Identification of differentially expressed proteins involved in the early larval development of the Pacific oyster <i>Crassostrea gigas</i> . <i>Journal of Proteomics</i> , 2012, 75, 3855-3865.	1.2	41
63	Toxic effects of benzo[a]pyrene (Bap) and Aroclor1254 on embryogenesis, larval growth, survival and metamorphosis of the bivalve <i>Meretrix meretrix</i> . <i>Ecotoxicology</i> , 2012, 21, 1617-1624.	1.1	11
64	Tissue-specific response of metallothionein and superoxide dismutase in the clam <i>Macraa veneriformis</i> under sublethal mercury exposure. <i>Ecotoxicology</i> , 2012, 21, 1593-1602.	1.1	16
65	RNAi assay in primary cells: a new method for gene function analysis in marine bivalve. <i>Molecular Biology Reports</i> , 2012, 39, 8209-8216.	1.0	15
66	Microsatellite-based genetic and growth analysis for a diallel mating design of two stocks of the clam, <i>Meretrix meretrix</i> . <i>Aquaculture Research</i> , 2012, 43, 260-270.	0.9	10
67	Genetic diversity of the sulfotransferase-like gene and one nonsynonymous SNP associated with growth traits of clam, <i>Meretrix meretrix</i> . <i>Molecular Biology Reports</i> , 2012, 39, 1323-1331.	1.0	9
68	Identification of a fructose-1,6-bisphosphate aldolase gene and association of the single nucleotide polymorphisms with growth traits in the clam <i>Meretrix meretrix</i> . <i>Molecular Biology Reports</i> , 2012, 39, 5017-5024.	1.0	10
69	Transcriptomic Analysis of the Clam <i>Meretrix meretrix</i> on Different Larval Stages. <i>Marine Biotechnology</i> , 2012, 14, 69-78.	1.1	84
70	The Potential Roles of a Laminin Receptor in Adhesion and Apoptosis of Cells of the Marine Bivalve <i>Meretrix meretrix</i> . <i>PLoS ONE</i> , 2012, 7, e47104.	1.1	7
71	Complete mtDNA of the <i>Meretrix lamarckii</i> (Bivalvia: Veneridae) and molecular identification of suspected <i>M. lamarckii</i> based on the whole mitochondrial genome. <i>Marine Genomics</i> , 2011, 4, 263-271.	0.4	19
72	Introduction of the Peruvian scallop and its hybridization with the bay scallop in China. <i>Aquaculture</i> , 2011, 310, 380-387.	1.7	75

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73	An effective method for parentage determination of the clam ( <i>Meretrix meretrix</i> ) based on SSR and COI markers. <i>Aquaculture</i> , 2011, 318, 223-228.	1.7	13
74	Large scale gene expression profiling during intestine and body wall regeneration in the sea cucumber <i>Apostichopus japonicus</i> . <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2011, 6, 195-205.	0.4	85
75	An i-type lysozyme from the Asiatic hard clam <i>Meretrix meretrix</i> potentially functioning in host immunity. <i>Fish and Shellfish Immunology</i> , 2011, 30, 550-558.	1.6	50
76	Cloning and characterization of a hsp70 gene from Asiatic hard clam <i>Meretrix meretrix</i> which is involved in the immune response against bacterial infection. <i>Fish and Shellfish Immunology</i> , 2011, 30, 791-799.	1.6	47
77	Molecular characterization of a glutathione peroxidase gene and its expression in the selected <i>Vibrio</i> -resistant population of the clam <i>Meretrix meretrix</i> . <i>Fish and Shellfish Immunology</i> , 2011, 30, 1294-1302.	1.6	31
78	Comparative proteomic analysis of challenged Zhikong scallop ( <i>Chlamys farreri</i> ): A new insight into the anti- <i>Vibrio</i> immune response of marine bivalves. <i>Fish and Shellfish Immunology</i> , 2011, 31, 1186-1192.	1.6	23
79	Isolation and characterization of a virulent <i>Vibrio</i> sp. bacterium from clams ( <i>Meretrix meretrix</i> ) with mass mortality. <i>Journal of Invertebrate Pathology</i> , 2011, 106, 242-249.	1.5	20
80	Mining of EST-SSR markers in clam <i>Meretrix meretrix</i> larvae from 454 shotgun transcriptome. <i>Genes and Genetic Systems</i> , 2011, 86, 197-205.	0.2	25
81	Estimation of genetic parameters for growth traits in cultured clam <i>Meretrix meretrix</i> (Bivalvia): Tj ETQq1 1 0.784314 rgBT /Overlock 0.9 36		
82	Recombinant expression, characterization and expressional analysis of clam <i>Meretrix meretrix</i> cathepsin B, an enzyme involved in nutrient digestion. <i>Molecular Biology Reports</i> , 2011, 38, 1861-1868.	1.0	8
83	Characterization of EST-SSR and genomic-SSR markers in the clam, <i>Meretrix meretrix</i> . <i>Conservation Genetics Resources</i> , 2011, 3, 655-658.	0.4	25
84	Potential role of cathepsin B in the embryonic and larval development of clam <i>Meretrix meretrix</i> . , 2011, 316B, 306-312.		5
85	Growth, survival and immune activity of scallops, <i>Chlamys farreri</i> Jones et Preston, compared between suspended and bottom culture in Haizhou Bay, China. <i>Aquaculture Research</i> , 2010, 41, 814-827.	0.9	10
86	Survival, growth and immune activity of scallop <i>Chlamys farreri</i> cultured at different depths in Haizhou Bay (Yellow Sea, China) during hot season. <i>Chinese Journal of Oceanology and Limnology</i> , 2010, 28, 498-507.	0.7	3
87	Seasonal variations in growth and clearance rate of the Zhikong scallop <i>Chlamys farreri</i> suspended in the deep water of Haizhou Bay, China. <i>Aquaculture International</i> , 2010, 18, 813-824.	1.1	5
88	Metallothionein and superoxide dismutase responses to sublethal cadmium exposure in the clam <i>Mactra veneriformis</i> . <i>Comparative Biochemistry and Physiology Part - C: Toxicology and Pharmacology</i> , 2010, 151, 325-333.	1.3	46
89	Identification and characterization of the pathogenic effect of a <i>Vibrio parahaemolyticus</i> -related bacterium isolated from clam <i>Meretrix meretrix</i> with mass mortality. <i>Journal of Invertebrate Pathology</i> , 2010, 103, 109-115.	1.5	67
90	Physiological and immune responses of zhikong scallop <i>Chlamys farreri</i> to the acute viral necrobiosis virus infection. <i>Fish and Shellfish Immunology</i> , 2010, 29, 42-48.	1.6	31

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91	Complete mtDNA of <i>Meretrix lusoria</i> (Bivalvia: Veneridae) reveals the presence of an <i>atp8</i> gene, length variation and heteroplasmy in the control region. <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2010, 5, 256-264.	0.4	18
92	Analysis of metallothionein expression and antioxidant enzyme activities in <i>Meretrix meretrix</i> larvae under sublethal cadmium exposure. <i>Aquatic Toxicology</i> , 2010, 100, 321-328.	1.9	23
93	Toxicity of lead, cadmium and mercury on embryogenesis, survival, growth and metamorphosis of <i>Meretrix meretrix</i> larvae. <i>Ecotoxicology</i> , 2009, 18, 829-837.	1.1	43
94	Cloning, characterization and expression of ferritin subunit from clam <i>Meretrix meretrix</i> in different larval stages. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2009, 154, 12-16.	0.7	33
95	Catecholaminergic responses to environmental stress in the hemolymph of zhikong scallop <i>Chlamys farreri</i> . <i>Journal of Experimental Zoology</i> , 2008, 309A, 289-296.	1.2	35
96	Molecular cloning and functional analysis of cathepsin B in nutrient metabolism during larval development in <i>Meretrix meretrix</i> . <i>Aquaculture</i> , 2008, 282, 41-46.	1.7	24
97	Preliminary Studies on Cryopreservation of Sydney Rock Oyster ( <i>Saccostrea glomerata</i> ) Larvae. <i>Journal of Shellfish Research</i> , 2008, 27, 1125-1128.	0.3	14
98	Particle-attached and free-living bacterial communities in a contrasting marine environment: Victoria Harbor, Hong Kong. <i>FEMS Microbiology Ecology</i> , 2007, 61, 496-508.	1.3	109
99	The phylogeny of native and exotic scallops cultured in China based on 16S rDNA sequences. <i>Chinese Journal of Oceanology and Limnology</i> , 2007, 25, 85-90.	0.7	4
100	Effects of various algal diets and starvation on larval growth and survival of <i>Meretrix meretrix</i> . <i>Aquaculture</i> , 2006, 254, 526-533.	1.7	81
101	Effect of stocking density on growth, settlement and survival of clam larvae, <i>Meretrix meretrix</i> . <i>Aquaculture</i> , 2006, 258, 344-349.	1.7	103
102	Pharmacological and immunocytochemical investigation of the role of catecholamines on larval metamorphosis by $\beta^2$ -adrenergic-like receptor in the bivalve <i>Meretrix meretrix</i> . <i>Aquaculture</i> , 2006, 258, 611-618.	1.7	34
103	Molecular Phylogeny and Species Identification of Pufferfish of the Genus <i>Takifugu</i> (Tetraodontiformes, Tetraodontidae). <i>Marine Biotechnology</i> , 2001, 3, 398-406.	1.1	27