Chenghua Li

List of Publications by Year in descending order

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99 99 3040
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#	Article	IF	CITATIONS
1	Radioprotective effect of Xâ€ray abdominal FLASH irradiation: Adaptation to oxidative damage and inflammatory response may be benefiting factors. Medical Physics, 2022, 49, 4812-4822.	3.0	18
2	CDK1 Promotes Epithelial–Mesenchymal Transition and Migration of Head and Neck Squamous Carcinoma Cells by Repressing â^†Np63α-Mediated Transcriptional Regulation. International Journal of Molecular Sciences, 2022, 23, 7385.	4.1	5
3	Copper metabolism in Saccharomyces cerevisiae: an update. BioMetals, 2021, 34, 3-14.	4.1	39
4	Pin1 and JNK1 cooperatively modulate TAp63γ. FEBS Open Bio, 2021, 11, 890-897.	2.3	1
5	Ultraviolet B irradiation upâ€regulates MM1 and induces photoageing of the epidermis. Photodermatology Photoimmunology and Photomedicine, 2021, 37, 395-403.	1.5	4
6	WW Domain-Containing E3 Ubiquitin Protein Ligase 1: A Self-Disciplined Oncoprotein. Frontiers in Cell and Developmental Biology, 2021, 9, 757493.	3.7	6
7	Prefoldin subunit MM1 promotes cell migration via facilitating filopodia formation. Biochemical and Biophysical Research Communications, 2020, 533, 613-619.	2.1	6
8	The molecular mechanisms of copper metabolism and its roles in human diseases. Pflugers Archiv European Journal of Physiology, 2020, 472, 1415-1429.	2.8	167
9	A double dealing tale of p63: an oncogene or a tumor suppressor. Cellular and Molecular Life Sciences, 2018, 75, 965-973.	5.4	71
10	Cullin3/ $<$ scp> $<$ KCTD $<$ /scp> $>$ 5 induces monoubiquitination of \hat{I} "Np63 \hat{I} \pm and impairs its activity. FEBS Letters, 2018, 592, 2334-2340.	2.8	8
11	ΒNp63α down-regulates c-Myc modulator MM1 via E3 ligase HERC3 in the regulation of cell senescence. Cell Death and Differentiation, 2018, 25, 2118-2129.	11.2	26
12	DNA damage induces expression of WWP1 to target î"Np63α to degradation. PLoS ONE, 2017, 12, e0176142.	2.5	8
13	Metabolic product response profiles of Cherax quadricarinatus towards white spot syndrome virus infection. Developmental and Comparative Immunology, 2016, 61, 236-241.	2.3	29
14	Cloning and comparative analysis the proximal promoter activities of arginase and agmatinase genes in Apostichopus japonicus. Developmental and Comparative Immunology, 2016, 65, 299-308.	2.3	4
15	Cloning and characterization of Vshppd, a gene inducing haemolysis and immune response of Apostichopus japonicus. Aquaculture, 2016, 464, 246-252.	3.5	24
16	A CgIFNLP receptor from Crassostrea gigas and its activation of the related genes in human JAK/STAT signaling pathway. Developmental and Comparative Immunology, 2016, 65, 98-106.	2.3	21
17	Long-term effects of di-octyl phthalate on the expression of immune-related genes in Tegillarca granosa. Chinese Journal of Oceanology and Limnology, 2016, 34, 423-429.	0.7	4
18	A \hat{l}^2 -integrin from sea cucumber Apostichopus japonicus exhibits LPS binding activity and negatively regulates coelomocyte apoptosis. Fish and Shellfish Immunology, 2016, 52, 103-110.	3.6	22

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19	Molecular cloning and characterization of four caspases members in Apostichopus japonicus. Fish and Shellfish Immunology, 2016, 55, 203-211.	3.6	26
20	Advances, challenges, and directions in shrimp disease control: the guidelines from an ecological perspective. Applied Microbiology and Biotechnology, 2016, 100, 6947-6954.	3.6	107
21	NF-κB/Rel, not STAT5, regulates nitric oxide synthase transcription in Apostichopus japonicus. Developmental and Comparative Immunology, 2016, 61, 42-47.	2.3	17
22	A preliminary study on the antibacterial mechanism of Tegillarca granosa hemoglobin by derived peptides and peroxidase activity. Fish and Shellfish Immunology, 2016, 51, 9-16.	3.6	34
23	Inhibition of marine Vibrio sp. by pyoverdine from Pseudomonas aeruginosa PA1. Journal of Hazardous Materials, 2016, 302, 217-224.	12.4	44
24	Nemo like kinase negatively regulates NF-κB activation andÂcoelomocytes apoptosis in Apostichopus japonicus. Developmental and Comparative Immunology, 2016, 54, 109-115.	2.3	22
25	p63α modulates c-Myc activity via direct interaction and regulation of MM1 protein stability. Oncotarget, 2016, 7, 44277-44287.	1.8	16
26	Description and phylogeny of a new prostomatid, Metacystis similis nov. spec. (Protista, Ciliophora) from the East China Sea. Zootaxa, 2015, 4033, 584-92.	0.5	3
27	MiR-31 modulates coelomocytes ROS production via targeting p105 in Vibrio splendidus challenged sea cucumber Apostichopus japonicus inÂvitro and inÂvivo. Fish and Shellfish Immunology, 2015, 45, 293-299.	3.6	44
28	Identification and characterization of a novel Foxo transcription factors in Apostichopus japonicus. Fish and Shellfish Immunology, 2015, 44, 164-171.	3.6	8
29	Molecular characterization of two novel molecular chaperones in bacterial-challenged Apostichopus japonicus. Gene, 2015, 570, 141-149.	2.2	5
30	Three members in JAK/STAT signal pathway from the sea cucumber Apostichopus japonicus: Molecular cloning, characterization and function analysis. Fish and Shellfish Immunology, 2015, 46, 523-536.	3.6	28
31	miR-200 modulates coelomocytes antibacterial activities and LPS priming via targeting Tollip in Apostichopus japonicus. Fish and Shellfish Immunology, 2015, 45, 431-436.	3.6	26
32	Cloning and characterization of two lipopolysaccharide-binding protein/bactericidal permeability–increasing protein (LBP/BPI) genes from the sea cucumber Apostichopus japonicus with diversified function in modulating ROS production. Developmental and Comparative Immunology, 2015, 52, 88-97.	2.3	37
33	The Roles of Two miRNAs in Regulating the Immune Response of Sea Cucumber. Genetics, 2015, 201, 1397-1410.	2.9	44
34	PKCδstabilizes TAp63 to promote cell apoptosis. FEBS Letters, 2015, 589, 2094-2099.	2.8	8
35	JNK1 inhibits transcriptional and proâ€apoptotic activity of TAp63γ. FEBS Letters, 2015, 589, 3686-3690.	2.8	3
36	Metabolomic responses of sea cucumber Apostichopus japonicus to thermal stresses. Aquaculture, 2015, 435, 390-397.	3.5	60

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37	Characterization of two regulators of the TNF-α signaling pathway in Apostichopus japonicus: LPS-induced TNF-α factor and baculoviral inhibitor of apoptosis repeat-containing 2. Developmental and Comparative Immunology, 2015, 48, 138-142.	2.3	14
38	Exploiting Quorum Sensing Interfering Strategies in Gram-Negative Bacteria for the Enhancement of Environmental Applications. Frontiers in Microbiology, 2015, 6, 1535.	3.5	106
39	iTRAQ-Based Proteomics Reveals Novel Members Involved in Pathogen Challenge in Sea Cucumber Apostichopus japonicus. PLoS ONE, 2014, 9, e100492.	2.5	53
40	Regulation of p63 Protein Stability via Ubiquitin-Proteasome Pathway. BioMed Research International, 2014, 2014, 1-8.	1.9	40
41	Proteomic identification of differentially expressed proteins in sea cucumber Apostichopus japonicus coelomocytes after Vibrio splendidus infection. Developmental and Comparative Immunology, 2014, 44, 370-377.	2.3	32
42	Identification and characterization of miR-92a and its targets modulating Vibrio splendidus challenged Apostichopus japonicus. Fish and Shellfish Immunology, 2014, 38, 383-388.	3.6	58
43	Cloning and characterization of hemerythrin gene from Sipuncula Phascolosoma esculenta. Genes and Genomics, 2013, 35, 95-100.	1.4	8
44	Two adaptor molecules of MyD88 and TRAF6 in Apostichopus japonicus Toll signaling cascade: Molecular cloning and expression analysis. Developmental and Comparative Immunology, 2013, 41, 498-504.	2.3	55
45	Divergent Metabolic Responses of Apostichopus japonicus Suffered from Skin Ulceration Syndrome and Pathogen Challenge. Journal of Agricultural and Food Chemistry, 2013, 61, 10766-10771.	5. 2	46
46	Characterisation of immune-related gene expression in clam (VenerupisÂphilippinarum) under exposure to di(2-ethylhexyl) phthalate. Fish and Shellfish Immunology, 2013, 34, 142-146.	3.6	37
47	Characterization of two negative regulators of the Toll-like receptor pathway in Apostichopus japonicus: Inhibitor of NF-κB and Toll-interacting protein. Fish and Shellfish Immunology, 2013, 35, 1663-1669.	3.6	41
48	The link between selenium binding protein from Sinonovacula constricta and environmental pollutions exposure. Fish and Shellfish Immunology, 2013, 35, 271-277.	3.6	12
49	MUC1 activates JNK1 and inhibits apoptosis under genotoxic stress. Biochemical and Biophysical Research Communications, 2013, 440, 179-183.	2.1	29
50	Identification of differential expressed proteins and characterization their mRNA expression in thermally stressed Apostichopus japonicus. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2013, 8, 194-200.	1.0	11
51	A small heat shock protein (sHSP) from Sinonovacula constricta against heavy metals stresses. Fish and Shellfish Immunology, 2013, 34, 1605-1610.	3.6	17
52	De Novo Assembly of the Sea Cucumber Apostichopus japonicus Hemocytes Transcriptome to Identify miRNA Targets Associated with Skin Ulceration Syndrome. PLoS ONE, 2013, 8, e73506.	2.5	66
53	Low-affinity copper transporter CTR2 is regulated by copper-sensing transcription factor Mac1p in Saccharomyces cerevisiae. Biochemical and Biophysical Research Communications, 2012, 420, 600-604.	2.1	15
54	DNA damage down-regulates î"Np63î± and induces apoptosis independent of wild type p53. Biochemical and Biophysical Research Communications, 2012, 423, 338-343.	2.1	13

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55	Metal-sensing transcription factors $Mac1p$ and $Aft1p$ coordinately regulate vacuolar copper transporter CTR2 in Saccharomyces cerevisiae. Biochemical and Biophysical Research Communications, 2012, 423, 424-428.	2.1	11
56	Two classes of glutathione S-transferase genes with different response profiles to bacterial challenge in Venerupis philippinarum. Fish and Shellfish Immunology, 2012, 32, 219-222.	3.6	19
57	Characterization of skin ulceration syndrome associated microRNAs in sea cucumber Apostichopus japonicus by deep sequencing. Fish and Shellfish Immunology, 2012, 33, 436-441.	3.6	114
58	Development and Application of Reverse Transcription Loopâ€Mediated Isothermal Amplification for Detecting Live <i>Shewanella putrefaciens</i> in Preserved Fish Sample. Journal of Food Science, 2012, 77, M226-30.	3.1	14
59	Molecular cloning and expression analysis of a selenium-independent glutathione peroxidase identified from Manila clam Venerupis philippinarum. Aquaculture Research, 2012, 43, 1176-1183.	1.8	1
60	A Ferritin from Dendrorhynchus zhejiangensis with Heavy Metals Detoxification Activity. PLoS ONE, 2012, 7, e51428.	2.5	18
61	Isolation of an ATP synthase cDNA from Sinonovacula constricta and its mRNA expression by thermal stress. African Journal of Biotechnology, 2012, 11, .	0.6	2
62	Haemocyte protein expression profiling of scallop Chlamys farreri response to acute viral necrosis virus (AVNV) infection. Developmental and Comparative Immunology, 2011, 35, 1135-1145.	2.3	26
63	Cloning and characterization of allograft inflammatory factor-1 (AIF-1) from manila clam Venerupis philippinarum. Fish and Shellfish Immunology, 2011, 30, 148-153.	3.6	45
64	Alternation of Venerupis philippinarum Hsp40 gene expression in response to pathogen challenge and heavy metal exposure. Fish and Shellfish Immunology, 2011, 30, 447-450.	3.6	24
65	Identification and characterization of a clam ferritin from Sinonovacula constricta. Fish and Shellfish Immunology, 2011, 30, 1147-1151.	3.6	32
66	Cloning and characterization of a sialic acid binding lectins (SABL) from Manila clam Venerupis philippinarum. Fish and Shellfish Immunology, 2011, 30, 1202-1206.	3.6	36
67	Transcriptional regulation of selenium-dependent glutathione peroxidase from Venerupis philippinarum in response to pathogen and contaminants challenge. Fish and Shellfish Immunology, 2011, 31, 831-837.	3.6	33
68	Toxicological responses to acute mercury exposure for three species of Manila clam Ruditapes philippinarum by NMR-based metabolomics. Environmental Toxicology and Pharmacology, 2011, 31, 323-332.	4.0	69
69	A manganese superoxide dismutase in blood clam Tegillarca granosa: Molecular cloning, tissue distribution and expression analysis. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2011, 159, 64-70.	1.6	26
70	Differential toxicological effects induced by mercury in gills from three pedigrees of Manila clam Ruditapes philippinarum by NMR-based metabolomics. Ecotoxicology, 2011, 20, 177-186.	2.4	89
71	Assessment of Clam <i>Ruditapes philippinarum</i> as Heavy Metal Bioindicators Using NMRâ€Based Metabolomics. Clean - Soil, Air, Water, 2011, 39, 759-766.	1.1	28
72	Metabolomic Study on the Halophyte <i>Suaeda salsa</i> in the Yellow River Delta. Clean - Soil, Air, Water, 2011, 39, 720-727.	1.1	13

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73	Identification and characterization of a Tegillarca granosa ferritin regulated by iron ion exposure and thermal stress. Developmental and Comparative Immunology, 2011, 35, 745-751.	2.3	31
74	The Metal Chelating and Chaperoning Effects of Clioquinol: Insights from Yeast Studies. Journal of Alzheimer's Disease, 2010, 21, 1249-1262.	2.6	41
75	Identification of two small heat shock proteins with different response profile to cadmium and pathogen stresses in Venerupis philippinarum. Cell Stress and Chaperones, 2010, 15, 897-904.	2.9	24
76	Identification of a cathepsin D potentially involved in H2A cleavage from scallop Chlamys farreri. Molecular Biology Reports, 2010, 37, 1451-1460.	2.3	5
77	Molecular Characterization of a Novel Big Defensin from Clam Venerupis philippinarum. PLoS ONE, 2010, 5, e13480.	2.5	72
78	Identification and characterization of an intracellular Cu, Zn-superoxide dismutase (icCu/Zn-SOD) gene from clam Venerupis philippinarum. Fish and Shellfish Immunology, 2010, 28, 499-503.	3.6	52
79	The first molluscan TCTP in Venerupis philippinarum: Molecular cloning and expression analysis. Fish and Shellfish Immunology, 2010, 29, 530-533.	3.6	14
80	Cloning and characterization of an invertebrate type lysozyme from Venerupis philippinarum. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2010, 156, 56-60.	1.6	47
81	Effects of manganese and hypoxia on coelomocyte renewal in the echinoderm, Asterias rubens (L.). Aquatic Toxicology, 2010, 100, 84-90.	4.0	22
82	Cloning and characterization of a novel C-type lectin gene from shrimp Litopenaeus vannamei. Fish and Shellfish Immunology, 2009, 26, 183-192.	3.6	82
83	Molecular cloning and characterization of peroxiredoxin 6 from Chinese mitten crab Eriocheir sinensis. Fish and Shellfish Immunology, 2009, 26, 821-827.	3.6	58
84	The expression of peptidoglycan recognition protein-S1 gene in the scallop Chlamys farreri was enhanced after a second challenge by Listonella anguillarum. Journal of Invertebrate Pathology, 2009, 100, 120-122.	3.2	15
85	Molecular cloning and responsive expression to injury stimulus of a defender against cell death 1 (DAD1) gene from bay scallops Argopecten irradians. Molecular Biology Reports, 2008, 35, 125-132.	2.3	22
86	A novel serine protease with clip domain from scallop Chlamys farreri. Molecular Biology Reports, 2008, 35, 257-264.	2.3	20
87	Molecular cloning and characterization of a catalase gene from Zhikong scallop Chlamys farreri. Fish and Shellfish Immunology, 2008, 24, 26-34.	3.6	85
88	A prophenoloxidase from the Chinese mitten crab Eriocheir sinensis: Gene cloning, expression and activity analysis. Fish and Shellfish Immunology, 2008, 24, 156-167.	3.6	88
89	A lectin (CfLec-2) aggregating Staphylococcus haemolyticus from scallop Chlamys farreri. Fish and Shellfish Immunology, 2008, 24, 286-293.	3.6	63
90	Molecular cloning and expression of a novel Kazal-type serine proteinase inhibitor gene from Zhikong scallop Chlamys farreri, and the inhibitory activity of its recombinant domain. Fish and Shellfish Immunology, 2008, 24, 629-637.	3.6	43

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91	A novel C1q-domain-containing protein from Zhikong scallop Chlamys farreri with lipopolysaccharide binding activity. Fish and Shellfish Immunology, 2008, 25, 281-289.	3.6	137
92	Molecular cloning, genomic organization and functional analysis of an anti-lipopolysaccharide factor from Chinese mitten crab Eriocheir sinensis. Developmental and Comparative Immunology, 2008, 32, 784-794.	2.3	95
93	Preliminary study on a potential antibacterial peptide derived from histone H2A in hemocytes of scallop Chlamys farreri. Fish and Shellfish Immunology, 2007, 22, 663-672.	3.6	75
94	Molecular cloning, expression of a big defensin gene from bay scallop Argopecten irradians and the antimicrobial activity of its recombinant protein. Molecular Immunology, 2007, 44, 360-368.	2.2	149
95	Cloning and characterization of a novel C-type lectin from Zhikong scallop Chlamys farreri. Molecular Immunology, 2007, 44, 722-731.	2.2	135
96	Molecular cloning of an invertebrate goose-type lysozyme gene from Chlamys farreri, and lytic activity of the recombinant protein. Molecular Immunology, 2007, 44, 1198-1208.	2.2	146
97	Molecular cloning and characterization of a thioester-containing protein from Zhikong scallop Chlamys farreri. Molecular Immunology, 2007, 44, 3492-3500.	2.2	69
98	Genomic organization, nucleotide sequence analysis of the core histone genes cluster inChlamys farreriand molecular evolution assessment of the H2A and H2B. DNA Sequence, 2006, 17, 440-451.	0.7	9