

List of Publications by Year in
Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

201 papers	5,107 citations	39 h-index	56 g-index
211 ext. papers	6,056 ext. citations	4.1 avg, IF	6.31 L-index

#	Paper	IF	Citations
201	Megalocytivirus Induces Complicated Fish Immune Response at Multiple RNA Levels Involving mRNA, miRNA, and circRNA. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	2
200	A Crustin from Hydrothermal Vent Shrimp: Antimicrobial Activity and Mechanism. <i>Marine Drugs</i> , 2021 , 19,	6	1
199	A deep-sea pathogenic <i>Bacillus subtilis</i> isolate employs different strategies to escape the killing of teleost and murine complements. <i>Developmental and Comparative Immunology</i> , 2021 , 119, 104037	3.2	1
198	Pol-miR-150 regulates anti-bacterial and viral infection in Japanese flounder (<i>Paralichthys olivaceus</i>) via the lysosomal protein LAMP2L. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2021 , 254, 110578	2.3	1
197	Identification and characterization of immune-related lncRNAs and lncRNA-miRNA-mRNA networks of <i>Paralichthys olivaceus</i> involved in <i>Vibrio anguillarum</i> infection. <i>BMC Genomics</i> , 2021 , 22, 447	4.5	3
196	A virulent strain from deep-sea cold seep induces pyroptosis in a manner that involves NLRP3 inflammasome, JNK pathway, and lysosomal rupture. <i>Virulence</i> , 2021 , 12, 1362-1376	4.7	3
195	Systematic Identification and Analysis of Circular RNAs of Japanese Flounder () in Response to Infection. <i>Genes</i> , 2021 , 12,	4.2	4
194	Characterization of a Deep Sea Isolate: Genomic and Pathogenic Features. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021 , 11, 629116	5.9	4
193	A Novel microRNA of Japanese Flounder Regulates Antimicrobial Immunity Involving a Bacteria-Binding CSF3. <i>Frontiers in Immunology</i> , 2021 , 12, 723401	8.4	1
192	An inactivated bivalent vaccine effectively protects turbot (<i>Scophthalmus maximus</i>) against <i>Vibrio anguillarum</i> and <i>Vibrio harveyi</i> infection. <i>Aquaculture</i> , 2021 , 544, 737158	4.4	5
191	High-Throughput Sequencing Reveals a Potentially Novel Species Dominating the Microbial Communities of the Seawater-Sediment Interface of a Deep-Sea Cold Seep in South China Sea. <i>Microorganisms</i> , 2020 , 8,	4.9	7
190	Transcriptome analysis reveals seven key immune pathways of Japanese flounder (<i>Paralichthys olivaceus</i>) involved in megalocytivirus infection. <i>Fish and Shellfish Immunology</i> , 2020 , 103, 150-158	4.3	6
189	IL-34 regulates the inflammatory response and anti-bacterial immune defense of Japanese flounder <i>Paralichthys olivaceus</i> . <i>Fish and Shellfish Immunology</i> , 2020 , 104, 228-236	4.3	2
188	Micro-Transcriptome Analysis Reveals Immune-Related MicroRNA Regulatory Networks of Induced by Infection. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	3
187	Pol-miR-363-3p plays a significant role in the immune defense of Japanese flounder <i>Paralichthys olivaceus</i> against bacterial and viral infection. <i>Fish and Shellfish Immunology</i> , 2020 , 104, 439-446	4.3	0
186	Japanese flounder <i>Paralichthys olivaceus</i> interleukin 21 induces inflammatory response and plays a vital role in the immune defense against bacterial pathogen. <i>Fish and Shellfish Immunology</i> , 2020 , 98, 364-373	4.3	5
185	pol-miR-7133 and pol-miR-3p-9227 of Japanese flounder <i>Paralichthys olivaceus</i> modulate <i>Streptococcus iniae</i> infection through regulation of the common target gene LAMP2. <i>Aquaculture</i> , 2020 , 520, 734980	4.4	3

184	The novel fish miRNA pol-miR-novel_171 and its target gene FAM49B play a critical role in apoptosis and bacterial infection. <i>Developmental and Comparative Immunology</i> , 2020 , 106, 103616	3.2	11
183	Japanese flounder (<i>Paralichthys olivaceus</i>) Bmal1 is involved in the regulation of inflammatory response and bacterial infection. <i>Aquaculture</i> , 2020 , 525, 735330	4.4	3
182	Transcriptome Analysis of Erythrocytes Reveals Profound Immune Responses Induced by Infection. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	4
181	A Fish Galectin-8 Possesses Direct Bactericidal Activity. <i>International Journal of Molecular Sciences</i> , 2020 , 22,	6.3	6
180	Japanese flounder pol-miR-3p-2 suppresses <i>Edwardsiella tarda</i> infection by regulation of autophagy via p53. <i>Developmental and Comparative Immunology</i> , 2020 , 103, 103531	3.2	6
179	Gene network analysis reveals a core set of genes involved in the immune response of Japanese flounder (<i>Paralichthys olivaceus</i>) against <i>Vibrio anguillarum</i> infection. <i>Fish and Shellfish Immunology</i> , 2020 , 98, 800-809	4.3	14
178	Characterization of <i>Streptococcus iniae</i> -induced microRNA profiles in <i>Paralichthys olivaceus</i> and identification of pol-3p-10740_175 as a regulator of antibacterial immune response. <i>Fish and Shellfish Immunology</i> , 2020 , 98, 860-867	4.3	5
177	Coral gasdermin triggers pyroptosis. <i>Science Immunology</i> , 2020 , 5,	2.8	22
176	The Translocation and Assembly Module (TAM) of Is Essential for Stress Resistance and Host Infection. <i>Frontiers in Microbiology</i> , 2020 , 11, 1743	5.7	8
175	Phosphatase and Tensin Homolog (PTEN) of Japanese Flounder-Its Regulation by miRNA and Role in Autophagy, Apoptosis and Pathogen Infection. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	2
174	A First Study of the Virulence Potential of a Isolate From Deep-Sea Hydrothermal Vent. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019 , 9, 183	5.9	19
173	Teleost Gasdermin E Is Cleaved by Caspase 1, 3, and 7 and Induces Pyroptosis. <i>Journal of Immunology</i> , 2019 , 203, 1369-1382	5.3	33
172	A Comparative Analysis of -Induced Transcriptome Profiles in RAW264.7 Cells Reveals New Insights into the Strategy of Bacterial Immune Evasion. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	8
171	Characterization of the Genome Feature and Toxic Capacity of a Isolate From the Hydrothermal Field in Okinawa Trough. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019 , 9, 370	5.9	6
170	Global profiling and characterization of Japanese flounder (<i>Paralichthys olivaceus</i>) kidney microRNAs regulated by <i>Edwardsiella tarda</i> infection in a time-dependent fashion. <i>Fish and Shellfish Immunology</i> , 2019 , 93, 766-780	4.3	11
169	Particle and bacteria uptake by Japanese flounder (<i>Paralichthys olivaceus</i>) red blood cells: Size dependence and pathway specificity. <i>Tissue and Cell</i> , 2019 , 61, 79-88	2.7	3
168	Global profiling of megalocytivirus-induced proteins in tongue sole (<i>Cynoglossus semilaevis</i>) spleen identifies cellular processes essential to viral infection. <i>Developmental and Comparative Immunology</i> , 2019 , 92, 150-159	3.2	6
167	pol-miR-194a of Japanese flounder (<i>Paralichthys olivaceus</i>) suppresses type I interferon response and facilitates <i>Edwardsiella tarda</i> infection. <i>Fish and Shellfish Immunology</i> , 2019 , 87, 220-225	4.3	24

166	Macropinocytosis-dependent endocytosis of Japanese flounder IgM B cells and its regulation by CD22. <i>Fish and Shellfish Immunology</i> , 2019 , 84, 138-147	4.3	15
165	First characterization of an anti-lipopolysaccharide factor (ALF) from hydrothermal vent shrimp: Insights into the immune function of deep-sea crustacean ALF. <i>Developmental and Comparative Immunology</i> , 2018 , 84, 382-395	3.2	14
164	Internalization of large particles by turbot (<i>Scophthalmus maximus</i>) IgM B cells mainly depends on macropinocytosis. <i>Developmental and Comparative Immunology</i> , 2018 , 82, 31-38	3.2	18
163	Comparative metagenomic analysis of the microbial communities in the surroundings of Iheya north and Iheya ridge hydrothermal fields reveals insights into the survival strategy of microorganisms in deep-sea environments. <i>Journal of Marine Systems</i> , 2018 , 180, 102-111	2.7	6
162	Bacterial communities associated with Shinkaia crosnieri from the Iheya North, Okinawa Trough: Microbial diversity and metabolic potentials. <i>Journal of Marine Systems</i> , 2018 , 180, 228-236	2.7	2
161	Characterization of a teleost membrane-associated protein that is involved in the regulation of complement activation and bacterial infection. <i>Developmental and Comparative Immunology</i> , 2018 , 79, 142-149	3.2	8
160	Sip2: A Serum-Induced Protein That Is Essential to Serum Survival, Acid Resistance, Intracellular Replication, and Host Infection. <i>Frontiers in Microbiology</i> , 2018 , 9, 1084	5.7	14
159	First characterization of two C-type lectins of the tubeworm <i>Alaysia</i> sp. from a deep-sea hydrothermal vent. <i>Developmental and Comparative Immunology</i> , 2018 , 86, 17-25	3.2	3
158	Description of <i>Bacillus kexueae</i> sp. nov. and <i>Bacillus manusensis</i> sp. nov., isolated from hydrothermal sediments. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018 , 68, 829-834	2.2	7
157	<i>Muricauda iocasae</i> sp. nov., isolated from deep sea sediment of the South China Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018 , 68, 2538-2544	2.2	5
156	Histones and chymotrypsin-like elastases play significant roles in the antimicrobial activity of tongue sole neutrophil extracellular traps. <i>Fish and Shellfish Immunology</i> , 2018 , 72, 470-476	4.3	9
155	<i>Fictibacillus iocasae</i> sp. nov., isolated from the deep-sea sediment in Pacmanus, Manus Basin. <i>Archives of Microbiology</i> , 2018 , 200, 1123-1128	3	1
154	A teleost complement factor Ba possesses antimicrobial activity and inhibits bacterial infection in fish. <i>Developmental and Comparative Immunology</i> , 2017 , 71, 49-58	3.2	15
153	Draft Genome Sequence of sp. HVEsp1, a Thermophilic Bacterium Isolated from a Deep-Sea Hydrothermal Vent in the Okinawa Trough. <i>Genome Announcements</i> , 2017 , 5,		1
152	Comparative transcriptome analysis of <i>Rimicaris</i> sp. reveals novel molecular features associated with survival in deep-sea hydrothermal vent. <i>Scientific Reports</i> , 2017 , 7, 2000	4.9	20
151	Comparative metagenomics reveals insights into the deep-sea adaptation mechanism of the microorganisms in Iheya hydrothermal fields. <i>World Journal of Microbiology and Biotechnology</i> , 2017 , 33, 86	4.4	10
150	Tongue Sole CD209: A Pattern-Recognition Receptor that Binds a Broad Range of Microbes and Promotes Phagocytosis. <i>International Journal of Molecular Sciences</i> , 2017 , 18,	6.3	8
149	Transcriptome analysis reveals temperature-regulated antiviral response in turbot <i>Scophthalmus maximus</i> . <i>Fish and Shellfish Immunology</i> , 2017 , 68, 359-367	4.3	14

148	First characterization of fish CD22: An inhibitory role in the activation of peripheral blood leukocytes. <i>Veterinary Immunology and Immunopathology</i> , 2017 , 190, 39-44	2	6
147	Edwardsiella tarda-induced miRNAs in a teleost host: Global profile and role in bacterial infection as revealed by integrative miRNA-mRNA analysis. <i>Virulence</i> , 2017 , 8, 1457-1464	4.7	15
146	Cytochemical identification of turbot myeloperoxidase-positive granulocytes by potassium iodide and oxidized pyronine Y staining. <i>Tissue and Cell</i> , 2017 , 49, 751-755	2.7	2
145	A teleost CD46 is involved in the regulation of complement activation and pathogen infection. <i>Scientific Reports</i> , 2017 , 7, 15028	4.9	10
144	Intracellular Trafficking Pathways of : From Clathrin- and Caveolin-Mediated Endocytosis to Endosome and Lysosome. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017 , 7, 400	5.9	41
143	Neutrophil Extracellular Traps of : Production Characteristics and Antibacterial Effect. <i>Frontiers in Immunology</i> , 2017 , 8, 290	8.4	31
142	A comparative study revealed first insights into the diversity and metabolisms of the microbial communities in the sediments of Pacmanus and Desmos hydrothermal fields. <i>PLoS ONE</i> , 2017 , 12, e0181048	3.7	4
141	Description of Algoriphagus iocasae sp. nov., isolated from deep-sea sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017 , 67, 243-249	2.2	8
140	Bacillus iocasae sp. nov., isolated from Pacmanus hydrothermal field, Manus Basin. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017 , 67, 3547-3552	2.2	5
139	Three novel B-type mannose-specific lectins of Cynoglossus semilaevis possess varied antibacterial activities against Gram-negative and Gram-positive bacteria. <i>Developmental and Comparative Immunology</i> , 2016 , 55, 194-202	3.2	25
138	pol-miR-731, a teleost miRNA upregulated by megalocytivirus, negatively regulates virus-induced type I interferon response, apoptosis, and cell cycle arrest. <i>Scientific Reports</i> , 2016 , 6, 28354	4.9	38
137	Toll-like receptor 2 of tongue sole Cynoglossus semilaevis: Signaling pathway and involvement in bacterial infection. <i>Fish and Shellfish Immunology</i> , 2016 , 51, 321-328	4.3	19
136	Comparative analysis of the expression patterns of eight suppressors of cytokine signaling in tongue sole, Cynoglossus semilaevis. <i>Fish and Shellfish Immunology</i> , 2016 , 55, 595-601	4.3	16
135	Neutrophils of Scopthalmus maximus produce extracellular traps that capture bacteria and inhibit bacterial infection. <i>Developmental and Comparative Immunology</i> , 2016 , 56, 7-12	3.2	23
134	A Teleost Bactericidal Permeability-Increasing Protein Kills Gram-Negative Bacteria, Modulates Innate Immune Response, and Enhances Resistance against Bacterial and Viral Infection. <i>PLoS ONE</i> , 2016 , 11, e0154045	3.7	7
133	Description of sp. nov., isolated from deep-sea sediment, and emended description of the genus. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016 , 66, 982-987	2.2	11
132	Pseudomonas oceani sp. nov., isolated from deep seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016 , 66, 4250-4255	2.2	13
131	Pseudomonas fluorescens Filamentous Hemagglutinin, an Iron-Regulated Protein, Is an Important Virulence Factor that Modulates Bacterial Pathogenicity. <i>Frontiers in Microbiology</i> , 2016 , 7, 1320	5.7	20

130	Edwardsiella tarda-Induced Inhibition of Apoptosis: A Strategy for Intracellular Survival. <i>Frontiers in Cellular and Infection Microbiology</i> , 2016 , 6, 76	5.9	22
129	First Comparative Analysis of the Community Structures and Carbon Metabolic Pathways of the Bacteria Associated with Alvinocaris longirostris in a Hydrothermal Vent of Okinawa Trough. <i>PLoS ONE</i> , 2016 , 11, e0154359	3.7	9
128	CsMAP34, a teleost MAP with dual role: A promoter of MASP-assisted complement activation and a regulator of immune cell activity. <i>Scientific Reports</i> , 2016 , 6, 39287	4.9	22
127	CsSAP, a teleost serum amyloid P component, interacts with bacteria, promotes phagocytosis, and enhances host resistance against bacterial and viral infection. <i>Developmental and Comparative Immunology</i> , 2016 , 55, 12-20	3.2	26
126	The global regulatory effect of Edwardsiella tarda Fur on iron acquisition, stress resistance, and host infection: A proteomics-based interpretation. <i>Journal of Proteomics</i> , 2016 , 140, 100-10	3.9	18
125	Molecular characterization reveals involvement of four caspases in the antibacterial immunity of tongue sole (Cynoglossus semilaevis). <i>Fish and Shellfish Immunology</i> , 2016 , 57, 340-349	4.3	10
124	Tongue sole (Cynoglossus semilaevis) CD59: A complement inhibitor that binds bacterial cells and promotes bacterial escape from the killing of fish serum. <i>Fish and Shellfish Immunology</i> , 2016 , 58, 442-448	4.3	11
123	Edwardsiella tarda MliC, a lysozyme inhibitor that participates in pathogenesis in a manner that parallels Ivy. <i>Infection and Immunity</i> , 2015 , 83, 583-90	3.7	31
122	Edwardsiella tarda-regulated proteins in Japanese flounder (Paralichthys olivaceus): Identification and evaluation of antibacterial potentials. <i>Journal of Proteomics</i> , 2015 , 124, 1-10	3.9	34
121	CD83 is required for the induction of protective immunity by a DNA vaccine in a teleost model. <i>Developmental and Comparative Immunology</i> , 2015 , 51, 141-7	3.2	22
120	Edwardsiella tarda Sip1: A serum-induced zinc metalloprotease that is essential to serum resistance and host infection. <i>Veterinary Microbiology</i> , 2015 , 177, 332-40	3.3	25
119	Pseudomonas fluorescens: iron-responsive proteins and their involvement in host infection. <i>Veterinary Microbiology</i> , 2015 , 176, 309-20	3.3	13
118	TLR7 is required for optimal immune defense against bacterial infection in tongue sole (Cynoglossus semilaevis). <i>Fish and Shellfish Immunology</i> , 2015 , 47, 93-9	4.3	18
117	Characteristics of the cultivable bacteria from sediments associated with two deep-sea hydrothermal vents in Okinawa Trough. <i>World Journal of Microbiology and Biotechnology</i> , 2015 , 31, 2025-37	4.4	13
116	Comparative study of four interleukin 17 cytokines of tongue sole Cynoglossus semilaevis: Genomic structure, expression pattern, and promoter activity. <i>Fish and Shellfish Immunology</i> , 2015 , 47, 321-30	4.3	15
115	A short-type peptidoglycan recognition protein from tongue sole (Cynoglossus semilaevis) promotes phagocytosis and defense against bacterial infection. <i>Fish and Shellfish Immunology</i> , 2015 , 47, 313-20	4.3	16
114	Tongue sole (Cynoglossus semilaevis) prothymosin alpha: Cytokine-like activities associated with the intact protein and the C-terminal region that lead to antiviral immunity via Myd88-dependent and -independent pathways respectively. <i>Developmental and Comparative Immunology</i> , 2015 , 53, 96-104	3.2	34
113	Pseudomonas fluorescens: fur is required for multiple biological properties associated with pathogenesis. <i>Veterinary Microbiology</i> , 2015 , 175, 145-9	3.3	7

112	Immune effects of R848: evidences that suggest an essential role of TLR7/8-induced, Myd88- and NF- κ B-dependent signaling in the antiviral immunity of Japanese flounder (<i>Paralichthys olivaceus</i>). <i>Developmental and Comparative Immunology</i> , 2015 , 49, 113-20	3.2	40
111	P247 and p523: two in vivo-expressed megalocytivirus proteins that induce protective immunity and are essential to viral infection. <i>PLoS ONE</i> , 2015 , 10, e0121282	3.7	11
110	CsBAFF, a Teleost B Cell Activating Factor, Promotes Pathogen-Induced Innate Immunity and Vaccine-Induced Adaptive Immunity. <i>PLoS ONE</i> , 2015 , 10, e0136015	3.7	22
109	Microbial diversity in the deep-sea sediments of Iheya North and Iheya Ridge, Okinawa Trough. <i>Microbiological Research</i> , 2015 , 177, 43-52	5.3	47
108	Ferritin M of <i>Paralichthys olivaceus</i> possesses antimicrobial and antioxidative properties. <i>Fish Physiology and Biochemistry</i> , 2015 , 41, 951-9	2.7	2
107	<i>Edwardsiella tarda</i> evades serum killing by preventing complement activation via the alternative pathway. <i>Fish and Shellfish Immunology</i> , 2015 , 43, 325-9	4.3	35
106	CsCTL1, a teleost C-type lectin that promotes antibacterial and antiviral immune defense in a manner that depends on the conserved EPN motif. <i>Developmental and Comparative Immunology</i> , 2015 , 50, 69-77	3.2	63
105	Nuclear factor 45 of tongue sole (<i>Cynoglossus semilaevis</i>): evidence for functional differentiation between two isoforms in immune defense against viral and bacterial pathogens. <i>Developmental and Comparative Immunology</i> , 2014 , 42, 125-31	3.2	6
104	CsIFIT1, an interferon-induced protein with tetratricopeptide repeat, inhibits viral infection in tongue sole (<i>Cynoglossus semilaevis</i>). <i>Fish and Shellfish Immunology</i> , 2014 , 41, 231-7	4.3	18
103	Junctional adhesion molecule A of red drum (<i>Sciaenops ocellatus</i>): a possible immunomodulator and a target for bacterial immune evasion. <i>Veterinary Immunology and Immunopathology</i> , 2014 , 161, 99-107	4.0	1
102	Turbot (<i>Scophthalmus maximus</i>) hepcidin-1 and hepcidin-2 possess antimicrobial activity and promote resistance against bacterial and viral infection. <i>Fish and Shellfish Immunology</i> , 2014 , 38, 127-34	4.3	36
101	<i>Edwardsiella tarda</i> Hfq: impact on host infection and global protein expression. <i>Veterinary Research</i> , 2014 , 45, 23	3.8	29
100	Rock bream (<i>Oplegnathus fasciatus</i>) viperin is a virus-responsive protein that modulates innate immunity and promotes resistance against megalocytivirus infection. <i>Developmental and Comparative Immunology</i> , 2014 , 45, 35-42	3.2	29
99	NKLP27: a teleost NK-lysin peptide that modulates immune response, induces degradation of bacterial DNA, and inhibits bacterial and viral infection. <i>PLoS ONE</i> , 2014 , 9, e106543	3.7	36
98	Poly(I:C) induces antiviral immune responses in Japanese flounder (<i>Paralichthys olivaceus</i>) that require TLR3 and MDA5 and is negatively regulated by Myd88. <i>PLoS ONE</i> , 2014 , 9, e112918	3.7	58
97	In-depth profiling and analysis of host and viral microRNAs in Japanese flounder (<i>Paralichthys olivaceus</i>) infected with megalocytivirus reveal involvement of microRNAs in host-virus interaction in teleost fish. <i>BMC Genomics</i> , 2014 , 15, 878	4.5	64
96	<i>Streptococcus iniae</i> SF1: complete genome sequence, proteomic profile, and immunoprotective antigens. <i>PLoS ONE</i> , 2014 , 9, e91324	3.7	21
95	Sil: a <i>Streptococcus iniae</i> bacteriocin with dual role as an antimicrobial and an immunomodulator that inhibits innate immune response and promotes <i>S. iniae</i> infection. <i>PLoS ONE</i> , 2014 , 9, e96222	3.7	17

94	C7: a CpG oligodeoxynucleotide that induces protective immune response against megalocytivirus in Japanese flounder (<i>Paralichthys olivaceus</i>) via Toll-like receptor 9-mediated signaling pathway. <i>Developmental and Comparative Immunology</i> , 2014 , 44, 124-32	3.2	16
93	Antibacterial and antiviral properties of tongue sole (<i>Cynoglossus semilaevis</i>) high mobility group B2 protein are largely independent on the acidic C-terminal domain. <i>Fish and Shellfish Immunology</i> , 2014 , 37, 66-74	4.3	49
92	Quantitative real time RT-PCR study of pathogen-induced gene expression in rock bream (<i>Oplegnathus fasciatus</i>): internal controls for data normalization. <i>Marine Genomics</i> , 2014 , 15, 75-84	1.9	12
91	Identification and analysis of three virulence-associated TonB-dependent outer membrane receptors of <i>Pseudomonas fluorescens</i> . <i>Diseases of Aquatic Organisms</i> , 2014 , 110, 181-91	1.7	8
90	Immunological study of the outer membrane proteins of <i>Vibrio harveyi</i> : insights that link immunoprotectivity to interference with bacterial infection. <i>Fish and Shellfish Immunology</i> , 2013 , 35, 1293-300	4.3	23
89	Identification of normalization factors for quantitative real-time RT-PCR analysis of gene expression in Pacific abalone <i>Haliotis discus hannai</i> . <i>Chinese Journal of Oceanology and Limnology</i> , 2013 , 31, 421-430		10
88	A NK-lysin from <i>Cynoglossus semilaevis</i> enhances antimicrobial defense against bacterial and viral pathogens. <i>Developmental and Comparative Immunology</i> , 2013 , 40, 258-65	3.2	41
87	Comparative study of four flagellins of <i>Vibrio anguillarum</i> : vaccine potential and adjuvanticity. <i>Fish and Shellfish Immunology</i> , 2013 , 34, 514-20	4.3	13
86	Edwardsiella tarda Ivy, a lysozyme inhibitor that blocks the lytic effect of lysozyme and facilitates host infection in a manner that is dependent on the conserved cysteine residue. <i>Infection and Immunity</i> , 2013 , 81, 3527-33	3.7	24
85	SagE induces highly effective protective immunity against <i>Streptococcus iniae</i> mainly through an immunogenic domain in the extracellular region. <i>Acta Veterinaria Scandinavica</i> , 2013 , 55, 78	2	4
84	Characterization of a c-type lysozyme of <i>Scophthalmus maximus</i> : expression, activity, and antibacterial effect. <i>Fish and Shellfish Immunology</i> , 2013 , 34, 46-54	4.3	43
83	The megalocytivirus-induced protein CsMig1 enhances <i>Cynoglossus semilaevis</i> resistance against viral infection. <i>Veterinary Immunology and Immunopathology</i> , 2013 , 151, 173-9	2	3
82	Megalocytivirus-induced proteins of turbot (<i>Scophthalmus maximus</i>): identification and antiviral potential. <i>Journal of Proteomics</i> , 2013 , 91, 430-43	3.9	19
81	Macrophage migration inhibitory factor of <i>Sciaenops ocellatus</i> regulates immune cell trafficking and is involved in pathogen-induced immune response. <i>Developmental and Comparative Immunology</i> , 2013 , 40, 232-9	3.2	23
80	SmCCL19, a CC chemokine of turbot <i>Scophthalmus maximus</i> , induces leukocyte trafficking and promotes anti-viral and anti-bacterial defense. <i>Fish and Shellfish Immunology</i> , 2013 , 35, 1677-82	4.3	42
79	Selection of normalization factors for quantitative real time RT-PCR studies in Japanese flounder (<i>Paralichthys olivaceus</i>) and turbot (<i>Scophthalmus maximus</i>) under conditions of viral infection. <i>Veterinary Immunology and Immunopathology</i> , 2013 , 152, 303-16	2	41
78	The galectin-3-binding protein of <i>Cynoglossus semilaevis</i> is a secreted protein of the innate immune system that binds a wide range of bacteria and is involved in host phagocytosis. <i>Developmental and Comparative Immunology</i> , 2013 , 39, 399-408	3.2	30
77	Nuclear factor 45 of half smooth tongue sole <i>Cynoglossus semilaevis</i> : gene structure, expression profile, and immunoregulatory property. <i>Fish and Shellfish Immunology</i> , 2013 , 35, 972-8	4.3	1

76	The C-reactive protein of tongue sole <i>Cynoglossus semilaevis</i> is an acute phase protein that interacts with bacterial pathogens and stimulates the antibacterial activity of peripheral blood leukocytes. <i>Fish and Shellfish Immunology</i> , 2013 , 34, 623-31	4.3	19
75	First characterization of a teleost Epstein-Barr virus-induced gene 3 (EBI3) reveals a regulatory effect of EBI3 on the innate immune response of peripheral blood leukocytes. <i>Developmental and Comparative Immunology</i> , 2013 , 41, 514-22	3.2	20
74	Complete genome sequence and transcription profiles of the rock bream iridovirus RBIV-C1. <i>Diseases of Aquatic Organisms</i> , 2013 , 104, 203-14	1.7	33
73	The major fimbrial subunit protein of <i>Edwardsiella tarda</i> : vaccine potential, adjuvant effect, and involvement in host infection. <i>Fish and Shellfish Immunology</i> , 2013 , 35, 858-65	4.3	26
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66	Construction and analysis of experimental DNA vaccines against megalocytivirus. <i>Fish and Shellfish Immunology</i> , 2012 , 33, 1192-8	4.3	41
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60	Inv1: an <i>Edwardsiella tarda</i> invasin and a protective immunogen that is required for host infection. <i>Fish and Shellfish Immunology</i> , 2012 , 32, 586-92	4.3	13
59	Molecular characterization of <i>Cynoglossus semilaevis</i> CD28. <i>Fish and Shellfish Immunology</i> , 2012 , 32, 934-8	4.3	19

58	A divalent DNA vaccine based on Sia10 and OmpU induces cross protection against <i>Streptococcus iniae</i> and <i>Vibrio anguillarum</i> in Japanese flounder. <i>Fish and Shellfish Immunology</i> , 2012 , 32, 1216-22	4.3	29
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56	A TonB-dependent outer membrane receptor of <i>Pseudomonas fluorescens</i> : virulence and vaccine potential. <i>Archives of Microbiology</i> , 2012 , 194, 795-802	3	29
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38	The Rab1 GTPase of <i>Sciaenops ocellatus</i> modulates intracellular bacterial infection. <i>Fish and Shellfish Immunology</i> , 2011 , 31, 1005-12	4.3	14
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32	Analysis of <i>Edwardsiella tarda</i> DegP, a serine protease and a protective immunogen. <i>Fish and Shellfish Immunology</i> , 2010 , 28, 672-7	4.3	20
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