

Xiao-hua Zhang

List of Publications by Year in descending order

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156
papers

4,935
citations

117625

34
h-index

144013

57
g-index

160
all docs

160
docs citations

160
times ranked

4582
citing authors

#	ARTICLE	IF	CITATIONS
1	Distinctive signatures of pathogenic and antibiotic resistant potentials in the hadal microbiome. <i>Environmental Microbiomes</i> , 2022, 17, 19.	5.0	6
2	<i>Marinifilum caeruleilacunae</i> sp. nov., isolated from Yongle Blue Hole in the South China Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	1.7	5
3	Viable but nonculturable bacteria and their resuscitation: implications for cultivating uncultured marine microorganisms. <i>Marine Life Science and Technology</i> , 2021, 3, 189-203.	4.6	44
4	Bacterial Dimethylsulfoniopropionate Biosynthesis in the East China Sea. <i>Microorganisms</i> , 2021, 9, 657.	3.6	12
5	Cultivation of uncultured marine microorganisms. <i>Marine Life Science and Technology</i> , 2021, 3, 117-120.	4.6	23
6	Oxidation of trimethylamine to trimethylamine <i>N</i> -oxide facilitates high hydrostatic pressure tolerance in a generalist bacterial lineage. <i>Science Advances</i> , 2021, 7, .	10.3	17
7	What do we mean by viability in terms of "viable but nonculturable"™ cells?. <i>Environmental Microbiology Reports</i> , 2021, 13, 248-252.	2.4	8
8	A novel ATP dependent dimethylsulfoniopropionate lyase in bacteria that releases dimethyl sulfide and acryloyl-CoA. <i>ELife</i> , 2021, 10, .	6.0	38
9	Vertical diversity and association pattern of total, abundant and rare microbial communities in deep-sea sediments. <i>Molecular Ecology</i> , 2021, 30, 2800-2816.	3.9	41
10	Reply to: "Questions remain about the biolability of dissolved black carbon along the combustion continuum". <i>Nature Communications</i> , 2021, 12, 4282.	12.8	3
11	DiTing: A Pipeline to Infer and Compare Biogeochemical Pathways From Metagenomic and Metatranscriptomic Data. <i>Frontiers in Microbiology</i> , 2021, 12, 698286.	3.5	21
12	Succession of marine bacteria in response to <i>Ulva prolifera</i> -derived dissolved organic matter. <i>Environment International</i> , 2021, 155, 106687.	10.0	24
13	Spatiotemporal distribution of bacterial dimethylsulfoniopropionate producing and catabolic genes in the Changjiang Estuary. <i>Environmental Microbiology</i> , 2021, 23, 7073-7092.	3.8	11
14	Dimethylsulfoniopropionate Biosynthetic Bacteria in the Subseafloor Sediments of the South China Sea. <i>Frontiers in Microbiology</i> , 2021, 12, 731524.	3.5	4
15	Comparative Genomic Analysis of <i>Labrenzia aggregata</i> (Alphaproteobacteria) Strains Isolated From the Mariana Trench: Insights Into the Metabolic Potentials and Biogeochemical Functions. <i>Frontiers in Microbiology</i> , 2021, 12, 770370.	3.5	6
16	Vertical variation in <i>Vibrio</i> community composition in Sansha Yongle Blue Hole and its ability to degrade macromolecules. <i>Marine Life Science and Technology</i> , 2020, 2, 60-72.	4.6	32
17	Comparative genomic and metabolic analysis of manganese-oxidizing mechanisms in <i>Celeribacter manganoxidans</i> DY25T: Its adaptation to the environment of polymetallic nodules. <i>Genomics</i> , 2020, 112, 2080-2091.	2.9	18
18	Dissolved black carbon is not likely a significant refractory organic carbon pool in rivers and oceans. <i>Nature Communications</i> , 2020, 11, 5051.	12.8	53

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19	Carbon cycling in the deep Mariana Trench in the western north Pacific Ocean: Insights from radiocarbon proxy data. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2020, 164, 103370.	1.4	12
20	Bacteria are important dimethylsulfoniopropionate producers in marine aphotic and high-pressure environments. <i>Nature Communications</i> , 2020, 11, 4658.	12.8	62
21	Insights into the Vertical Stratification of Microbial Ecological Roles across the Deepest Seawater Column on Earth. <i>Microorganisms</i> , 2020, 8, 1309.	3.6	18
22	Methane production in oxic seawater of the western North Pacific and its marginal seas. <i>Limnology and Oceanography</i> , 2020, 65, 2352-2365.	3.1	19
23	Ancestral niche separation and evolutionary rate differentiation between sister marine flavobacteria lineages. <i>Environmental Microbiology</i> , 2020, 22, 3234-3247.	3.8	8
24	Novel insights into the Thaumarchaeota in the deepest oceans: their metabolism and potential adaptation mechanisms. <i>Microbiome</i> , 2020, 8, 78.	11.1	47
25	Two Highly Similar Chitinases from Marine <i>Vibrio</i> Species have Different Enzymatic Properties. <i>Marine Drugs</i> , 2020, 18, 139.	4.6	12
26	Spatiotemporal dynamics of the archaeal community in coastal sediments: assembly process and co-occurrence relationship. <i>ISME Journal</i> , 2020, 14, 1463-1478.	9.8	153
27	Spatiotemporal dynamics of the total and active <i>Vibrio</i> spp. populations throughout the Changjiang estuary in China. <i>Environmental Microbiology</i> , 2020, 22, 4438-4455.	3.8	22
28	Metagenomic Insights Into the Cycling of Dimethylsulfoniopropionate and Related Molecules in the Eastern China Marginal Seas. <i>Frontiers in Microbiology</i> , 2020, 11, 157.	3.5	22
29	Carbon Cycling in the World's Deepest Blue Hole. <i>Journal of Geophysical Research G: Biogeosciences</i> , 2020, 125, e2019JG005307.	3.0	17
30	Enhanced Activity against Multidrug-Resistant Bacteria through Coapplication of an Analogue of Tachyplesin I and an Inhibitor of the QseC/B Signaling Pathway. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 3475-3484.	6.4	20
31	Diversity of culturable heterotrophic bacteria from the Mariana Trench and their ability to degrade macromolecules. <i>Marine Life Science and Technology</i> , 2020, 2, 181-193.	4.6	28
32	<i>Vibrio harveyi</i> : a serious pathogen of fish and invertebrates in mariculture. <i>Marine Life Science and Technology</i> , 2020, 2, 231-245.	4.6	147
33	DMSP-Producing Bacteria Are More Abundant in the Surface Microlayer than Subsurface Seawater of the East China Sea. <i>Microbial Ecology</i> , 2020, 80, 350-365.	2.8	28
34	Metagenomic Insights Into the Microbial Assemblage Capable of Quorum Sensing and Quorum Quenching in Particulate Organic Matter in the Yellow Sea. <i>Frontiers in Microbiology</i> , 2020, 11, 602010.	3.5	7
35	Shift and Metabolic Potentials of Microbial Eukaryotic Communities Across the Full Depths of the Mariana Trench. <i>Frontiers in Microbiology</i> , 2020, 11, 603692.	3.5	5
36	<i>Vibrio ouci</i> sp. nov. and <i>Vibrio aquaticus</i> sp. nov., two marine bacteria isolated from the East China Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 172-179.	1.7	14

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37	<i>Glycocalyx profundus</i> sp. nov., a marine bacterium isolated from seawater of the Mariana Trench. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 814-819.	1.7	12
38	<i>Vibrio sinensis</i> sp. nov. and <i>Vibrio viridaestus</i> sp. nov., two marine bacteria isolated from the East China Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 889-896.	1.7	15
39	<i>Puteibacter caeruleilacunae</i> gen. nov., sp. nov., a facultatively anaerobic bacterium isolated from Yongle Blue Hole in the South China Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 1623-1629.	1.7	9
40	<i>Marinobacter salinexigens</i> sp. nov., a marine bacterium isolated from hadal seawater of the Mariana Trench. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 3794-3800.	1.7	12
41	Bacteria are important dimethylsulfoniopropionate producers in coastal sediments. <i>Nature Microbiology</i> , 2019, 4, 1815-1825.	13.3	67
42	<i>Corallincola luteus</i> sp. nov., a marine bacterium isolated from surface sediment of Bohai Sea of China. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 1691-1697.	1.7	0
43	Biogenic production of DMSP and its degradation to DMS—their roles in the global sulfur cycle. <i>Science China Life Sciences</i> , 2019, 62, 1296-1319.	4.9	68
44	Microbial assembly, interaction, functioning, activity and diversification: a review derived from community compositional data. <i>Marine Life Science and Technology</i> , 2019, 1, 112-128.	4.6	104
45	A novel heterologous expression strategy for the quorum-quenching enzyme MomL in <i>Lysobacter</i> enzymogenes to the inhibit pathogenicity of <i>Pectobacterium</i> . <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 8889-8898.	3.6	3
46	Interspecies and Intraspecies Signals Synergistically Regulate <i>Lysobacter</i> enzymogenes Twitching Motility. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	15
47	Effect of ZnO on Morphology and Breakdown Characteristics of Polyethylene. , 2019, , .		0
48	2D few-layer iron phosphosulfide: a self-buffer heterophase structure induced by irreversible breakage of P—S bonds for high-performance lithium/sodium storage. <i>Journal of Materials Chemistry A</i> , 2019, 7, 1529-1538.	10.3	48
49	Mechanistic insight into 3-methylmercaptopropionate metabolism and kinetical regulation of demethylation pathway in marine dimethylsulfoniopropionate-catabolizing bacteria. <i>Molecular Microbiology</i> , 2019, 111, 1057-1073.	2.5	18
50	Activity Improvement and Vital Amino Acid Identification on the Marine-Derived Quorum Quenching Enzyme MomL by Protein Engineering. <i>Marine Drugs</i> , 2019, 17, 300.	4.6	15
51	Indole Reverses Intrinsic Antibiotic Resistance by Activating a Novel Dual-Function Importer. <i>MBio</i> , 2019, 10, .	4.1	31
52	Dynamic mechanism of breakdown in polypropylene-based nano-dielectric. <i>AIP Advances</i> , 2019, 9, 015135.	1.3	8
53	Spatiotemporal Dynamics of Free-Living and Particle-Associated <i>Vibrio</i> Communities in the Northern Chinese Marginal Seas. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	46
54	Spatial Heterogeneity of <i>Vibrio</i> spp. in Sediments of Chinese Marginal Seas. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	18

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55	Proliferation of hydrocarbon-degrading microbes at the bottom of the Mariana Trench. <i>Microbiome</i> , 2019, 7, 47.	11.1	128
56	Heterologous Expression of the Marine-Derived Quorum Quenching Enzyme MomL Can Expand the Antibacterial Spectrum of <i>Bacillus brevis</i> . <i>Marine Drugs</i> , 2019, 17, 128.	4.6	10
57	Role of RpoN from <i>Labrenzia aggregata</i> LZB033 (<i>Rhodobacteraceae</i>) in Formation of Flagella and Biofilms, Motility, and Environmental Adaptation. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	19
58	Carbohydrate catabolic capability of a <i>Flavobacteriia</i> bacterium isolated from hadal water. <i>Systematic and Applied Microbiology</i> , 2019, 42, 263-274.	2.8	25
59	The Mechanisms and Applications of Quorum Sensing (QS) and Quorum Quenching (QQ). <i>Journal of Ocean University of China</i> , 2019, 18, 1427-1442.	1.2	34
60	Marine life science and technology (MLST): a new journal highlighting all aspects of marine biology and biotechnology research. <i>Marine Life Science and Technology</i> , 2019, 1, 1-3.	4.6	2
61	Silicon promotes seedling growth and alters endogenous IAA, GA ₃ and ABA concentrations in <i>Glycyrrhiza uralensis</i> under 100 mM NaCl stress. <i>Journal of Horticultural Science and Biotechnology</i> , 2019, 94, 87-93.	1.9	31
62	<i>Thalassococcus profundus</i> sp. nov., a marine bacterium isolated from deep seawater of the Okinawa Trough. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 920-925.	1.7	7
63	<i>Winogradskyella ouciana</i> sp. nov., isolated from the hadal seawater of the Mariana Trench. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 71, .	1.7	9
64	Characterization and overexpression of a glycosyl hydrolase family 16 beta-agarase YM01-1 from marine bacterium <i>Catenovulum agarivorans</i> YM01 T. <i>Protein Expression and Purification</i> , 2018, 143, 1-8.	1.3	20
65	A Culture-Dependent Method for the Identification of Quorum Quenching Enzymes of Microbial Origin. <i>Methods in Molecular Biology</i> , 2018, 1673, 297-309.	0.9	4
66	Novel Insights Into Bacterial Dimethylsulfoniopropionate Catabolism in the East China Sea. <i>Frontiers in Microbiology</i> , 2018, 9, 3206.	3.5	35
67	A nearly uniform distributional pattern of heterotrophic bacteria in the Mariana Trench interior. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2018, 142, 116-126.	1.4	27
68	Characterization of a Novel N-Acylhomoserine Lactonase RmmL from <i>Ruegeria mobilis</i> YJ3. <i>Marine Drugs</i> , 2018, 16, 370.	4.6	13
69	Distribution patterns of ammonia-oxidizing archaea and bacteria in sediments of the eastern China marginal seas. <i>Systematic and Applied Microbiology</i> , 2018, 41, 658-668.	2.8	19
70	Sediment Depth-Dependent Spatial Variations of Bacterial Communities in Mud Deposits of the Eastern China Marginal Seas. <i>Frontiers in Microbiology</i> , 2018, 9, 1128.	3.5	32
71	Comparative genomic analysis reveals the evolution and environmental adaptation strategies of vibrios. <i>BMC Genomics</i> , 2018, 19, 135.	2.8	71
72	Significance of <i>Vibrio</i> species in the marine organic carbon cycle—A review. <i>Science China Earth Sciences</i> , 2018, 61, 1357-1368.	5.2	99

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73	<i>Flavobacterium ovatum</i> sp. nov., a marine bacterium isolated from an Antarctic intertidal sandy beach. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 795-800.	1.7	15
74	<i>Bacillus alkalitolerans</i> sp. nov., isolated from marine sediment near a hydrothermal vent. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1184-1189.	1.7	8
75	<i>Euzebya rosea</i> sp. nov., a rare actinobacterium isolated from the East China Sea and analysis of two genome sequences in the genus <i>Euzebya</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2900-2905.	1.7	21
76	<i>Abyssibacter profundus</i> gen. nov., sp. nov., a marine bacterium isolated from seawater of the Mariana Trench. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3424-3429.	1.7	13
77	<i>Marinifilum breve</i> sp. nov., a marine bacterium isolated from the Yongle Blue Hole in the South China Sea and emended description of the genus <i>Marinifilum</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3540-3545.	1.7	13
78	Bacterial community structure in intertidal sediments of Fildes Peninsula, maritime Antarctica. <i>Polar Biology</i> , 2017, 40, 339-349.	1.2	28
79	Dimethylsulfoniopropionate biosynthesis in marine bacteria and identification of the key gene in this process. <i>Nature Microbiology</i> , 2017, 2, 17009.	13.3	222
80	Early diagenesis and authigenic mineral formation in mobile muds of the Changjiang Estuary and adjacent shelf. <i>Journal of Marine Systems</i> , 2017, 172, 64-74.	2.1	26
81	Diversity and Abundance of the Denitrifying Microbiota in the Sediment of Eastern China Marginal Seas and the Impact of Environmental Factors. <i>Microbial Ecology</i> , 2017, 73, 602-615.	2.8	14
82	A novel stress response mechanism, triggered by indole, involved in quorum quenching enzyme MomL and iron-sulfur cluster in <i>Muricauda olearia</i> Th120. <i>Scientific Reports</i> , 2017, 7, 4252.	3.3	24
83	Bacterial Community Associated with Healthy and Diseased Pacific White Shrimp (<i>Litopenaeus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 8, 1362.	3.5	105
84	<i>Photobacterium alginatilyticum</i> sp. nov., a marine bacterium isolated from bottom seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1912-1917.	1.7	23
85	<i>Roseibium sediminis</i> sp. nov., isolated from sea surface sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 2862-2867.	1.7	13
86	<i>Polaribacter pacificus</i> sp. nov., isolated from a deep-sea polymetallic nodule from the Eastern Pacific Ocean. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 3203-3208.	1.7	7
87	<i>Croceitalea marina</i> sp. nov., isolated from marine particles of Yellow Sea, and emended description of the genera <i>Croceitalea</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 4253-4259.	1.7	9
88	PfmA, a novel quorum-quenching N-acylhomoserine lactone acylase from <i>Pseudoalteromonas flavipulchra</i> . <i>Microbiology (United Kingdom)</i> , 2017, 163, 1389-1398.	1.8	24
89	Reclassification of <i>Xuhuaishuia manganoxidans</i> Wang et al. 2015 as a later heterotypic synonym of <i>Brevirhabdus pacifica</i> Wu et al. 2015 and emendation of the species description. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 3095-3098.	1.7	3
90	Diversity, Abundance, and Niche Differentiation of Ammonia-Oxidizing Prokaryotes in Mud Deposits of the Eastern China Marginal Seas. <i>Frontiers in Microbiology</i> , 2016, 7, 137.	3.5	40

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91	Marine Microbiological Enzymes: Studies with Multiple Strategies and Prospects. <i>Marine Drugs</i> , 2016, 14, 171.	4.6	30
92	LaaA, a novel high-active alkalophilic alpha-amylase from deep-sea bacterium <i>Luteimonas abyssi</i> XH031T. <i>Enzyme and Microbial Technology</i> , 2016, 90, 83-92.	3.2	15
93	Degradation properties of various macromolecules of cultivable psychrophilic bacteria from the deep-sea water of the South Pacific Gyre. <i>Extremophiles</i> , 2016, 20, 663-671.	2.3	18
94	Comparison of cultivable bacterial communities associated with Pacific white shrimp (<i>Litopenaeus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	3.5	93
95	<i>Aquimarina hainanensis</i> sp. nov., isolated from diseased Pacific white shrimp <i>Litopenaeus vannamei</i> larvae. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 70-75.	1.7	18
96	<i>Enterovibrio pacificus</i> sp. nov., isolated from seawater, and emended descriptions of <i>Enterovibrio coralii</i> and the genus <i>Enterovibrio</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 319-325.	1.7	11
97	<i>Arcobacter pacificus</i> sp. nov., isolated from seawater of the South Pacific Gyre. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 542-547.	1.7	31
98	<i>Xuhuaishuia manganoxidans</i> gen. nov., sp. nov., a manganese-oxidizing bacterium isolated from deep-sea sediments from the Pacific Polymetallic Nodule Province. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 1521-1526.	1.7	23
99	<i>Marinicella pacifica</i> sp. nov., isolated from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 2313-2318.	1.7	17
100	<i>Allohadella marinimesophila</i> gen. nov., sp. nov., isolated from seawater and reclassification of <i>Hahella antarctica</i> as <i>Allohadella antarctica</i> comb. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 3207-3213.	1.7	12
101	<i>Bacterioplanoides pacificum</i> gen. nov., sp. nov., isolated from seawater of South Pacific Gyre. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 5010-5015.	1.7	15
102	Genomic insight into <i>Aquimarina longa</i> SW024T: its ultra-oligotrophic adapting mechanisms and biogeochemical functions. <i>BMC Genomics</i> , 2015, 16, 772.	2.8	13
103	Genomic analysis of <i>Luteimonas abyssi</i> XH031T: insights into its adaption to the subseafloor environment of South Pacific Gyre and ecological role in biogeochemical cycle. <i>BMC Genomics</i> , 2015, 16, 1092.	2.8	22
104	Effect of silicon on seed germination and the physiological characteristics of <i>Glycyrrhizauralensis</i> under different levels of salinity. <i>Journal of Horticultural Science and Biotechnology</i> , 2015, 90, 439-443.	1.9	36
105	<i>Loktanella sediminum</i> sp. nov., isolated from marine surface sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 686-691.	1.7	17
106	<i>Deinococcus antarcticus</i> sp. nov., isolated from soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 331-335.	1.7	20
107	Phylogenetic diversity and biological activities of marine actinomycetes isolated from sediments of the Yellow Sea Cold Water Mass, China. <i>Marine Biology Research</i> , 2015, 11, 551-560.	0.7	1
108	<i>Ichthyenterobacterium magnum</i> gen. nov., sp. nov., a member of the family Flavobacteriaceae isolated from olive flounder (<i>Paralichthys olivaceus</i>). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1186-1192.	1.7	17

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109	Genome analysis of <i>Flaviramulus ichthyoenteri</i> Th78T in the family Flavobacteriaceae: insights into its quorum quenching property and potential roles in fish intestine. <i>BMC Genomics</i> , 2015, 16, 38.	2.8	22
110	<i>Flavirhabdus iliipiscaria</i> gen. nov., sp. nov., isolated from intestine of flounder (<i>Paralichthys</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 707 Tc <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1347-1353.	1.7	22
111	Phylogenetic shifts of bacterioplankton community composition along the Pearl Estuary: the potential impact of hypoxia and nutrients. <i>Frontiers in Microbiology</i> , 2015, 6, 64.	3.5	135
112	Bacterial and Archaeal Communities in Sediments of the North Chinese Marginal Seas. <i>Microbial Ecology</i> , 2015, 70, 105-117.	2.8	133
113	<i>Salipiger nanhaiensis</i> sp. nov., a bacterium isolated from deep sea water. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1122-1126.	1.7	18
114	Quorum sensing in marine snow and its possible influence on production of extracellular hydrolytic enzymes in marine snow bacterium <i>Pantoea ananatis</i> B9. <i>FEMS Microbiology Ecology</i> , 2015, 91, 1-13.	2.7	65
115	Shift of anammox bacterial community structure along the Pearl Estuary and the impact of environmental factors. <i>Journal of Geophysical Research: Oceans</i> , 2015, 120, 2869-2883.	2.6	28
116	Studies on bacterial pathogens isolated from diseased torafugu (<i>Takifugu rubripes</i>) cultured in marine industrial recirculation aquaculture system in Shandong Province, China. <i>Aquaculture Research</i> , 2015, 46, 736-744.	1.8	11
117	MomL, a Novel Marine-Derived <i>N</i> -Acyl Homoserine Lactonase from <i>Muricauda olearia</i> . <i>Applied and Environmental Microbiology</i> , 2015, 81, 774-782.	3.1	104
118	<i>Dokdonia pacifica</i> sp. nov., isolated from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 2222-2226.	1.7	13
119	<i>Leucothrix pacifica</i> sp. nov., isolated from seawater, and emended description of the genus <i>Leucothrix</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 2397-2402.	1.7	10
120	<i>Aureibacillus halotolerans</i> gen. nov., sp. nov., isolated from marine sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 3950-3958.	1.7	11
121	<i>Muricauda pacifica</i> sp. nov., isolated from seawater of the South Pacific Gyre. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 4087-4092.	1.7	30
122	Overexpression and Characterization of a Novel Thermostable β -Agarase YM01-3, from Marine Bacterium <i>Catenovulum agarivorans</i> YM01T. <i>Marine Drugs</i> , 2014, 12, 2731-2747.	4.6	42
123	Quorum Quenching Agents: Resources for Antivirulence Therapy. <i>Marine Drugs</i> , 2014, 12, 3245-3282.	4.6	141
124	The influence of proteolytic and cytolytic enzymes on starch degradation during mashing. <i>Journal of the Institute of Brewing</i> , 2014, 120, n/a-n/a.	2.3	8
125	<i>Aquimarina pacifica</i> sp. nov., isolated from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 1991-1997.	1.7	46
126	Spatial distribution patterns of benthic microbial communities along the Pearl Estuary, China. <i>Systematic and Applied Microbiology</i> , 2014, 37, 578-589.	2.8	89

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127	<i>Roseivivax marinus</i> sp. nov., isolated from deep water. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 2540-2544.	1.7	17
128	Description of <i>Thalassotalea piscium</i> gen. nov., sp. nov., isolated from flounder (<i>Paralichthys</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 712 <i>Thalassotalea</i> gen. nov. and emended description of the genus <i>Thalassomonas</i> . International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1223-1228.	1.7	46
129	<i>Luteococcus sediminum</i> sp. nov., isolated from deep seafloor sediment of the South Pacific Gyre. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 2522-2527.	1.7	14
130	A mutation in <i>rcsB</i> , a gene encoding the core component of the Rcs cascade, enhances the virulence of <i>Edwardsiella tarda</i> . Research in Microbiology, 2014, 165, 226-232.	2.1	9
131	<i>Achromobacter sediminum</i> sp. nov., isolated from deep seafloor sediment of South Pacific Gyre. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 2244-2249.	1.7	21
132	<i>Aquimarina megaterium</i> sp. nov., isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 122-127.	1.7	34
133	Shifts in archaeoplankton community structure along ecological gradients of Pearl Estuary. FEMS Microbiology Ecology, 2014, 90, n/a-n/a.	2.7	41
134	<i>Luteimonas abyssi</i> sp. nov., isolated from deep-sea sediment. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 668-674.	1.7	36
135	<i>Nocardioides pacificus</i> sp. nov., isolated from deep sub-seafloor sediment. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 2217-2222.	1.7	16
136	<i>Oceanobacillus pacificus</i> sp. nov., isolated from a deep-sea sediment. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1278-1283.	1.7	26
137	<i>Flaviramulus ichthyoenteri</i> sp. nov., an N-acylhomoserine lactone-degrading bacterium isolated from the intestine of a flounder (<i>Paralichthys olivaceus</i>), and emended descriptions of the genus <i>Flaviramulus</i> and <i>Flaviramulus basaltis</i> . International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 4477-4483.	1.7	33
138	Genome analysis of <i>Pseudoalteromonas flavipulchra</i> JG1 reveals various survival advantages in marine environment. BMC Genomics, 2013, 14, 707.	2.8	23
139	<i>Ferrimonas sediminum</i> sp. nov., isolated from coastal sediment of an amphioxus breeding zone. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 977-981.	1.7	13
140	<i>Aquimarina longa</i> sp. nov., isolated from seawater, and emended description of <i>Aquimarina muelleri</i> . International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 1235-1240.	1.7	42
141	Evaluation of a new high-throughput method for identifying quorum quenching bacteria. Scientific Reports, 2013, 3, 2935.	3.3	66
142	Spatial Variations in Microbial Community Composition in Surface Seawater from the Ultra-Oligotrophic Center to Rim of the South Pacific Gyre. PLoS ONE, 2013, 8, e55148.	2.5	76
143	Purification and characterization of antibacterial compounds of <i>Pseudoalteromonas flavipulchra</i> JG1. Microbiology (United Kingdom), 2012, 158, 835-842.	1.8	47
144	<i>Salinicoccus qingdaonensis</i> sp. nov., isolated from coastal seawater during a bloom of green algae. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 545-549.	1.7	14

#	ARTICLE	IF	CITATIONS
145	<i>Myroides phaeus</i> sp. nov., isolated from human saliva, and emended descriptions of the genus <i>Myroides</i> and the species <i>Myroides profundus</i> Zhang et al. 2009 and <i>Myroides marinus</i> Cho et al. 2011. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 770-775.	1.7	29
146	<i>Lentibacter algarum</i> gen. nov., sp. nov., isolated from coastal water during a massive green algae bloom. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 1042-1047.	1.7	23
147	<i>Salinactinospora qingdaonensis</i> gen. nov., sp. nov., a halophilic actinomycete isolated from a salt pond. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 954-959.	1.7	25
148	<i>Huaishuia halophila</i> gen. nov., sp. nov., isolated from coastal seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 223-228.	1.7	18
149	Gel microbead cultivation with a subenrichment procedure can yield better bacterial cultivability from a seawater sample than standard plating method. <i>Journal of Ocean University of China</i> , 2012, 11, 45-51.	1.2	12
150	<i>Catenovulum agarivorans</i> gen. nov., sp. nov., a peritrichously flagellated, chain-forming, agar-hydrolysing gammaproteobacterium from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2866-2873.	1.7	34
151	<i>Spinactinospora alkalitolerans</i> gen. nov., sp. nov., an actinomycete isolated from marine sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2805-2810.	1.7	28
152	The linear and nonlinear optical effects of white light. <i>Science in China Series G: Physics, Mechanics and Astronomy</i> , 2009, 52, 649-664.	0.2	1
153	Retention of Virulence in a Viable but Nonculturable <i>Edwardsiella tarda</i> Isolate. <i>Applied and Environmental Microbiology</i> , 2007, 73, 1349-1354.	3.1	98
154	Antibacterial activity of gallic acid from the flowers of <i>Rosa chinensis</i> Jacq. against fish pathogens. <i>Aquaculture Research</i> , 2007, 38, 1110-1112.	1.8	19
155	Characterization and resuscitation of viable but nonculturable <i>Vibrio alginolyticus</i> VIB283. <i>Archives of Microbiology</i> , 2007, 188, 283-288.	2.2	45
156	MomL inhibits bacterial antibiotic resistance through the starvation stringent response pathway. , 0, , .		1