Wei-Jia Zhang

List of Publications by Year in descending order

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		687363	580821
26	652	13	25
papers	citations	h-index	g-index
26	26	26	600
26	26	26	602
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Proliferation of hydrocarbon-degrading microbes at the bottom of the Mariana Trench. Microbiome, 2019, 7, 47.	11.1	128
2	Comparative genomic analysis provides insights into the evolution and niche adaptation of marine <i><scp>M</scp>agnetospira</i> sp. <scp>QH</scp> â€2 strain. Environmental Microbiology, 2014, 16, 525-544.	3.8	66
3	A novel genus of multicellular magnetotactic prokaryotes from the Yellow Sea. Environmental Microbiology, 2012, 14, 405-413.	3.8	64
4	The chimeric nature of the genomes of marine magnetotactic coccoidâ€ovoid bacteria defines a novel group of <scp><i>P</i></scp> <i>roteobacteria</i> . Environmental Microbiology, 2017, 19, 1103-1119.	3.8	60
5	Swimming behaviour and magnetotaxis function of the marine bacterium strain <scp>MO</scp> â€1. Environmental Microbiology Reports, 2014, 6, 14-20.	2.4	34
6	High Hydrostatic Pressure Inducible Trimethylamine N-Oxide Reductase Improves the Pressure Tolerance of Piezosensitive Bacteria Vibrio fluvialis. Frontiers in Microbiology, 2017, 8, 2646.	3 . 5	33
7	Architecture of a flagellar apparatus in the fast-swimming magnetotactic bacterium MO-1. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20643-20648.	7.1	29
8	Bacterial community structure and novel species of magnetotactic bacteria in sediments from a seamount in the Mariana volcanic arc. Scientific Reports, 2017, 7, 17964.	3.3	29
9	Complex Spatial Organization and Flagellin Composition of Flagellar Propeller from Marine Magnetotactic Ovoid Strain MO-1. Journal of Molecular Biology, 2012, 416, 558-570.	4.2	27
10	Configuration of redox gradient determines magnetotactic polarity of the marine bacteria MOâ€1. Environmental Microbiology Reports, 2010, 2, 646-650.	2.4	25
11	Ultrastructure of ellipsoidal magnetotactic multicellular prokaryotes depicts their complex assemblage and cellular polarity in the context of magnetotaxis. Environmental Microbiology, 2017, 19, 2151-2163.	3.8	22
12	Calcium ionâ€mediated assembly and function of glycosylated flagellar sheath of marine magnetotactic bacterium. Molecular Microbiology, 2010, 78, 1304-1312.	2.5	19
13	Genomic and physiological analysis reveals versatile metabolic capacity of deep-sea Photobacterium phosphoreum ANT-2200. Extremophiles, 2016, 20, 301-310.	2.3	18
14	Flagella and Swimming Behavior of Marine Magnetotactic Bacteria. Biomolecules, 2020, 10, 460.	4.0	14
15	Analysis of the Antigenic Properties of Membrane Proteins of Mycobacterium tuberculosis. Scientific Reports, 2019, 9, 3042.	3.3	13
16	Complete genome sequence of Shewanella benthica DB21MT-2, an obligate piezophilic bacterium isolated from the deepest Mariana Trench sediment. Marine Genomics, 2019, 44, 52-56.	1.1	12
17	Crassaminicella thermophila sp. nov., a moderately thermophilic bacterium isolated from a deep-sea hydrothermal vent chimney and emended description of the genus Crassaminicella. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	12
18	Pressure-Regulated Gene Expression and Enzymatic Activity of the Two Periplasmic Nitrate Reductases in the Deep-Sea Bacterium Shewanella piezotolerans WP3. Frontiers in Microbiology, 2018, 9, 3173.	3. 5	9

#	Article	IF	CITATIONS
19	Thermococcus aciditolerans sp. nov., a piezotolerant, hyperthermophilic archaeon isolated from a deep-sea hydrothermal vent chimney in the Southwest Indian Ridge. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	9
20	Measurement of Free-Swimming Motility and Magnetotactic Behavior of Magnetococcus massalia Strain MO-1. Methods in Molecular Biology, 2017, 1593, 305-320.	0.9	7
21	Resazurin as an indicator of reducing capacity for analyzing the physiologic status of deep-sea bacterium Photobacterium phosphoreum ANT-2200. Journal of Oceanology and Limnology, 2021, 39, 297-305.	1.3	5
22	Genome analysis of Crassaminicella sp. SY095, an anaerobic mesophilic marine bacterium isolated from a deep-sea hydrothermal vent on the Southwest Indian Ridge. Marine Genomics, 2020, 52, 100733.	1.1	4
23	Comparative genomic analysis of obligately piezophilic Moritella yayanosii DB21MT-5 reveals bacterial adaptation to the Challenger Deep, Mariana Trench. Microbial Genomics, 2021, 7, .	2.0	4
24	Metagenomic analysis reveals wide distribution of phototrophic bacteria in hydrothermal vents on the ultraslow-spreading Southwest Indian Ridge. Marine Life Science and Technology, 2022, 4, 255-267.	4.6	4
25	Distinct influence of trimethylamine N-oxide and high hydrostatic pressure on community structure and culturable deep-sea bacteria. Journal of Oceanology and Limnology, 2020, 38, 364-377.	1.3	3
26	Complete genome sequence of Crassaminicella sp. 143-21,isolated from a deep-sea hydrothermal vent. Marine Genomics, 2021, 62, 100899.	1.1	2