

# Zongze Shao

## List of Publications by Year in descending order

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195  
papers

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87723

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203  
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203  
docs citations

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times ranked

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#	ARTICLE	IF	CITATIONS
1	Acremolin D, a new acremolin alkaloid from the deep-sea sediment derived <i>Aspergillus sydowii</i> fungus. <i>Natural Product Research</i> , 2022, 36, 4936-4942.	1.0	8
2	Structure elucidation of a novel cyclic tripeptide from the marine-derived fungus <i>Aspergillus ochraceopetaliformis</i> DSW-2. <i>Natural Product Research</i> , 2022, 36, 3572-3578.	1.0	4
3	Genome-resolved evidence for functionally redundant communities and novel nitrogen fixers in the deyin-1 hydrothermal field, Mid-Atlantic Ridge. <i>Microbiome</i> , 2022, 10, 8.	4.9	5
4	Volatile Organic Compounds from <i>Bacillus aryabhatai</i> MCCC 1K02966 with Multiple Modes against <i>Meloidogyne incognita</i> . <i>Molecules</i> , 2022, 27, 103.	1.7	13
5	Core microbiome involved in nitrite removal in shrimp culture ponds. <i>Aquaculture Research</i> , 2022, 53, 1663-1675.	0.9	10
6	Characterization of a bioactive meroterpenoid isolated from the marine-derived fungus <i>Talaromyces</i> sp.. <i>Applied Microbiology and Biotechnology</i> , 2022, 106, 2927-2935.	1.7	11
7	Complete genome sequence of <i>Thermosulfurimonas marina</i> SU872T, an anaerobic thermophilic chemolithoautotrophic bacterium isolated from a shallow marine hydrothermal vent. <i>Marine Genomics</i> , 2021, 55, 100800.	0.4	2
8	A novel SAR324 bacterium associated with abalone, <i>Haliotis diversicolor</i> . <i>Aquaculture Research</i> , 2021, 52, 1945-1953.	0.9	0
9	Elemental sulfur reduction by a deep-sea hydrothermal vent <i>Campylobacterium</i> <i>Sulfurimonas</i> sp. <i>Environmental Microbiology</i> , 2021, 23, 965-979.	1.8	17
10	Microorganisms from deep-sea hydrothermal vents. <i>Marine Life Science and Technology</i> , 2021, 3, 204-230.	1.8	34
11	Diversity and Antiaflatoxic Activities of Culturable Filamentous Fungi from Deep-Sea Sediments of the South Atlantic Ocean. <i>Mycobiology</i> , 2021, 49, 151-160.	0.6	2
12	Complete genome sequence of a denitrifying bacterium <i>Halomonas</i> sp. SS10-MC5 isolated from hydrothermal vent of Indian Ocean. <i>Marine Genomics</i> , 2021, 58, 100849.	0.4	3
13	<i>Alcanivorax profundimaris</i> sp. nov., a Novel Marine Hydrocarbonoclastic Bacterium Isolated from Seawater and Deep-Sea Sediment. <i>Current Microbiology</i> , 2021, 78, 1053-1060.	1.0	14
14	Characterization of <i>Sulfurimonas hydrogeniphila</i> sp. nov., a Novel Bacterium Predominant in Deep-Sea Hydrothermal Vents and Comparative Genomic Analyses of the Genus <i>Sulfurimonas</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 626705.	1.5	18
15	<i>Sulfurimonas sediminis</i> sp. nov., a novel hydrogen- and sulfur-oxidizing chemolithoautotroph isolated from a hydrothermal vent at the Longqi system, southwestern Indian ocean. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 813-822.	0.7	8
16	Anti-Food Allergic Compounds from <i>Penicillium griseofulvum</i> MCCC 3A00225, a Deep-Sea-Derived Fungus. <i>Marine Drugs</i> , 2021, 19, 224.	2.2	12
17	<i>Croceimicrobium hydrocarbonivorans</i> gen. nov., sp. nov., a novel marine bacterium isolated from a bacterial consortium that degrades polyethylene terephthalate. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	10
18	<i>Halomonas diversa</i> sp. nov., isolated from deep-sea sediment of the Pacific Ocean. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	12

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19	<i>Bacillus pumilus</i> Group Comparative Genomics: Toward Pangenome Features, Diversity, and Marine Environmental Adaptation. <i>Frontiers in Microbiology</i> , 2021, 12, 571212.	1.5	9
20	<i>Thermococcus henrietii</i> sp. nov., a novel extreme thermophilic and piezophilic sulfur-reducing archaeon isolated from a deep-sea hydrothermal chimney. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	8
21	<i>Thermosipho ferrireducens</i> sp.nov., an anaerobic thermophilic iron(III)-reducing bacterium isolated from a deep-sea hydrothermal sulfide deposits. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	5
22	<i>Mesonia hitae</i> sp. nov., isolated from the seawater of the South Atlantic Ocean. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	5
23	Characterization of two novel psychrophilic and piezotolerant strains, <i>Shewanella psychropiezotolerans</i> sp. nov. and <i>Shewanella eurypsychrophilus</i> sp. nov, adapted to an extreme deep-sea environment. <i>Systematic and Applied Microbiology</i> , 2021, 44, 126266.	1.2	9
24	<i>Solimonas marina</i> sp. nov., isolated from deep seawater of the Pacific Ocean. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	7
25	An Intracellular Sensing and Signal Transduction System That Regulates the Metabolism of Polycyclic Aromatic Hydrocarbons in Bacteria. <i>MSystems</i> , 2021, 6, e0063621.	1.7	8
26	Chemical epigenetic manipulation triggers the production of sesquiterpenes from the deep-sea derived <i>Eutypella</i> fungus. <i>Phytochemistry</i> , 2021, 192, 112978.	1.4	10
27	<i>Nitrogeniibacter aestuarii</i> sp. nov., a Novel Nitrogen-Fixing Bacterium Affiliated to the Family Zoogloeaceae and Phylogeny of the Family Zoogloeaceae Revisited. <i>Frontiers in Microbiology</i> , 2021, 12, 755908.	1.5	40
28	<i>Fusibacter ferrireducens</i> sp. nov., an anaerobic, Fe(â..ç)- and sulphur-reducing bacterium isolated from mangrove sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	12
29	<i>Marinobacter mangrovi</i> sp. nov., isolated from mangrove sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	6
30	New Monoterpenoids and Polyketides from the Deep-Sea Sediment-Derived Fungus <i>Aspergillus sydowii</i> MCCC 3A00324. <i>Marine Drugs</i> , 2020, 18, 561.	2.2	10
31	Genomic Characterization and Environmental Distribution of a Thermophilic Anaerobe <i>Dissulfurirhabdus thermomarina</i> SH388T Involved in Disproportionation of Sulfur Compounds in Shallow Sea Hydrothermal Vents. <i>Microorganisms</i> , 2020, 8, 1132.	1.6	12
32	Genome sequencing of deep-sea hydrothermal vent snails reveals adaptations to extreme environments. <i>GigaScience</i> , 2020, 9, .	3.3	5
33	Phenolic bisabolane and cuparene sesquiterpenoids with anti-inflammatory activities from the deep-sea-derived <i>Aspergillus sydowii</i> MCCC 3A00324 fungus. <i>Bioorganic Chemistry</i> , 2020, 105, 104420.	2.0	21
34	Aspeterreureone A, a Cytotoxic Dihydrobenzofuranâ€“Phenyl Acrylate Hybrid from the Deep-Sea-Derived Fungus <i>Aspergillus terreus</i> CC-S06-18. <i>Journal of Natural Products</i> , 2020, 83, 1998-2003.	1.5	26
35	Prokaryotic communities vary with floc size in a biofloc-technology based aquaculture system. <i>Aquaculture</i> , 2020, 529, 735632.	1.7	28
36	Cytotoxic Nitrogenated Azaphilones from the Deep-Sea-Derived Fungus <i>Chaetomium globosum</i> MP4-S01-7. <i>Journal of Natural Products</i> , 2020, 83, 1157-1166.	1.5	39

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37	Metabolic Adaptation to Sulfur of Hyperthermophilic <i>Palaeococcus pacificus</i> DY20341T from Deep-Sea Hydrothermal Sediments. <i>International Journal of Molecular Sciences</i> , 2020, 21, 368.	1.8	8
38	<i>Parahalaea maris</i> sp. nov., isolated from surface seawater and emended description of the genus <i>Parahalaea</i> . <i>Journal of Microbiology</i> , 2020, 58, 92-98.	1.3	13
39	The oxidation of hydrocarbons by diverse heterotrophic and mixotrophic bacteria that inhabit deep-sea hydrothermal ecosystems. <i>ISME Journal</i> , 2020, 14, 1994-2006.	4.4	46
40	âœœ <i>Candidatus</i> <i>Desulfobulbus rimicarenis</i> , an Uncultivated Deltaproteobacterial Epibiont from the Deep-Sea Hydrothermal Vent Shrimp <i>Rimicaris exoculata</i> . <i>Applied and Environmental Microbiology</i> , 2020, 86, .	1.4	24
41	A novel alphaproteobacterium with a small genome identified from the digestive gland of multiple species of abalone. <i>Environmental Microbiology Reports</i> , 2020, 12, 387-395.	1.0	3
42	<i>Sphingorhabdus soli</i> sp. nov., isolated from Arctic soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 1610-1616.	0.8	8
43	<i>Sulfurimonas xiamenensis</i> sp. nov. and <i>Sulfurimonas lithotrophica</i> sp. nov., hydrogen- and sulfur-oxidizing chemolithoautotrophs within the Epsilonproteobacteria isolated from coastal sediments, and an emended description of the genus <i>Sulfurimonas</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 2657-2663.	0.8	23
44	<i>Devosia marina</i> sp. nov., isolated from deep seawater of the South China Sea, and reclassification of <i>Devosia subaequoris</i> as a later heterotypic synonym of <i>Devosia soli</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 3062-3068.	0.8	16
45	<i>Pseudoceanicola aestuarii</i> sp. nov., isolated from the Jiulong River Estuary in PR China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 6220-6225.	0.8	7
46	Identification and Characterization of Nematicidal Volatile Organic Compounds from Deep-Sea <i>Virgibacillus dokdonensis</i> MCCC 1A00493. <i>Molecules</i> , 2020, 25, 744.	1.7	33
47	<i>Marinobacter changyiensis</i> , sp. nov., isolated from offshore sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 3004-3011.	0.8	7
48	<i>Mangrovibacterium lignilyticum</i> sp. nov., a facultatively anaerobic lignin-degrading bacterium isolated from mangrove sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 4502-4507.	0.8	9
49	Aphidicolin Chemistry of the Deep-Sea-Derived Fungus <i>Botryotinia fuckeliana</i> MCCC 3A00494. <i>Journal of Natural Products</i> , 2019, 82, 2307-2331.	1.5	32
50	Graphostromols Aâ€“K, Eleven New Chained Polyketides from the Deepâ€“Seaâ€“Derived <i>Graphostroma</i> sp. <i>Chemistry and Biodiversity</i> , 2019, 16, e1900326.	1.0	2
51	AhIX, an N-acylhomoserine Lactonase with Unique Properties. <i>Marine Drugs</i> , 2019, 17, 387.	2.2	6
52	<i>Nonlabens xiamenensis</i> sp. nov., isolated from coastal seawater. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 1263-1271.	0.7	11
53	Cytotoxic Polyketides Isolated from the Deep-Sea-Derived Fungus <i>Penicillium chrysogenum</i> MCCC 3A00292. <i>Marine Drugs</i> , 2019, 17, 686.	2.2	20
54	Citrinin Monomer and Dimer Derivatives with Antibacterial and Cytotoxic Activities Isolated from the Deep Sea-Derived Fungus <i>Penicillium citrinum</i> NLG-S01-P1. <i>Marine Drugs</i> , 2019, 17, 46.	2.2	36

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55	<i>Sinomicrobium soli</i> sp. nov., isolated from arctic soil. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1070-1074.	0.8	8
56	<i>Altererythrobacter aerophilus</i> sp. nov., isolated from deep-sea water of the north-west Pacific. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1689-1695.	0.8	16
57	<i>Cohaesibacter intestini</i> sp. nov., isolated from the intestine of abalone, <i>Haliotis discus hannai</i> . International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 3202-3206.	0.8	10
58	<i>Pseudidiomarina maritima</i> Wu et al. 2009 is a later heterotypic synonym of <i>Pseudidiomarina tainanensis</i> Jean et al. 2009 and emended description of the species. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 3765-3768.	0.8	5
59	<i>Tsuneonella suprasediminis</i> sp. nov., isolated from the Pacific Ocean. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	16
60	Genetic diversity and population structure of the <i>Bacillus cereus</i> group bacteria from diverse marine environments. Scientific Reports, 2017, 7, 689.	1.6	47
61	Complete genome sequence and whole-genome phylogeny of <i>Kosmotoga pacifica</i> type strain SLHJ1T from an East Pacific hydrothermal sediment. Standards in Genomic Sciences, 2017, 12, 3.	1.5	4
62	A Multilocus Sequence Analysis Scheme for Phylogeny of <i>Thioclava</i> Bacteria and Proposal of Two Novel Species. Frontiers in Microbiology, 2017, 8, 1321.	1.5	24
63	The Pelagic Bacterium <i>Paraphotobacterium marinum</i> Has the Smallest Complete Genome Within the Family Vibrionaceae. Frontiers in Microbiology, 2017, 8, 1994.	1.5	3
64	Sulfur Metabolism of <i>Hydrogenovibrio thermophilus</i> Strain S5 and Its Adaptations to Deep-Sea Hydrothermal Vent Environment. Frontiers in Microbiology, 2017, 8, 2513.	1.5	24
65	<i>Anaeromicrobium sediminis</i> gen. nov., sp. nov., a fermentative bacterium isolated from deep-sea sediment. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 1462-1467.	0.8	15
66	<i>Nioella sediminis</i> sp. nov., isolated from surface sediment and emended description of the genus <i>Nioella</i> . International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 1271-1274.	0.8	17
67	<i>Thioclava nitratireducens</i> sp. nov., isolated from surface seawater. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2109-2113.	0.8	11
68	<i>Shewanella intestini</i> sp. nov., isolated from the intestine of abalone, <i>Haliotis diversicolor</i> . International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 1901-1905.	0.8	6
69	Proposal for transfer of <i>Oceanibulbus indolifex</i> Wagner-Dãbler et al. 2004 to the genus <i>Sulfitobacter</i> as <i>Sulfitobacter indolifex</i> comb. nov.. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2328-2331.	0.8	11
70	<i>Notoacmeibacter marinus</i> gen. nov., sp. nov., isolated from the gut of a limpet and proposal of <i>Notoacmeibacteraceae</i> fam. nov. in the order Rhizobiales of the class Alphaproteobacteria. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2527-2531.	0.8	20
71	<i>Salipiger nanhaiensis</i> Dai et al. 2015 is a later heterotypic synonym of <i>Thiobacimonas profunda</i> Li et al. 2015. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2043-2045.	0.8	4
72	<i>Defluviimonas nitratireducens</i> sp. nov., isolated from surface seawater. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2752-2757.	0.8	13

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73	<i>Croceivirga radialis</i> gen. nov., sp. nov., isolated from a rotten tropical mangrove root. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3733-3738.	0.8	15
74	<i>Agaribacterium haliotis</i> gen. nov., sp. nov., isolated from abalone faeces. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3819-3823.	0.8	11
75	<i>Oceaniglobus indicus</i> gen. nov., sp. nov., a member of the family Rhodobacteraceae isolated from surface seawater. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 4930-4935.	0.8	18
76	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. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3095-3098.	0.8	3
77	<i>Pleomorphobacterium xiamenense</i> Yin et al. 2013 is a later heterotypic synonym of <i>Oceanicella actignis</i> Albuquerque et al. 2012. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3532-3534.	0.8	3
78	<i>Oceanibaculum nanhaiense</i> sp. nov., isolated from surface seawater. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 4842-4845.	0.8	7
79	Changes in the intestinal bacterial community during the growth of white shrimp, <i>Litopenaeus vannamei</i> . Aquaculture Research, 2016, 47, 1737-1746.	0.9	157
80	Cytoglobosins H and I, New Antiproliferative Cytochalasans from Deep-Sea-Derived Fungus <i>Chaetomium globosum</i> . Marine Drugs, 2016, 14, 233.	2.2	29
81	Delta-proteobacterial SAR324 group in hydrothermal plumes on the South Mid-Atlantic Ridge. Scientific Reports, 2016, 6, 22842.	1.6	21
82	Family-wide Structural Characterization and Genomic Comparisons Decode the Diversity-oriented Biosynthesis of Thalassospiramides by Marine Proteobacteria. Journal of Biological Chemistry, 2016, 291, 27228-27238.	1.6	11
83	Comparative genomic and functional analyses: unearthing the diversity and specificity of nematocidal factors in <i>Pseudomonas putida</i> strain 1A00316. Scientific Reports, 2016, 6, 29211.	1.6	15
84	Identification of <i>Bacillus cellulosensis</i> strain NIO-1130T as a member of <i>Bacillus altitudinis</i> and emendation of the latter. Archives of Microbiology, 2016, 198, 835-838.	1.0	6
85	Spiromastilactones: A new class of influenza virus inhibitors from deep-sea fungus. European Journal of Medicinal Chemistry, 2016, 108, 229-244.	2.6	46
86	<i>Paraphotobacterium marinum</i> gen. nov., sp. nov., a member of the family Vibrionaceae, isolated from surface seawater. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 3050-3056.	0.8	23
87	<i>Alcanivorax nanhaiticus</i> sp. nov., isolated from deep sea sediment. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 3651-3655.	0.8	20
88	<i>Pseudotenacibaculum haliotis</i> gen. nov., sp. nov., a new member within the <i>Tenacibaculum-Polaribacter</i> clade of the family Flavobacteriaceae, isolated from the intestine of adult abalones, <i>Haliotis diversicolor</i> and <i>H. discus hannai</i> . International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 3718-3724.	0.8	9
89	<i>Thermodesulfator autotrophicus</i> sp. nov., a thermophilic sulfate-reducing bacterium from the Indian Ocean. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 3978-3982.	0.8	13
90	<i>Wukongibacter baidiensis</i> gen. nov., sp. nov., an anaerobic bacterium isolated from hydrothermal sulfides, and proposal for the reclassification of the closely related <i>Clostridium halophilum</i> and <i>Clostridium caminithermale</i> within <i>Maledivibacter</i> gen. nov. and <i>Paramaledivibacter</i> gen. nov., respectively. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4355-4361.	0.8	37

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91	Genomic insights into the taxonomic status of the <i>Bacillus cereus</i> group. <i>Scientific Reports</i> , 2015, 5, 14082.	1.6	220
92	Complete Genome Sequence of Hyperthermophilic Piezophilic Archaeon <i>Palaeococcus pacificus</i> DY20341 T, Isolated from Deep-Sea Hydrothermal Sediments. <i>Genome Announcements</i> , 2015, 3, .	0.8	5
93	The diversity of PAH-degrading bacteria in a deep-sea water column above the Southwest Indian Ridge. <i>Frontiers in Microbiology</i> , 2015, 6, 853.	1.5	72
94	New Polyphenols from a Deep Sea <i>Spiromastix</i> sp. Fungus, and Their Antibacterial Activities. <i>Marine Drugs</i> , 2015, 13, 2526-2540.	2.2	26
95	The status of the species <i>Hyphomonas rosenbergii</i> Weiner et al. 2000. Request for an Opinion. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 321-321.	0.8	8
96	Phylogenomics characterization of a highly virulent <i>Edwardsiella</i> strain ET080813T encoding two distinct T3SS and three T6SS gene clusters: Propose a novel species as <i>Edwardsiella anguillarum</i> sp. nov.. <i>Systematic and Applied Microbiology</i> , 2015, 38, 36-47.	1.2	126
97	<i>Edwardsiella anguillarum</i> sp. nov., isolated from surface seawater of the Atlantic Ocean and reclassification of <i>Oceanicola batsensis</i> , <i>Oceanicola marinus</i> , <i>Oceanicola nitratireducens</i> , <i>Oceanicola nanhaiensis</i> , <i>Oceanicola antarcticus</i> and <i>Oceanicola flagellatus</i> , as <i>Pseudoceanicola batsensis</i> comb. nov., <i>Pseudoceanicola marinus</i> comb. nov., <i>Pseudoceanicola nitratireducens</i> comb. nov., <i>Pseudoceanicola nanhaiensis</i> comb. nov., <i>Pseudoceanicola antarcticus</i> comb. nov., and <i>Pseudoceanicola flagellatus</i> com. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 1065-1074.	0.7	46
98	<i>Emcibacter nanhaiensis</i> gen. nov. sp. nov., isolated from sediment of the South China Sea. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 893-900.	0.7	18
99	Distribution and diversity of bacterioplankton communities in subtropical seawater around Xiamen Island, China. <i>Microbiological Research</i> , 2015, 175, 16-23.	2.5	20
100	Thermophilic hydrogen-producing bacteria inhabiting deep-sea hydrothermal environments represented by <i>Caloranaerobacter</i> . <i>Research in Microbiology</i> , 2015, 166, 677-687.	1.0	9
101	<i>Mameliella atlantica</i> sp. nov., a marine bacterium of the <i>Roseobacter</i> clade isolated from deep-sea sediment of the South Atlantic Ocean. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 2255-2259.	0.8	13
102	Reclassification of <i>Bacillus invictae</i> as a later heterotypic synonym of <i>Bacillus altitudinis</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 2769-2773.	0.8	10
103	<i>Tamlana nanhaiensis</i> sp. nov., isolated from surface seawater collected from the South China Sea. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 1189-1196.	0.7	13
104	<i>Caloranaerobacter ferrireducens</i> sp. nov., an anaerobic, thermophilic, iron (III)-reducing bacterium isolated from deep-sea hydrothermal sulfide deposits. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1714-1718.	0.8	23
105	<i>Kiloniella litopenaei</i> sp. nov., isolated from the gut microflora of Pacific white shrimp, <i>Litopenaeus vannamei</i> . <i>Antonie Van Leeuwenhoek</i> , 2015, 108, 1293-1299.	0.7	10
106	Effective harvesting of the microalgae <i>Chlorella vulgaris</i> via flocculation—flotation with bioflocculant. <i>Bioresource Technology</i> , 2015, 198, 922-925.	4.8	80
107	Phylogenetic diversity of nitrogen-utilizing genes in hydrothermal chimneys from 3 middle ocean ridges. <i>Extremophiles</i> , 2015, 19, 1173-1182.	0.9	8
108	Draft genome sequence of the denitrifying strain <i>Kiloniella</i> sp. P1-1 isolated from the gut microflora of Pacific white shrimp, <i>Litopenaeus vannamei</i> . <i>Marine Genomics</i> , 2015, 24, 261-263.	0.4	2

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109	<i>Anoxybacter fermentans</i> gen. nov., sp. nov., a piezophilic, thermophilic, anaerobic, fermentative bacterium isolated from a deep-sea hydrothermal vent. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 710-715.	0.8	19
110	<i>Thioclava indica</i> sp. nov., isolated from surface seawater of the Indian Ocean. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 297-304.	0.7	14
111	<i>Halovulum dunhuangense</i> gen. nov., sp. nov., isolated from a saline terrestrial spring. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 2810-2816.	0.8	18
112	<i>Aestuariivita atlantica</i> sp. nov., isolated from deep-sea sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 3281-3285.	0.8	10
113	Identification of strains <i>Bacillus aerophilus</i> MTCC 7304T as <i>Bacillus altitudinis</i> and <i>Bacillus stratosphericus</i> MTCC 7305T as a <i>Proteus</i> sp. and the status of the species <i>Bacillus aerius</i> Shivaji et al. 2006. Request for an Opinion. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 3228-3231.	0.8	26
114	<i>Erythrobacter atlanticus</i> sp. nov., a bacterium from ocean sediment able to degrade polycyclic aromatic hydrocarbons. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 3714-3719.	0.8	45
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117	Oil degradation and biosurfactant production by the deep sea bacterium <i>Dietzia maris</i> As-13-3. <i>Frontiers in Microbiology</i> , 2014, 5, 711.	1.5	81
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