

Min Wu

List of Publications by Year in descending order

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120
papers

2,593
citations

185998

28
h-index

276539

41
g-index

122
all docs

122
docs citations

122
times ranked

2085
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Halomonas saccharevitans</i> sp. nov., <i>Halomonas arcis</i> sp. nov. and <i>Halomonas subterranea</i> sp. nov., halophilic bacteria isolated from hypersaline environments of China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 1619-1624.	0.8	116
2	<i>Parabacteroides chartae</i> sp. nov., an obligately anaerobic species from wastewater of a paper mill. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 2613-2617.	0.8	89
3	Isolation and Complete Genome Sequence of <i>Algibacter alginolytica</i> sp. nov., a Novel Seaweed-Degrading Bacteroidetes Bacterium with Diverse Putative Polysaccharide Utilization Loci. <i>Applied and Environmental Microbiology</i> , 2016, 82, 2975-2987.	1.4	87
4	<i>Pelagibacterium halotolerans</i> gen. nov., sp. nov. and <i>Pelagibacterium luteolum</i> sp. nov., novel members of the family Hyphomicrobiaceae. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 1817-1822.	0.8	85
5	<i>Haloferax larsenii</i> sp. nov., an extremely halophilic archaeon from a solar saltern. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 717-720.	0.8	78
6	<i>Brassicibacter mesophilus</i> gen. nov., sp. nov., a strictly anaerobic bacterium isolated from food industry wastewater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 3018-3023.	0.8	55
7	<i>Natrinema altunense</i> sp. nov., an extremely halophilic archaeon isolated from a salt lake in Altun Mountain in Xinjiang, China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2005, 55, 1311-1314.	0.8	53
8	<i>Sinimariniobacterium flocculans</i> gen. nov., sp. nov., a gammaproteobacterium from offshore surface seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 3541-3546.	0.8	46
9	Cloning, expression and characterization of a halotolerant esterase from a marine bacterium <i>Pelagibacterium halotolerans</i> B2T. <i>Extremophiles</i> , 2012, 16, 427-435.	0.9	45
10	Identification and characterization of novel esterases from a deep-sea sediment metagenome. <i>Archives of Microbiology</i> , 2012, 194, 207-214.	1.0	45
11	<i>Amphibacillus jilinensis</i> sp. nov., a facultatively anaerobic, alkaliphilic bacillus from a soda lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2540-2543.	0.8	44
12	<i>Gracilibacillus ureilyticus</i> sp. nov., a halotolerant bacterium from a saline alkaline soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1383-1386.	0.8	44
13	<i>Pseudoroseovarius zhejiangensis</i> gen. nov., sp. nov., a novel alpha-proteobacterium isolated from the chemical wastewater, and reclassification of <i>Roseovarius crassostreae</i> as <i>Pseudoroseovarius crassostreae</i> comb. nov., <i>Roseovarius sediminilitoris</i> as <i>Pseudoroseovarius sediminilitoris</i> comb. nov. and <i>Roseovarius halocynthiae</i> as <i>Pseudoroseovarius halocynthiae</i> comb. nov.. <i>Antonie Van Leeuwenhoek</i> , 2015, 108, 281-288.	0.7	44
14	<i>Paracoccus sediminis</i> sp. nov., isolated from Pacific Ocean marine sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 2512-2516.	0.8	43
15	<i>Maripseudobacter aurantiacus</i> gen. nov., sp. nov., a novel member of the family Flavobacteriaceae isolated from a sedimentation basin. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 778-783.	0.8	43
16	<i>Thermus arciformis</i> sp. nov., a thermophilic species from a geothermal area. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 834-839.	0.8	42
17	<i>Halolamina salifodinae</i> sp. nov. and <i>Halolamina salina</i> sp. nov., two extremely halophilic archaea isolated from a salt mine. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4380-4385.	0.8	40
18	<i>Haloterrigena saccharevitans</i> sp. nov., an extremely halophilic archaeon from Xin-Jiang, China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2005, 55, 2539-2542.	0.8	37

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19	<i>Kordiimonas lacus</i> sp. nov., isolated from a ballast water tank, and emended description of the genus <i>Kordiimonas</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 422-426.	0.8	36
20	<i>Maribacter thermophilus</i> sp. nov., isolated from an algal bloom in an intertidal zone, and emended description of the genus <i>Maribacter</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 36-41.	0.8	35
21	<i>Muricauda antarctica</i> sp. nov., a marine member of the Flavobacteriaceae isolated from Antarctic seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 3451-3456.	0.8	34
22	<i>Aequorivita viscosa</i> sp. nov., isolated from an intertidal zone, and emended descriptions of <i>Aequorivita antarctica</i> and <i>Aequorivita capsosiphonis</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 3192-3196.	0.8	34
23	<i>Meiothermus cateniformans</i> sp. nov., a slightly thermophilic species from north-eastern China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 840-844.	0.8	33
24	<i>Pseudohongiella nitratireducens</i> sp. nov., isolated from seawater, and emended description of the genus <i>Pseudohongiella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 5155-5160.	0.8	33
25	<i>Pseudoalteromonas lipolytica</i> sp. nov., isolated from the Yangtze River estuary. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2176-2181.	0.8	31
26	<i>Ruegeria marina</i> sp. nov., isolated from Marine Sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 347-350.	0.8	31
27	<i>Gemmobacter megaterium</i> sp. nov., isolated from coastal planktonic seaweeds. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 66-71.	0.8	31
28	<i>Thalassomonas eurytherma</i> sp. nov., a marine proteobacterium. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 2079-2083.	0.8	31
29	<i>Pseudoceanicola lipolyticus</i> sp. nov., a marine alphaproteobacterium, reclassification of <i>Oceanicola flagellatus</i> as <i>Pseudoceanicola flagellatus</i> comb. nov. and emended description of the genus <i>Pseudoceanicola</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 409-415.	0.8	31
30	<i>Halobiforma lacisalsi</i> sp. nov., isolated from a salt lake in China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2005, 55, 1949-1952.	0.8	30
31	<i>Devosia pacifica</i> sp. nov., isolated from deep-sea sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 2637-2641.	0.8	30
32	<i>Halorubrum arcis</i> sp. nov., an extremely halophilic archaeon isolated from a saline lake on the Qinghai-Tibet Plateau, China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 1069-1072.	0.8	29
33	<i>Streptomyces nigra</i> sp. nov. Is a Novel Actinobacterium Isolated From Mangrove Soil and Exerts a Potent Antitumor Activity in Vitro. <i>Frontiers in Microbiology</i> , 2018, 9, 1587.	1.5	29
34	<i>Terasakiella brassicae</i> sp. nov., isolated from the wastewater of a pickle-processing factory, and emended descriptions of <i>Terasakiella pusilla</i> and the genus <i>Terasakiella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 1807-1812.	0.8	29
35	Characterization of alcohol dehydrogenase from the haloalkaliphilic archaeon <i>Natronomonas pharaonis</i> . <i>Extremophiles</i> , 2008, 12, 471-476.	0.9	28
36	<i>Oceanobacillus neutriphilus</i> sp. nov., isolated from activated sludge in a bioreactor. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2409-2414.	0.8	28

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37	<i>Parvularcula flava</i> sp. nov., an alphaproteobacterium isolated from surface seawater of the South China Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 3498-3502.	0.8	28
38	<i>Microbulbifer marinus</i> sp. nov. and <i>Microbulbifer yueqingensis</i> sp. nov., isolated from marine sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 505-510.	0.8	26
39	Image-based promoter prediction: a promoter prediction method based on evolutionarily generated patterns. <i>Scientific Reports</i> , 2018, 8, 17695.	1.6	25
40	Bacterial and archaeal communities in the surface sediment from the northern slope of the South China Sea. <i>Journal of Zhejiang University: Science B</i> , 2009, 10, 890-901.	1.3	24
41	<i>Tepidibacter mesophilus</i> sp. nov., a mesophilic fermentative anaerobe isolated from soil polluted by crude oil, and emended description of the genus <i>Tepidibacter</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 66-70.	0.8	24
42	<i>Anoxybacillus vitaminiphilus</i> sp. nov., a strictly aerobic and moderately thermophilic bacterium isolated from a hot spring. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4064-4071.	0.8	24
43	Activating cryptic biosynthetic gene cluster through a CRISPR-Cas12a-mediated direct cloning approach. <i>Nucleic Acids Research</i> , 2022, 50, 3581-3592.	6.5	23
44	<i>Alkaliphilus halophilus</i> sp. nov., a strictly anaerobic and halophilic bacterium isolated from a saline lake, and emended description of the genus <i>Alkaliphilus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2898-2902.	0.8	22
45	<i>Rosevivax pacificus</i> sp. nov., isolated from deep-sea sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4574-4579.	0.8	22
46	<i>Mesorhizobium oceanicum</i> sp. nov., isolated from deep seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 2739-2745.	0.8	22
47	<i>Extensimonas vulgaris</i> gen. nov., sp. nov., a member of the family Comamonadaceae. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 2062-2068.	0.8	21
48	<i>Gilvimarinus polysaccharolyticus</i> sp. nov., an agar-digesting bacterium isolated from seaweed, and emended description of the genus <i>Gilvimarinus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 562-569.	0.8	21
49	Draft Genome Sequence of Microbacterium profundus Shh49, an Actinobacterium Isolated from Deep-Sea Sediment of a Polymetallic Nodule Environment. <i>Genome Announcements</i> , 2015, 3, .	0.8	21
50	Three New Isoflavonoid Glycosides from the Mangrove-Derived Actinomycete <i>Micromonospora aurantiaca</i> 110B. <i>Marine Drugs</i> , 2019, 17, 294.	2.2	21
51	<i>Muricauda maritima</i> sp. nov., <i>Muricauda aequoris</i> sp. nov. and <i>Muricauda oceanensis</i> sp. nov., three marine bacteria isolated from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 6240-6250.	0.8	21
52	<i>Halomonas endophytica</i> sp. nov., isolated from liquid in the stems of <i>Populus euphratica</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1633-1638.	0.8	20
53	<i>Sphingomonas rubra</i> sp. nov., isolated from bioreactor wastewater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 1028-1032.	0.8	18
54	Draft genome sequence of <i>Microbulbifer elongatus</i> strain HZ11, a brown seaweed-degrading bacterium with potential ability to produce bioethanol from alginate. <i>Marine Genomics</i> , 2014, 18, 83-85.	0.4	18

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55	<i>Marinobacterium zhoushanense</i> sp. nov., isolated from surface seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 3437-3442.	0.8	18
56	<i>Luteolibacter flavescens</i> sp. nov., isolated from deep seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 729-735.	0.8	18
57	<i>Actibacterium pelagium</i> sp. nov., a novel alphaproteobacterium, and emended description of the genus <i>Actibacterium</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 5080-5086.	0.8	18
58	<i>Citricoccus zhacaiensis</i> sp. nov., isolated from a bioreactor for saline wastewater treatment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 495-499.	0.8	17
59	<i>Pectinatus brassicae</i> sp. nov., a Gram-negative, anaerobic bacterium isolated from salty wastewater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 2145-2149.	0.8	17
60	Cloning, expression and characterization of a new enantioselective esterase from a marine bacterium <i>Pelagibacterium halotolerans</i> B2T. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2013, 97, 270-277.	1.8	17
61	<i>Salimesophilobacter vulgaris</i> gen. nov., sp. nov., an anaerobic bacterium isolated from paper-mill wastewater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 1317-1322.	0.8	17
62	Ecological functions of uncultured microorganisms in the cobalt-rich ferromanganese crust of a seamount in the central Pacific are elucidated by fosmid sequencing. <i>Acta Oceanologica Sinica</i> , 2015, 34, 92-113.	0.4	17
63	<i>Halopiger salifodinae</i> sp. nov., an extremely halophilic archaeon isolated from a salt mine. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 3563-3567.	0.8	16
64	<i>Haloimpatiens lingqiaonensis</i> gen. nov., sp. nov., an anaerobic bacterium isolated from paper-mill wastewater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 628-632.	0.8	16
65	<i>Algicoccus marinus</i> gen. nov. sp. nov., a marine bacterium isolated from the surface of brown seaweed <i>Laminaria japonica</i> . <i>Archives of Microbiology</i> , 2019, 201, 943-950.	1.0	15
66	Aldehyde dehydrogenase of the haloalkaliphilic archaeon <i>Natronomonas pharaonis</i> and its function in ethanol metabolism. <i>Extremophiles</i> , 2008, 12, 849-854.	0.9	14
67	<i>Oceanirhabdus sediminicola</i> gen. nov., sp. nov., an anaerobic bacterium isolated from sea sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4277-4283.	0.8	14
68	<i>Rhodohalobacter barkolensis</i> sp. nov., isolated from a saline lake and emended description of the genus <i>Rhodohalobacter</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1949-1954.	0.8	14
69	<i>Pseudomonas mangrovi</i> sp. nov., isolated from mangrove soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 377-383.	0.8	14
70	<i>Aquibacillus salifodinae</i> sp. nov., a novel bacterium isolated from a salt mine in Xinjiang province, China. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 367-374.	0.7	13
71	Polyphasic taxonomic characterisation of a novel strain as <i>Pararhizobium haloflavum</i> sp. nov., isolated from soil samples near a sewage treatment tank. <i>Antonie Van Leeuwenhoek</i> , 2018, 111, 485-491.	0.7	13
72	<i>Pseudomonas pharmafabricae</i> sp. nov., Isolated From Pharmaceutical Wastewater. <i>Current Microbiology</i> , 2018, 75, 1119-1125.	1.0	13

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73	<i>Roseivirga marina</i> sp. nov., isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 4239-4243.	0.8	13
74	<i>Muriicola marianensis</i> sp. nov., isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 407-411.	0.8	12
75	<i>Kordiimonas pumila</i> sp. nov., isolated from coastal sediment. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 1743-1748.	0.8	12
76	<i>Saccharospirillum mangrovi</i> sp. nov., a bacterium isolated from mangrove sediment. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2813-2818.	0.8	12
77	<i>Meridianimarinicoccus roseus</i> gen. nov., sp. nov., a novel genus of the family Rhodobacteraceae isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 504-510.	0.8	12
78	<i>Halomonas populi</i> sp. nov. isolated from <i>Populus euphratica</i> . Archives of Microbiology, 2022, 204, 86.	1.0	12
79	<i>Halopelagius longus</i> sp. nov., a member of the family Halobacteriaceae isolated from a salt mine, and emended description of the genus <i>Halopelagius</i> . International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3585-3590.	0.8	11
80	<i>Thalassobaculum fulvum</i> sp. nov., isolated from deep seawater. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 2186-2191.	0.8	11
81	<i>Marinibaculum pumilum</i> gen. nov., sp. nov., isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4844-4849.	0.8	11
82	<i>Marinicaulis flavus</i> gen. nov., sp. nov., a novel stalked bacterium of the family Parvularculaceae. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2061-2067.	0.8	11
83	Culturable halophilic archaeal diversity of Ayakekumu salt lake located in Xinjiang, China. Acta Ecologica Sinica, 2007, 27, 3119-3123.	0.9	10
84	<i>Aestuarium zhoushanense</i> gen. nov., sp. nov., Isolated from the Tidal Flat. Current Microbiology, 2017, 74, 1469-1476.	1.0	10
85	Description of <i>Erythrobacter mangrovi</i> sp. nov., an aerobic bacterium from rhizosphere soil of mangrove plant (<i>Kandelia candel</i>). Antonie Van Leeuwenhoek, 2020, 113, 1425-1435.	0.7	10
86	<i>Idiomarina mangrovi</i> sp. nov., isolated from rhizosphere soil of a mangrove <i>Avicennia marina</i> forest. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1662-1668.	0.8	10
87	Microbial community structure and nitrogenase gene diversity of sediment from a deep-sea hydrothermal vent field on the Southwest Indian Ridge. Acta Oceanologica Sinica, 2014, 33, 94-104.	0.4	9
88	<i>Bacillus salitolerans</i> sp. nov., a novel bacterium isolated from a salt mine in Xinjiang province, China. Antonie Van Leeuwenhoek, 2015, 108, 443-451.	0.7	9
89	<i>Gallaecimonas mangrovi</i> sp. nov., a novel bacterium isolated from mangrove sediment. Antonie Van Leeuwenhoek, 2018, 111, 1855-1862.	0.7	9
90	<i>Chryseobacterium lineare</i> sp. nov., isolated from a limpid stream. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 800-805.	0.8	9

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91	<i>Roseovarius nitratireducens</i> sp. nov., a halotolerant bacterium isolated from a saline lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1204-1209.	0.8	9
92	Description of two novel anaerobic members in the family Clostridiaceae, <i>Anaeromonas gelatinilytica</i> gen.nov., sp. nov., and <i>Anaeromonas frigorigresistens</i> sp. nov., isolated from saline lake sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	9
93	<i>Micromonospora zhangzhouensis</i> sp. nov., a Novel Actinobacterium Isolated from Mangrove Soil, Exerts a Cytotoxic Activity in vitro. <i>Scientific Reports</i> , 2020, 10, 3889.	1.6	8
94	<i>Alkalilacustris brevis</i> gen. nov., sp. nov., isolated from a soda lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 1669-1675.	0.8	8
95	<i>Marortus luteolus</i> gen. nov., sp. nov., isolated from surface seawater of the East Sea in China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 1490-1495.	0.8	8
96	<i>Emcibacter congregatus</i> sp. nov., isolated from sediment cultured in situ. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2846-2850.	0.8	7
97	Discovery of Three 22-Membered Macrolides by Deciphering the Streamlined Genome of Mangrove-Derived <i>Streptomyces</i> sp. HM190. <i>Frontiers in Microbiology</i> , 2020, 11, 1464.	1.5	6
98	Description of <i>Wenzhouxiangella salilacus</i> sp. nov., a moderate halophilic bacterium isolated from a salt lake in Xinjiang Province, China. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 847-855.	0.7	5
99	Complete genome sequence of <i>Salinigranum rubrum</i> CX10T, an extremely halophilic archaeon isolated from a marine solar saltern. <i>Marine Genomics</i> , 2019, 44, 57-60.	0.4	5
100	<i>Nocardioides malaquae</i> sp. nov., a novel actinobacterium isolated from sewage sludge of a fisheries processing factory. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	5
101	<i>Seohaecicola zhoushanensis</i> sp. nov., isolated from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 3029-3033.	0.8	5
102	<i>Nesterenkonia muleiensis</i> sp. nov., a novel actinobacterium isolated from sap of <i>Populus euphratica</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 1888-1894.	0.8	5
103	<i>Agromyces kandeliae</i> sp. nov., isolated from rhizosphere soil of <i>Kandelia candel</i> in a mangrove. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 5861-5867.	0.8	5
104	<i>Nitratireductor mangrovi</i> sp. nov., a Nitrate-Reducing Bacterium Isolated from Mangrove Soil. <i>Current Microbiology</i> , 2020, 77, 1334-1340.	1.0	4
105	<i>Marinobacter caseinilyticus</i> sp. nov., Isolated from Saline Soil. <i>Current Microbiology</i> , 2021, 78, 1045-1052.	1.0	4
106	<i>Agromyces mangrovi</i> sp. nov., a Novel Actinobacterium Isolated from Mangrove Soil. <i>Current Microbiology</i> , 2018, 75, 1055-1061.	1.0	3
107	<i>Nesterenkonia ebinurensis</i> sp. nov., a Novel Actinobacterium Isolated From <i>Populus euphratica</i> . <i>Current Microbiology</i> , 2021, 78, 3804-3810.	1.0	3
108	<i>Gramella crocea</i> sp. nov., isolated from activated sludge of a seafood processing plant. <i>Antonie Van Leeuwenhoek</i> , 2022, 115, 969-978.	0.7	3

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109	Isolation and Phylogenetic Analysis of Halophilic Archaeon AJ6. <i>Frontiers of Biology in China: Selected Publications From Chinese Universities</i> , 2006, 1, 29-34.	0.2	2
110	Draft genome sequence of Halopiger salifodinae KCY07-B2T, an extremely halophilic archaeon isolated from a salt mine. <i>Standards in Genomic Sciences</i> , 2015, 10, 124.	1.5	2
111	Chemical Oxygen Demand Removal in Na ₂ SO ₄ Saturated Wastewater by Cultivable Halotolerant Bacterial Community. <i>Clean - Soil, Air, Water</i> , 2016, 44, 593-598.	0.7	2
112	The Complete Genome of <i>Emcibacter congregatus</i> ZYLT, a Marine Bacterium Encoding a CRISPR-Cas 9 Immune System. <i>Current Microbiology</i> , 2020, 77, 762-768.	1.0	2
113	<i>Pseudomonas lopnurensis</i> sp. nov., an endophytic bacterium isolated from <i>Populus euphratica</i> at the ancient Ugan river. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 399-410.	0.7	2
114	Complete genome sequence of marine Roseobacter lineage member <i>Monaibacterium</i> sp. ALG8 with six plasmids isolated from seawater around brown algae. <i>Marine Genomics</i> , 2021, 60, 100878.	0.4	2
115	Draft Genome Sequence of <i>Amphibacillus jilinensis</i> Y1T, a Facultatively Anaerobic, Alkaliphilic and Halotolerant Bacterium. <i>Standards in Genomic Sciences</i> , 2013, 8, 491-499.	1.5	1
116	Draft genomic DNA sequence of strain <i>Halomonas</i> sp. FS-N4 exhibiting high catalase activity. <i>Marine Genomics</i> , 2014, 18, 119-121.	0.4	1
117	Genome sequence of facultatively anaerobic marine bacterium <i>Maribacter thermophilus</i> strain HT7-2T. <i>Marine Genomics</i> , 2015, 24, 265-268.	0.4	1
118	High quality draft genome sequence of an extremely halophilic archaeon <i>Natrinema altunense</i> strain AJ2T. <i>Standards in Genomic Sciences</i> , 2017, 12, 25.	1.5	1
119	<i>Anaerophilus nitritogenes</i> gen. nov., sp. nov., isolated from salt lake sediment in Xinjiang Province, China. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 417-425.	0.7	1
120	<i>Confluentibacter flavum</i> sp. nov., Isolated from the Saline Lake. <i>Current Microbiology</i> , 2018, 75, 1447-1452.	1.0	0