

Jin-Woo Bae

List of Publications by Year in descending order

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230
papers

14,014
citations

57631

44
h-index

26548

107
g-index

239
all docs

239
docs citations

239
times ranked

17044
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteobacteria: microbial signature of dysbiosis in gut microbiota. Trends in Biotechnology, 2015, 33, 496-503.	4.9	2,453
2	An increase in the <i>Akkermansia</i> spp. population induced by metformin treatment improves glucose homeostasis in diet-induced obese mice. Gut, 2014, 63, 727-735.	6.1	1,288
3	Innate Immune Homeostasis by the Homeobox Gene <i>Caudal</i> and Commensal-Gut Mutualism in <i>Drosophila</i> . Science, 2008, 319, 777-782.	6.0	766
4	Insect Gut Bacterial Diversity Determined by Environmental Habitat, Diet, Developmental Stage, and Phylogeny of Host. Applied and Environmental Microbiology, 2014, 80, 5254-5264.	1.4	591
5	Metagenomic Analysis of Kimchi, a Traditional Korean Fermented Food. Applied and Environmental Microbiology, 2011, 77, 2264-2274.	1.4	416
6	Transfer of a healthy microbiota reduces amyloid and tau pathology in an Alzheimer's disease animal model. Gut, 2020, 69, 283-294.	6.1	336
7	Diversity of ammonium-oxidizing bacteria in a granular sludge anaerobic ammonium-oxidizing (anammox) reactor. Environmental Microbiology, 2008, 10, 3130-3139.	1.8	283
8	Amplification Methods Bias Metagenomic Libraries of Uncultured Single-Stranded and Double-Stranded DNA Viruses. Applied and Environmental Microbiology, 2011, 77, 7663-7668.	1.4	259
9	Investigation of archaeal and bacterial diversity in fermented seafood using barcoded pyrosequencing. ISME Journal, 2010, 4, 1-16.	4.4	256
10	Diversity and Abundance of Single-Stranded DNA Viruses in Human Feces. Applied and Environmental Microbiology, 2011, 77, 8062-8070.	1.4	207
11	Comparative Analysis of Korean Human Gut Microbiota by Barcoded Pyrosequencing. PLoS ONE, 2011, 6, e22109.	1.1	199
12	Bacterial community analysis during fermentation of ten representative kinds of kimchi with barcoded pyrosequencing. Food Microbiology, 2012, 30, 197-204.	2.1	198
13	Strict vegetarian diet improves the risk factors associated with metabolic diseases by modulating gut microbiota and reducing intestinal inflammation. Environmental Microbiology Reports, 2013, 5, 765-775.	1.0	171
14	Impact of Pelvic Radiotherapy on Gut Microbiota of Gynecological Cancer Patients Revealed by Massive Pyrosequencing. PLoS ONE, 2013, 8, e82659.	1.1	171
15	Enteric Viruses Ameliorate Gut Inflammation via Toll-like Receptor 3 and Toll-like Receptor 7-Mediated Interferon- β Production. Immunity, 2016, 44, 889-900.	6.6	170
16	Influence of Soil Components on the Biodegradation of Benzene, Toluene, Ethylbenzene, and <i>o</i> -, <i>m</i> -, and <i>p</i> -Xylenes by the Newly Isolated Bacterium <i>Pseudoxanthomonas spadix</i> BD-a59. Applied and Environmental Microbiology, 2008, 74, 7313-7320.	1.4	149
17	Amplification of Uncultured Single-Stranded DNA Viruses from Rice Paddy Soil. Applied and Environmental Microbiology, 2008, 74, 5975-5985.	1.4	148
18	Comparing microarrays and next-generation sequencing technologies for microbial ecology research. Trends in Biotechnology, 2010, 28, 291-299.	4.9	142

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19	Chronic Repression of mTOR Complex 2 Induces Changes in the Gut Microbiota of Diet-induced Obese Mice. <i>Scientific Reports</i> , 2016, 6, 30887.	1.6	142
20	Lysogeny is prevalent and widely distributed in the murine gut microbiota. <i>ISME Journal</i> , 2018, 12, 1127-1141.	4.4	140
21	Development and Evaluation of Genome-Probing Microarrays for Monitoring Lactic Acid Bacteria. <i>Applied and Environmental Microbiology</i> , 2005, 71, 8825-8835.	1.4	114
22	Analysis of yeast and archaeal population dynamics in kimchi using denaturing gradient gel electrophoresis. <i>International Journal of Food Microbiology</i> , 2008, 126, 159-166.	2.1	113
23	Metagenomic Analysis of the Viral Communities in Fermented Foods. <i>Applied and Environmental Microbiology</i> , 2011, 77, 1284-1291.	1.4	108
24	<i>Blautia faecis</i> sp. nov., isolated from human faeces. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 599-603.	0.8	107
25	Metagenomic Characterization of Airborne Viral DNA Diversity in the Near-Surface Atmosphere. <i>Journal of Virology</i> , 2012, 86, 8221-8231.	1.5	103
26	Host habitat is the major determinant of the gut microbiome of fish. <i>Microbiome</i> , 2021, 9, 166.	4.9	100
27	Unexpected convergence of fungal and bacterial communities during fermentation of traditional Korean alcoholic beverages inoculated with various natural starters. <i>Food Microbiology</i> , 2012, 30, 112-123.	2.1	96
28	Bacterial, archaeal, and eukaryal diversity in the intestines of Korean people. <i>Journal of Microbiology</i> , 2008, 46, 491-501.	1.3	85
29	Phylogenetic Characterization of Two Novel Commensal Bacteria Involved with Innate Immune Homeostasis in <i>Drosophila melanogaster</i> . <i>Applied and Environmental Microbiology</i> , 2008, 74, 6171-6177.	1.4	85
30	<i>Geobacillus toebii</i> sp. nov., a novel thermophilic bacterium isolated from hay compost. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2002, 52, 2251-2255.	0.8	80
31	<i>Arthrobacter soli</i> sp. nov., a novel bacterium isolated from wastewater reservoir sediment. <i>Journal of Microbiology</i> , 2008, 46, 40-44.	1.3	77
32	Longitudinal evaluation of fecal microbiota transplantation for ameliorating calf diarrhea and improving growth performance. <i>Nature Communications</i> , 2021, 12, 161.	5.8	76
33	Spatial disturbances in altered mucosal and luminal gut viromes of diet-induced obese mice. <i>Environmental Microbiology</i> , 2016, 18, 1498-1510.	1.8	73
34	Metatranscriptome analysis of lactic acid bacteria during kimchi fermentation with genome-probing microarrays. <i>International Journal of Food Microbiology</i> , 2009, 130, 140-146.	2.1	71
35	Obesogenic diet-induced gut barrier dysfunction and pathobiont expansion aggravate experimental colitis. <i>PLoS ONE</i> , 2017, 12, e0187515.	1.1	71
36	<i>Halalkalicoccus jeotgali</i> sp. nov., a halophilic archaeon from shrimp jeotgal, a traditional Korean fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 2296-2298.	0.8	69

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37	Statistical superiority of genome-probing microarrays as genomic DNA-DNA hybridization in revealing the bacterial phylogenetic relationship compared to conventional methods. <i>Journal of Microbiological Methods</i> , 2008, 75, 523-530.	0.7	67
38	Autophagy deficiency in myeloid cells increases susceptibility to obesity-induced diabetes and experimental colitis. <i>Autophagy</i> , 2016, 12, 1390-1403.	4.3	65
39	The human gut archaeome: identification of diverse haloarchaea in Korean subjects. <i>Microbiome</i> , 2020, 8, 114.	4.9	65
40	Social status shapes the bacterial and fungal gut communities of the honey bee. <i>Scientific Reports</i> , 2018, 8, 2019.	1.6	64
41	<i>Blautia stercoris</i> sp. nov., isolated from human faeces. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 776-779.	0.8	57
42	<i>Endozoicomonas atrinae</i> sp. nov., isolated from the intestine of a comb pen shell <i>Atrina pectinata</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 2312-2318.	0.8	52
43	<i>Brachybacterium squillarum</i> sp. nov., isolated from salt-fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 1118-1122.	0.8	49
44	<i>Pedobacter agri</i> sp. nov., from soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 1640-1643.	0.8	47
45	<i>Methylobacterium platani</i> sp. nov., isolated from a leaf of the tree <i>Platanus orientalis</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 2849-2853.	0.8	44
46	<i>Pseudoalteromonas marina</i> sp. nov., a marine bacterium isolated from tidal flats of the Yellow Sea, and reclassification of <i>Pseudoalteromonas sagamiensis</i> as <i>Algicola sagamiensis</i> comb. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 12-18.	0.8	44
47	<i>Acinetobacter apis</i> sp. nov., isolated from the intestinal tract of a honey bee, <i>Apis mellifera</i> . <i>Journal of Microbiology</i> , 2014, 52, 639-645.	1.3	44
48	<i>Bombella apis</i> sp. nov., an acetic acid bacterium isolated from the midgut of a honey bee. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 2184-2188.	0.8	44
49	<i>Marinobacterium halophilum</i> sp. nov., a marine bacterium isolated from the Yellow Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 77-80.	0.8	43
50	<i>Natronococcus jeotgali</i> sp. nov., a halophilic archaeon isolated from shrimp jeotgal, a traditional fermented seafood from Korea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 2129-2131.	0.8	42
51	<i>Sulfitobacter litoralis</i> sp. nov., a marine bacterium isolated from the East Sea, Korea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 692-695.	0.8	42
52	Determination of Cyanobacterial Diversity during Algal Blooms in Daechung Reservoir, Korea, on the Basis of <i>cpcBA</i> Intergenic Spacer Region Analysis. <i>Applied and Environmental Microbiology</i> , 2006, 72, 3252-3258.	1.4	41
53	<i>Mucilaginibacter oryzae</i> sp. nov., isolated from soil of a rice paddy. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 1451-1454.	0.8	41
54	<i>Haloterrigena jeotgali</i> sp. nov., an extremely halophilic archaeon from salt-fermented food. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 2359-2363.	0.8	41

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55	<i>Oceanobacillus kimchii</i> sp. nov. Isolated from a traditional Korean fermented food. <i>Journal of Microbiology</i> , 2010, 48, 862-866.	1.3	41
56	<i>Paracoccus aestuarii</i> sp. nov., isolated from tidal flat sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 790-794.	0.8	41
57	<i>Luteimonas aestuarii</i> sp. nov., isolated from tidal flat sediment. <i>Journal of Microbiology</i> , 2008, 46, 525-529.	1.3	40
58	<i>Joostella marina</i> gen. nov., sp. nov., a novel member of the family Flavobacteriaceae isolated from the East Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 1388-1392.	0.8	40
59	Application of quantitative real-time PCR for enumeration of total bacterial, archaeal, and yeast populations in kimchi. <i>Journal of Microbiology</i> , 2009, 47, 682-685.	1.3	38
60	<i>Kocuria atrinae</i> sp. nov., isolated from traditional Korean fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 914-918.	0.8	37
61	<i>Kocuria koreensis</i> sp. nov., isolated from fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 140-143.	0.8	37
62	ESRRA (estrogen related receptor alpha) is a critical regulator of intestinal homeostasis through activation of autophagic flux via gut microbiota. <i>Autophagy</i> , 2021, 17, 2856-2875.	4.3	37
63	<i>Leucobacter celer</i> sp. nov., isolated from Korean fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2353-2357.	0.8	36
64	<i>Stappia marina</i> sp. nov., a marine bacterium isolated from the Yellow Sea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2006, 56, 75-79.	0.8	35
65	<i>Halapricum salinum</i> gen. nov., sp. nov., an extremely halophilic archaeon isolated from non-purified solar salt. <i>Antonie Van Leeuwenhoek</i> , 2014, 105, 979-986.	0.7	35
66	<i>Alishewanella jeotgali</i> sp. nov., isolated from traditional fermented food, and emended description of the genus <i>Alishewanella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 2313-2316.	0.8	34
67	<i>Halorubrum cibi</i> sp. nov., an extremely halophilic archaeon from salt-fermented seafood. <i>Journal of Microbiology</i> , 2009, 47, 162-166.	1.3	33
68	<i>Cobetia crustatorum</i> sp. nov., a novel slightly halophilic bacterium isolated from traditional fermented seafood in Korea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 620-626.	0.8	33
69	<i>Shimia haliotis</i> sp. nov., a bacterium isolated from the gut of an abalone, <i>Haliotis discus hannai</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4248-4253.	0.8	33
70	<i>Acidovorax soli</i> sp. nov., isolated from landfill soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2715-2718.	0.8	32
71	<i>Paenibacillus oceanisediminis</i> sp. nov. isolated from marine sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 428-434.	0.8	31
72	<i>Halomonas jeotgali</i> sp. nov., a new moderate halophilic bacterium isolated from a traditional fermented seafood. <i>Journal of Microbiology</i> , 2010, 48, 404-410.	1.3	30

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73	<i>Leucobacter salsicius</i> sp. nov., from a salt-fermented food. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 502-506.	0.8	30
74	<i>Orbus sasakiae</i> sp. nov., a bacterium isolated from the gut of the butterfly <i>Sasakia charonda</i> , and emended description of the genus <i>Orbus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 1766-1770.	0.8	30
75	<i>Vagococcus martis</i> sp. nov., isolated from the small intestine of a marten, <i>Martes flavigula</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 3398-3402.	0.8	30
76	Homogeneous versus heterogeneous probes for microbial ecological microarrays. <i>Trends in Biotechnology</i> , 2006, 24, 318-323.	4.9	29
77	<i>Alishewanella aestuarii</i> sp. nov., isolated from tidal flat sediment, and emended description of the genus <i>Alishewanella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 421-424.	0.8	29
78	<i>Pseudomonas sabulinigri</i> sp. nov., isolated from black beach sand. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 38-41.	0.8	29
79	<i>Sphingopyxis soli</i> sp. nov., isolated from landfill soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1682-1686.	0.8	29
80	<i>Weissella diestrammenae</i> sp. nov., isolated from the gut of a camel cricket (<i>Diestrammena coreana</i>). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 2951-2956.	0.8	29
81	<i>Blastopirellula cremea</i> sp. nov., isolated from a dead ark clam. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 2314-2319.	0.8	29
82	Phenotypic and Genomic Properties of <i>Brachybacterium vulturis</i> sp. nov. and <i>Brachybacterium avium</i> sp. nov.. <i>Frontiers in Microbiology</i> , 2018, 9, 1809.	1.5	29
83	<i>Clostridium hastiforme</i> is a later synonym of <i>Tissierella praeacuta</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2004, 54, 947-949.	0.8	29
84	A GLP α 1/GLP α 2 receptor dual agonist to treat NASH: Targeting the gut-liver axis and microbiome. <i>Hepatology</i> , 2022, 75, 1523-1538.	3.6	29
85	Characterization of the depth-related changes in the microbial communities in Lake Hovsgol sediment by 16S rRNA gene-based approaches. <i>Journal of Microbiology</i> , 2008, 46, 125-136.	1.3	28
86	<i>Nitratireductor basaltis</i> sp. nov., isolated from black beach sand. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 135-138.	0.8	28
87	<i>Henriciella marina</i> gen. nov., sp. nov., a novel member of the family Hyphomonadaceae isolated from the East Sea. <i>Journal of Microbiology</i> , 2009, 47, 156-161.	1.3	28
88	<i>Alishewanella agri</i> sp. nov., isolated from landfill soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2199-2203.	0.8	28
89	<i>Kocuria salsicia</i> sp. nov., isolated from salt-fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 286-289.	0.8	28
90	<i>Ruegeria conchae</i> sp. nov., isolated from the ark clam <i>Scapharca broughtonii</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 2851-2857.	0.8	28

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91	Viral community predicts the geographical origin of fermented vegetable foods more precisely than bacterial community. <i>Food Microbiology</i> , 2018, 76, 319-327.	2.1	28
92	Design of long oligonucleotide probes for functional gene detection in a microbial community. <i>Bioinformatics</i> , 2005, 21, 4092-4100.	1.8	27
93	<i>Nocardioides basaltis</i> sp. nov., isolated from black beach sand. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 42-47.	0.8	27
94	<i>Halorubrum halophilum</i> sp. nov., an extremely halophilic archaeon isolated from a salt-fermented seafood. <i>Antonie Van Leeuwenhoek</i> , 2014, 105, 603-612.	0.7	27
95	<i>Proteus cibarius</i> sp. nov., a swarming bacterium from Jeotgal, a traditional Korean fermented seafood, and emended description of the genus <i>Proteus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 2158-2164.	0.8	27
96	<i>Arthrobacter subterraneus</i> sp. nov., isolated from deep subsurface water of the South Coast of Korea. <i>Journal of Microbiology and Biotechnology</i> , 2007, 17, 1875-9.	0.9	27
97	Characterization of <i>Symbiobacterium toebii</i> , an obligate commensal thermophile isolated from compost. <i>Extremophiles</i> , 2002, 6, 57-64.	0.9	26
98	Generation of subspecies level-specific microbial diagnostic microarrays using genes amplified from subtractive suppression hybridization as microarray probes. <i>Nucleic Acids Research</i> , 2005, 33, e113-e113.	6.5	26
99	<i>Polaribacter atrinae</i> sp. nov., isolated from the intestine of a comb pen shell, <i>Atrina pectinata</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 1654-1661.	0.8	26
100	Oral microbiome associated with lymph node metastasis in oral squamous cell carcinoma. <i>Scientific Reports</i> , 2021, 11, 23176.	1.6	26
101	<i>Haladaptatus cibarius</i> sp. nov., an extremely halophilic archaeon from seafood, and emended description of the genus <i>Haladaptatus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1187-1190.	0.8	25
102	<i>Actibacter haliotis</i> sp. nov., isolated from the gut of an abalone, <i>Haliotis discus hannai</i> , and emended description of the genus <i>Actibacter</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 49-55.	0.8	25
103	Development of Microbial Genome-Probing Microarrays Using Digital Multiple Displacement Amplification of Uncultivated Microbial Single Cells. <i>Environmental Science & Technology</i> , 2008, 42, 6058-6064.	4.6	24
104	Phenotypic characterization and genomic analysis of the <i>Shigella sonnei</i> bacteriophage SP18. <i>Journal of Microbiology</i> , 2010, 48, 213-222.	1.3	24
105	<i>Dietzia alimentaria</i> sp. nov., isolated from a traditional Korean food. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2254-2258.	0.8	24
106	<i>Bacteroides faecis</i> sp. nov., isolated from human faeces. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 2572-2576.	0.8	23
107	<i>Virgibacillus alimentarius</i> sp. nov., isolated from a traditional Korean food. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2851-2855.	0.8	23
108	<i>Kistimonas scapharcae</i> sp. nov., isolated from a dead ark clam (<i>Scapharca broughtonii</i>), and emended description of the genus <i>Kistimonas</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 2865-2869.	0.8	23

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109	<i>Dyella jejuensis</i> sp. nov., isolated from soil of Hallasan Mountain in Jeju Island. <i>Journal of Microbiology</i> , 2014, 52, 373-377.	1.3	23
110	Description of <i>Nocardioides piscis</i> sp. nov., <i>Sphingomonas piscis</i> sp. nov. and <i>Sphingomonas sinipercae</i> sp. nov., isolated from the intestine of fish species <i>Odontobutis interrupta</i> (Korean spotted sleeper) and <i>Siniperca scherzeri</i> (leopard mandarin fish). <i>Journal of Microbiology</i> , 2021, 59, 552-562.	1.3	23
111	<i>Vibrio litoralis</i> sp. nov., isolated from a Yellow Sea tidal flat in Korea. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 562-565.	0.8	23
112	Quantitative real time PCR assays for the enumeration of <i>Saccharomyces cerevisiae</i> and the <i>Saccharomyces sensu stricto</i> complex in human feces. <i>Journal of Microbiological Methods</i> , 2007, 71, 191-201.	0.7	22
113	<i>Aliihoeflea aestuarii</i> gen. nov., sp. nov., a novel bacterium isolated from tidal flat sediment. <i>Journal of Microbiology</i> , 2008, 46, 594-598.	1.3	22
114	<i>Marinobacter goseongensis</i> sp. nov., from seawater. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 2866-2870.	0.8	22
115	<i>Agromyces atrinae</i> sp. nov., isolated from fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1056-1059.	0.8	22
116	<i>Lentibacillus jeotgali</i> sp. nov., a halophilic bacterium isolated from traditional Korean fermented seafood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1017-1022.	0.8	22
117	The effects of sequencing platforms on phylogenetic resolution in 16S rRNA gene profiling of human feces. <i>Scientific Data</i> , 2018, 5, 180068.	2.4	22
118	<i>Paenibacillus apis</i> sp. nov. and <i>Paenibacillus intestini</i> sp. nov., isolated from the intestine of the honey bee <i>Apis mellifera</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1918-1924.	0.8	22
119	<i>Blautia hominis</i> sp. nov., isolated from human faeces. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1059-1064.	0.8	22
120	<i>Sphingomonas aestuarii</i> sp. nov., isolated from tidal flat sediment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 1359-1363.	0.8	21
121	<i>Brevundimonas basaltis</i> sp. nov., isolated from black sand. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1488-1492.	0.8	21
122	Complete Genome Sequence of <i>Halalkalicoccus jeotgali</i> B3 ^T , an Extremely Halophilic Archaeon. <i>Journal of Bacteriology</i> , 2010, 192, 4528-4529.	1.0	21
123	<i>Neptunomonas concharum</i> sp. nov., isolated from a dead ark clam, and emended description of the genus <i>Neptunomonas</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 2657-2661.	0.8	21
124	<i>Rhodopirellula rosea</i> sp. nov., a novel bacterium isolated from an ark clam <i>Scapharca broughtonii</i> . <i>Journal of Microbiology</i> , 2013, 51, 301-304.	1.3	21
125	<i>Cloacibacterium haliotis</i> sp. nov., isolated from the gut of an abalone, <i>Haliotis discus hannai</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 72-77.	0.8	21
126	<i>Leucobacter coleopterorum</i> sp. nov., <i>Leucobacter insecticola</i> sp. nov., and <i>Leucobacter viscericola</i> sp. nov., isolated from the intestine of the diving beetles, <i>Cybister brevis</i> and <i>Cybister lewisianus</i> , and emended description of the genus <i>Leucobacter</i> . <i>Journal of Microbiology</i> , 2021, 59, 360-368.	1.3	21

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127	The green tea modulates large intestinal microbiome and exo/endogenous metabolome altered through chronic UVB-exposure. <i>PLoS ONE</i> , 2017, 12, e0187154.	1.1	21
128	Arginine-mediated gut microbiome remodeling promotes host pulmonary immune defense against nontuberculous mycobacterial infection. <i>Gut Microbes</i> , 2022, 14, 2073132.	4.3	21
129	Isolation of uncultivated anaerobic thermophiles from compost by supplementing cell extract of <i>Geobacillus toebii</i> in enrichment culture medium. <i>Extremophiles</i> , 2005, 9, 477-485.	0.9	20
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