## Ok-Sun Kim

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Introducing EzTaxon-e: a prokaryotic 16S rRNA gene sequence database with phylotypes that represent uncultured species. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 716-721.	0.8	4,898
2	Phylogenetic and functional marker genes to study ammonia-oxidizing microorganisms (AOM) in the environment. Applied Microbiology and Biotechnology, 2010, 85, 425-440.	1.7	144
3	Bacterial community structure and soil properties of a subarctic tundra soil in Council, Alaska. FEMS Microbiology Ecology, 2014, 89, 465-475.	1.3	121
4	Use of Barcoded Pyrosequencing and Shared OTUs To Determine Sources of Fecal Bacteria in Watersheds. Environmental Science & Technology, 2010, 44, 7777-7782.	4.6	108
5	Influence of Soil Characteristics and Proximity to Antarctic Research Stations on Abundance of Antibiotic Resistance Genes in Soils. Environmental Science & Technology, 2016, 50, 12621-12629.	4.6	107
6	Algal and Fungal Diversity in Antarctic Lichens. Journal of Eukaryotic Microbiology, 2015, 62, 196-205.	0.8	53
7	Highly Heterogeneous Soil Bacterial Communities around Terra Nova Bay of Northern Victoria Land, Antarctica. PLoS ONE, 2015, 10, e0119966.	1.1	51
8	Comparative in silico analysis of PCR primers suited for diagnostics and cloning of ammonia monooxygenase genes from ammonia-oxidizing bacteria. FEMS Microbiology Ecology, 2008, 64, 141-152.	1.3	50
9	Niche specialization of bacteria in permanently ice overed lakes of the <scp>M</scp> cMurdo Dry Valleys, <scp>A</scp> ntarctica. Environmental Microbiology, 2017, 19, 2258-2271.	1.8	49
10	Bacterial Communities of Surface Mixed Layer in the Pacific Sector of the Western Arctic Ocean during Sea-Ice Melting. PLoS ONE, 2014, 9, e86887.	1.1	40
11	Evaluation of PCR Primer Selectivity and Phylogenetic Specificity by Using Amplification of 16S rRNA Genes from Betaproteobacterial Ammonia-Oxidizing Bacteria in Environmental Samples. Applied and Environmental Microbiology, 2008, 74, 5231-5236.	1.4	32
12	Distribution of denitrifying bacterial communities in the stratified water column and sediment–water interface in two freshwater lakes and the Baltic Sea. Aquatic Ecology, 2011, 45, 99-112.	0.7	32
13	Local-scale variation of soil bacterial communities in ice-free regions of maritime Antarctica. Soil Biology and Biochemistry, 2019, 133, 165-173.	4.2	32
14	Comparative analysis of ammonia monooxygenase (amoA) genes in the water column and sediment–water interface of two lakes and the Baltic Sea. FEMS Microbiology Ecology, 2008, 66, 367-378.	1.3	30
15	Complete genome sequence of Pseudomonas antarctica PAMC 27494, a bacteriocin-producing psychrophile isolated from Antarctica. Journal of Biotechnology, 2017, 259, 15-18.	1.9	28
16	Bacterial communities in Antarctic lichens. Antarctic Science, 2016, 28, 455-461.	0.5	26
17	Comparative approach to capture bacterial diversity of coastal waters. Journal of Microbiology, 2011, 49, 729-740.	1.3	25
18	Bacterial diversity in ornithogenic soils compared to mineral soils on King George Island, Antarctica. Journal of Microbiology, 2012, 50, 1081-1085.	1.3	25

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19	Soil development and bacterial community shifts along the chronosequence of the Midtre Lovénbreen glacier foreland in Svalbard. Journal of Ecology and Environment, 2015, 38, 461-476.	1.6	19
20	Community analysis of betaproteobacterial ammonia-oxidizing bacteria using the amoCAB operon. Applied Microbiology and Biotechnology, 2009, 83, 175-188.	1.7	16
21	Methane production in the oxygenated water column of a perennially iceâ€covered Antarctic lake. Limnology and Oceanography, 2020, 65, 143-156.	1.6	14
22	Endophytic bacterial diversity of an Antarctic moss, <i>Sanionia uncinata</i> . Antarctic Science, 2013, 25, 51-54.	0.5	13
23	The latitudinal gradient in rock-inhabiting bacterial community compositions in Victoria Land, Antarctica. Science of the Total Environment, 2019, 657, 731-738.	3.9	12
24	Hydrogeological characteristics of groundwater and surface water associated with two small lake systems on King George Island, Antarctica. Journal of Hydrology, 2020, 590, 125537.	2.3	11
25	Complete Genome Sequence of Cryobacterium arcticum Strain PAMC 27867, Isolated from a Sedimentary Rock Sample in Northern Victoria Land, Antarctica. Genome Announcements, 2016, 4, .	0.8	10
26	Monitoring of soil bacterial community and some inoculated bacteria after prescribed fire in microcosm. Journal of Microbiology, 2004, 42, 285-91.	1.3	10
27	Complete genome sequence of Pedobacter cryoconitis PAMC 27485, a CRISPR-Cas system-containing psychrophile isolated from Antarctica. Journal of Biotechnology, 2016, 226, 74-75.	1.9	9
28	Complete Genome Sequence of a Psychrotolerant Denitrifying Bacterium, Janthinobacterium svalbardensis PAMC 27463. Genome Announcements, 2017, 5, .	0.8	8
29	Monitoring of bacterial community in a coniferous forest soil after a wildfire. Journal of Microbiology, 2004, 42, 278-84.	1.3	6
30	Statistical understanding for snow cover effects on near-surface ground temperature at the margin of maritime Antarctica, King George Island. Geoderma, 2022, 410, 115661.	2.3	4
31	Proposal to transferFlavobacterium oceanosedimentum Carty and Litchfield 1978 to the genusCurtobacterium asCurtobacterium oceanosedimentumcomb. nov FEMS Microbiology Letters, 2009, 296, 137-141.	0.7	3
32	Betaproteobacterial ammonia oxidizers in root zones of aquatic macrophytes. Fundamental and Applied Limnology, 2010, 177, 241-255.	0.4	3
33	Complete Genome Sequence of <i>Psychrobacter alimentarius</i> PAMC 27889, a Psychrophile Isolated from an Antarctic Rock Sample. Genome Announcements, 2016, 4, .	0.8	3
34	Effect of salinity on cyanobacterial community composition along a transect from Fuliya spring into the water of Lake Kinneret, Israel. Fundamental and Applied Limnology, 2013, 182, 99-107.	0.4	2
35	Draft Genome Sequence of the Psychrotolerant Bacterium <i>Methylobacterium</i> sp. Strain BTF04, Isolated from Freshwater in Antarctica. Microbiology Resource Announcements, 2020, 9, .	0.3	1
36	Draft Genome Sequence of the Chitin-Degrading Psychrotolerant Bacterium Pedobacter jejuensis TN23, Isolated from Antarctic Soil. Microbiology Resource Announcements, 2021, 10, e0052321.	0.3	0