Hanbyul Lee

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Biosurfactant-assisted bioremediation of crude oil by indigenous bacteria isolated from Taean beach sediment. Environmental Pollution, 2018, 241, 254-264.	3.7	128
2	Biotechnological procedures to select white rot fungi for the degradation of PAHs. Journal of Microbiological Methods, 2014, 97, 56-62.	0.7	116
3	Microbial community composition and PAHs removal potential of indigenous bacteria in oil contaminated sediment of Taean coast, Korea. Environmental Pollution, 2018, 234, 503-512.	3.7	111
4	Investigation of Marine-Derived Fungal Diversity and Their Exploitable Biological Activities. Marine Drugs, 2015, 13, 4137-4155.	2.2	77
5	Importance of functional diversity in assessing the recovery of the microbial community after the Hebei Spirit oil spill in Korea. Environment International, 2019, 128, 89-94.	4.8	35
6	Diversity of Wood-Inhabiting Polyporoid and Corticioid Fungi in Odaesan National Park, Korea. Mycobiology, 2016, 44, 217-236.	0.6	34
7	Enhanced removal of PAHs by Peniophora incarnata and ascertainment of its novel ligninolytic enzyme genes. Journal of Environmental Management, 2015, 164, 10-18.	3.8	32
8	Biofilm development of Bacillus siamensis ATKU1 on pristine short chain low-density polyethylene: A case study on microbe-microplastics interaction. Journal of Hazardous Materials, 2021, 409, 124516.	6.5	32
9	An integrative process for obtaining lipids and glucose from Chlorella vulgaris biomass with a single treatment of cell disruption. Algal Research, 2017, 27, 286-294.	2.4	31
10	Bioaccumulation of Polycyclic Aromatic Hydrocarbons (PAHs) by the Marine Clam, <i>Mactra veneriformis</i> , Chronically Exposed to Oil-Suspended Particulate Matter Aggregates. Environmental Science & Technology, 2018, 52, 7910-7920.	4.6	26
11	Heterologous expression of a new manganese-dependent peroxidase gene from Peniophora incarnata KUC8836 and its ability to remove anthracene in Saccharomyces cerevisiae. Journal of Bioscience and Bioengineering, 2016, 122, 716-721.	1.1	21
12	Investigation of Filamentous Fungi Producing Safe, Functional Water-Soluble Pigments. Mycobiology, 2018, 46, 269-277.	0.6	21
13	Fungal Diversity in Intertidal Mudflats and Abandoned Solar Salterns as a Source for Biological Resources. Marine Drugs, 2019, 17, 601.	2.2	20
14	Gemmobacter lutimaris sp. nov., a marine bacterium isolated from a tidal flat. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1676-1681.	0.8	19
15	Desorption and solubilization of anthracene by a rhamnolipid biosurfactant from <i>Rhodococcus fascians</i> . Water Environment Research, 2019, 91, 739-747.	1.3	18
16	Environmental drivers affecting the bacterial community of intertidal sediments in the Yellow Sea. Science of the Total Environment, 2021, 755, 142726.	3.9	18
17	The genus Arthrinium (Ascomycota, Sordariomycetes, Apiosporaceae) from marine habitats from Korea, with eight new species. IMA Fungus, 2021, 12, 13.	1.7	18
18	Echinicola sediminis sp. nov., a marine bacterium isolated from coastal sediment. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3351-3357.	0.8	17

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19	Isolation and Analysis of the Enzymatic Properties of Thermophilic Fungi from Compost. Mycobiology, 2014, 42, 181-184.	0.6	15
20	Fungal Diversity and Enzyme Activity Associated with the Macroalgae, <i>Agarum clathratum</i> . Mycobiology, 2019, 47, 50-58.	0.6	15
21	A proposed stepwise screening framework for the selection of polycyclic aromatic hydrocarbon (PAH)-degrading white rot fungi. Bioprocess and Biosystems Engineering, 2020, 43, 767-783.	1.7	15
22	Halo-tolerance of Marine-derived Fungi and their Enzymatic Properties. BioResources, 2015, 10, .	0.5	13
23	Comparison of the Diversity of Basidiomycetes from Dead Wood of the Manchurian fir (Abies) Tj ETQq1 1 0.7843 Microbial Ecology, 2015, 70, 634-645.	14 rgBT /C 1.4	Overlock 10 13
24	Optimization of Fungal Enzyme Production by <i>Trichoderma harzianum</i> KUC1716 through Surfactant-Induced Morphological Changes. Mycobiology, 2017, 45, 48-51.	0.6	13
25	Oceanimonas marisflavi sp. nov., a polycyclic aromatic hydrocarbon-degrading marine bacterium. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2990-2995.	0.8	13
26	Blastococcus litoris sp. nov., isolated from sea-tidal flat sediment. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3435-3440.	0.8	13
27	Marinobacter halodurans sp. nov., a halophilic bacterium isolated from sediment of a salt flat. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 6294-6300.	0.8	12
28	Wood Decay Fungi in South Korea: Polypores from Seoul. Mycobiology, 2014, 42, 140-146.	0.6	11
29	Recovery of the benthic bacterial community in coastal abandoned saltern requires over 35Âyears: A comparative case study in the Yellow Sea. Environment International, 2020, 135, 105412.	4.8	11
30	Successional Variation in the Soil Microbial Community in Odaesan National Park, Korea. Sustainability, 2020, 12, 4795.	1.6	11
31	Miniaturized enzyme production and development of micro-assays for cellulolytic and xylanolytic enzymes. Journal of Microbiological Methods, 2011, 86, 124-127.	0.7	10
32	Phylogenetic analysis of major molds inhabiting woods. Part 4. Genus <i>Alternaria</i> . Holzforschung, 2014, 68, 247-251.	0.9	10
33	Five New Records of Soil-Derived <i>Trichoderma</i> in Korea: <i>T. albolutescens</i> , <i>T. asperelloides</i> , <i>T. orientale</i> , <i>T. spirale</i> , and <i>T. tomentosum</i> . Mycobiology, 2017, 45, 1-8.	0.6	10
34	New Report of Three Unrecorded Species in <i>Trichoderma harzianum</i> Species Complex in Korea. Mycobiology, 2018, 46, 177-184.	0.6	10
35	Diversity of Trichoderma spp. in Marine Environments and Their Biological Potential for Sustainable Industrial Applications. Sustainability, 2020, 12, 4327.	1.6	10
36	Novosphingobium aureum sp. nov., a marine bacterium isolated from salt flat sediment. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	10

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37	Zobellella maritima sp. nov., a polycyclic aromatic hydrocarbon-degrading bacterium, isolated from beach sediment. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2279-2284.	0.8	10
38	Multiple evaluation of the potential toxic effects of sediments and biota collected from an oil-polluted area around Abu Ali Island, Saudi Arabia, Arabian Gulf. Ecotoxicology and Environmental Safety, 2019, 183, 109547.	2.9	9
39	Nocardioides litoris sp. nov., isolated from the Taean seashore. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2332-2336.	0.8	9
40	Screening for xylanase and β-xylosidase production from wood-inhabiting <i>Penicillium</i> strains for potential use in biotechnological applications. Holzforschung, 2012, 66, 267-271.	0.9	8
41	Comparative evaluation of bioremediation techniques on oil contaminated sediments in long-term recovery of benthic community health. Environmental Pollution, 2019, 252, 137-145.	3.7	8
42	Maribacter litoralis sp. nov. a marine bacterium isolated from seashore. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3471-3478.	0.8	8
43	Two Unrecorded <i>Apiospora</i> Species Isolated from Marine Substrates in Korea with Eight New Combinations (<i>A. piptatheri</i> and <i>A. rasikravindrae</i>). Mycobiology, 2022, 50, 46-54.	0.6	6
44	Influence of Tree Vegetation on Soil Microbial Communities in Temperate Forests and Their Potential as a Proactive Indicator of Vegetation Shift Due to Climate Change. Sustainability, 2020, 12, 10591.	1.6	5
45	Serinicoccus sediminis sp. nov., isolated from tidal flat sediment. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1998-2003.	0.8	5
46	Phylogenetic analysis of wood-inhabiting molds and assessment of soft-rot wood deterioration. Part 5. Genus <i>Aureobasidium</i> . Holzforschung, 2017, 71, 437-443.	0.9	4
47	Changes in Archaeal Community and Activity by the Invasion of Spartina anglica Along Soil Depth Profiles of a Coastal Wetland. Microbial Ecology, 2022, 83, 436-446.	1.4	4
48	Phylogenetic analysis of the genus Fusarium and their antifungal activity against wood-decay and sapstain fungi. Holzforschung, 2013, 67, 473-478.	0.9	3
49	Transcriptional analysis of genes encoding β-glucosidase of Schizophyllum commune KUC9397 under optimal conditions. Folia Microbiologica, 2017, 62, 191-196.	1.1	1