

Jaekyeong Song

List of Publications by Year in descending order

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13
papers

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1040056

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425
citing authors

#	ARTICLE	IF	CITATIONS
1	A native conjugative plasmid confers potential selective advantages to plant growth-promoting <i>Bacillus velezensis</i> strain GH1-13. <i>Communications Biology</i> , 2021, 4, 582.	4.4	3
2	Dispute cases related to pain management in Korea: analysis of Korea Medical Dispute Mediation and Arbitration Agency data. <i>Anesthesia and Pain Medicine</i> , 2020, 15, 96-102.	1.4	4
3	Structure and Mechanism of Surfactin Peptide from <i>Bacillus velezensis</i> Antagonistic to Fungi Plant Pathogens. <i>Bulletin of the Korean Chemical Society</i> , 2019, 40, 704-709.	1.9	26
4	Plant growth-promoting activity of beta-propeller protein YxaL secreted from <i>Bacillus velezensis</i> strain GH1-13. <i>PLoS ONE</i> , 2019, 14, e0207968.	2.5	16
5	Mass Cultivation and Characterization of Multifunctional <i>Bacillus velezensis</i> GH1-13. <i>Korean Journal of Organic Agriculture</i> , 2019, 27, 65-76.	0.1	3
6	Isolation and structural elucidation of pelgipeptin E, a novel pore-forming pelgipeptin analog from <i>Paenibacillus elgii</i> with low hemolytic activity. <i>Journal of Antibiotics</i> , 2018, 71, 1008-1017.	2.0	10
7	Complete genome sequence of <i>Bacillus velezensis</i> M75, a biocontrol agent against fungal plant pathogens, isolated from cotton waste. <i>Journal of Biotechnology</i> , 2017, 241, 112-115.	3.8	47
8	Effects of diet type, developmental stage, and gut compartment in the gut bacterial communities of two <i>Cerambycidae</i> species (Coleoptera). <i>Journal of Microbiology</i> , 2017, 55, 21-30.	2.8	66
9	The complete genome sequence of <i>Bacillus velezensis</i> strain GH1-13 reveals agriculturally beneficial properties and a unique plasmid. <i>Journal of Biotechnology</i> , 2017, 259, 221-227.	3.8	40
10	Characterization of Multifunctional <i>Bacillus</i> sp. GH1-13. <i>Nong'yag Gwahag Hoeji</i> , 2016, 20, 189-196.	0.5	11
11	Rapid discrimination of potato scab-causing <i>Streptomyces</i> species based on the RNase P RNA gene sequences. <i>Journal of Microbiology</i> , 2011, 49, 791-796.	2.8	0
12	Phylogenetic analysis of <i>Streptomyces</i> spp. isolated from potato scab lesions in Korea on the basis of 16S rRNA gene and 16S-23S rDNA internally transcribed spacer sequences. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2004, 54, 203-209.	1.7	70
13	Phylogenetic diversity of thermophilic actinomycetes and <i>Thermoactinomyces</i> spp. isolated from mushroom composts in Korea based on 16S rRNA gene sequence analysis. <i>FEMS Microbiology Letters</i> , 2001, 202, 97-102.	1.8	42