

Hyun-Hee Lee

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

380
citations

840776

11
h-index

794594

19
g-index

23
all docs

23
docs citations

23
times ranked

582
citing authors

#	ARTICLE	IF	CITATIONS
1	The In Vitro and In Planta Interspecies Interactions Among Rice-Pathogenic <i>Burkholderia</i> Species. <i>Plant Disease</i> , 2021, 105, 134-143.	1.4	7
2	Pan-Genome Analysis Reveals Host-Specific Functional Divergences in <i>Burkholderia gladioli</i> . <i>Microorganisms</i> , 2021, 9, 1123.	3.6	13
3	Type VI secretion systems of plant-pathogenic <i>Burkholderia glumae</i> BGR1 play a functionally distinct role in interspecies interactions and virulence. <i>Molecular Plant Pathology</i> , 2020, 21, 1055-1069.	4.2	20
4	Transcriptome analysis to understand the effects of the toxoflavin and tropolone produced by phytopathogenic <i>Burkholderia</i> on <i>Escherichia coli</i> . <i>Journal of Microbiology</i> , 2019, 57, 781-794.	2.8	8
5	Hepatoprotective Effect of Kombucha Tea in Rodent Model of Nonalcoholic Fatty Liver Disease/Nonalcoholic Steatohepatitis. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2369.	4.1	26
6	Stress Tolerance and Virulence-Related Roles of Lipopolysaccharide in <i>Burkholderia glumae</i> . <i>Plant Pathology Journal</i> , 2019, 35, 445-458.	1.7	16
7	Characterization of <i>Burkholderia glumae</i> Putative Virulence Factor 11 (PVF11) via Yeast Two-Hybrid Interaction and Phenotypic Analysis. <i>Plant Pathology Journal</i> , 2019, 35, 280-286.	1.7	3
8	Roles of three FurA paralogs in the regulation of genes pertaining to peroxide defense in <i>Mycobacterium smegmatis</i> mc ² 155. <i>Molecular Microbiology</i> , 2018, 108, 661-682.	2.5	14
9	Cooperative interactions between seed-borne bacterial and air-borne fungal pathogens on rice. <i>Nature Communications</i> , 2018, 9, 31.	12.8	46
10	Profiling of glucose-induced transcription in <i>Sulfolobus acidocaldarius</i> DSM 639. <i>Genes and Genomics</i> , 2018, 40, 1157-1167.	1.4	3
11	Genomics-based Sensitive and Specific Novel Primers for Simultaneous Detection of <i>Burkholderia glumae</i> and <i>Burkholderia gladioli</i> in Rice Seeds. <i>Plant Pathology Journal</i> , 2018, 34, 490-498.	1.7	8
12	Specific and Sensitive Primers Developed by Comparative Genomics to Detect Bacterial Pathogens in Grains. <i>Plant Pathology Journal</i> , 2018, 34, 104-112.	1.7	10
13	Genome-Wide Analysis of Type VI System Clusters and Effectors in <i>Burkholderia</i> Species. <i>Plant Pathology Journal</i> , 2018, 34, 11-22.	1.7	20
14	The Roles of Two hfq Genes in the Virulence and Stress Resistance of <i>Burkholderia glumae</i> . <i>Plant Pathology Journal</i> , 2018, 34, 412-425.	1.7	20
15	Development of High Cordycepin-Producing <i>Cordyceps militaris</i> Strains. <i>Mycobiology</i> , 2017, 45, 31-38.	1.7	31
16	Characterization of Newly Bred <i>Cordyceps militaris</i> Strains for Higher Production of Cordycepin through HPLC and URP-PCR Analysis. <i>Journal of Microbiology and Biotechnology</i> , 2017, 27, 1223-1232.	2.1	11
17	Comparative Genome Analysis of <i>Rathayibacter tritici</i> NCPPB 1953 with <i>Rathayibacter toxicus</i> Strains Can Facilitate Studies on Mechanisms of Nematode Association and Host Infection. <i>Plant Pathology Journal</i> , 2017, 33, 370-381.	1.7	6
18	Computational Identification and Comparative Analysis of Secreted and Transmembrane Proteins in Six <i>Burkholderia</i> Species. <i>Plant Pathology Journal</i> , 2017, 33, 148-162.	1.7	6

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19	MicroRNA Expression Profiling in CCl4-Induced Liver Fibrosis of <i>Mus musculus</i> . <i>International Journal of Molecular Sciences</i> , 2016, 17, 961.	4.1	32
20	Understanding the direction of evolution in <i>Burkholderia glumae</i> through comparative genomics. <i>Current Genetics</i> , 2016, 62, 115-123.	1.7	15
21	Comparative Analyses of Tomato yellow leaf curl virus C4 Protein-Interacting Host Proteins in Healthy and Infected Tomato Tissues. <i>Plant Pathology Journal</i> , 2016, 32, 377-387.	1.7	9
22	Comparative genome analysis of rice-pathogenic <i>Burkholderia</i> provides insight into capacity to adapt to different environments and hosts. <i>BMC Genomics</i> , 2015, 16, 349.	2.8	45
23	Complete genome sequence of <i>Bacillus velezensis</i> G341, a strain with a broad inhibitory spectrum against plant pathogens. <i>Journal of Biotechnology</i> , 2015, 211, 97-98.	3.8	11