

Gung Pyo Lee

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8373857/publications.pdf>

Version: 2024-02-01

10
papers

126
citations

1478505

6
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

138
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of QTLs and Candidate Genes for Days to Heading in Rice Recombinant Inbred Lines. <i>Genes</i> , 2020, 11, 957.	2.4	6
2	Exploring molecular markers and candidate genes responsible for watermelon dwarfism. <i>Horticulture Environment and Biotechnology</i> , 2020, 61, 173-182.	2.1	5
3	An evolutionarily conserved non-synonymous SNP in a leucine-rich repeat domain determines anthracnose resistance in watermelon. <i>Theoretical and Applied Genetics</i> , 2019, 132, 473-488.	3.6	17
4	Identification of the pleiotropic function of TOUSLED kinase in tomato (<i>Solanum lycopersicum</i> L.) using a Cucumber mosaic virus-based vector. <i>Horticulture Environment and Biotechnology</i> , 2018, 59, 105-114.	2.1	4
5	De novo-based transcriptome profiling of male-sterile and fertile watermelon lines. <i>PLoS ONE</i> , 2017, 12, e0187147.	2.5	7
6	Identification of the subgenomic promoter of the coat protein gene of cucumber fruit mottle mosaic virus and development of a heterologous expression vector. <i>Archives of Virology</i> , 2016, 161, 1527-1538.	2.1	13
7	Transcriptome profiling of differentially expressed genes in floral buds and flowers of male sterile and fertile lines in watermelon. <i>BMC Genomics</i> , 2015, 16, 914.	2.8	54
8	Construction of a genetic linkage map using a frame set of simple sequence repeat and high-resolution melting markers for watermelon (<i>Citrullus</i> spp.). <i>Horticulture Environment and Biotechnology</i> , 2015, 56, 669-676.	2.1	9
9	Infectivity and complete nucleotide sequence of cucumber fruit mottle mosaic virus isolate Cm cDNA. <i>Archives of Virology</i> , 2014, 159, 1807-1811.	2.1	7
10	Brief report: genome sequence and construction of an infectious cDNA clone of Ribgrass mosaic virus from Chinese cabbage in Korea. <i>Virus Genes</i> , 2012, 44, 345-348.	1.6	4