

Hai Yan Fan

List of Publications by Year in descending order

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papers

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citations

1163117

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144
citing authors

#	ARTICLE	IF	CITATIONS
1	Regulation of Growth and Salt Resistance in Cucumber Seedlings by Hydrogen-Rich Water. <i>Journal of Plant Growth Regulation</i> , 2023, 42, 134-153.	5.1	10
2	Nucleotide-Binding Leucine-Rich Repeat Genes CsRSF1 and CsRSF2 Are Positive Modulators in the <i>Cucumis sativus</i> Defense Response to <i>Sphaerotheca fuliginea</i> . <i>International Journal of Molecular Sciences</i> , 2021, 22, 3986.	4.1	15
3	Genetic diversity of diazotrophs and total bacteria in the phyllosphere of <i>Pyrus serotina</i> , <i>Prunus armeniaca</i> , <i>Prunus avium</i> , and <i>Vitis vinifera</i> . <i>Canadian Journal of Microbiology</i> , 2019, 65, 642-652.	1.7	10
4	Mildew Resistance Locus O Genes CsMLO1 and CsMLO2 Are Negative Modulators of the <i>Cucumis sativus</i> Defense Response to <i>Corynespora cassiicola</i> . <i>International Journal of Molecular Sciences</i> , 2019, 20, 4793.	4.1	9
5	Functional Identification of <i>Corynespora cassiicola</i> -Responsive miRNAs and Their Targets in Cucumber. <i>Frontiers in Plant Science</i> , 2019, 10, 668.	3.6	9
6	Cucumber Mildew Resistance Locus O Interacts with Calmodulin and Regulates Plant Cell Death Associated with Plant Immunity. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2995.	4.1	11
7	Transcriptome and miRNA analyses of the response to <i>Corynespora cassiicola</i> in cucumber. <i>Scientific Reports</i> , 2018, 8, 7798.	3.3	43
8	The Two Translationally Controlled Tumor Protein Genes, CsTCTP1 and CsTCTP2, Are Negative Modulators in the <i>Cucumis sativus</i> Defense Response to <i>Sphaerotheca fuliginea</i> . <i>Frontiers in Plant Science</i> , 2018, 9, 544.	3.6	21
9	Molecular characterization, expression analysis and heterologous expression of two translationally controlled tumor protein genes from <i>Cucumis sativus</i> . <i>PLoS ONE</i> , 2017, 12, e0184872.	2.5	7
10	Comparative proteomic analysis of cucumber roots infected by <i>Fusarium oxysporum</i> f. sp. <i>cucumerium</i> Owen. <i>Physiological and Molecular Plant Pathology</i> , 2016, 96, 77-84.	2.5	17
11	A comparative cell wall proteomic analysis of cucumber leaves under <i>Sphaerotheca fuliginea</i> stress. <i>Acta Physiologiae Plantarum</i> , 2016, 38, 1.	2.1	9
12	Draft Genome Sequence of <i>Paenibacillus polymyxa</i> EBL06, a Plant Growth-Promoting Bacterium Isolated from Wheat Phyllosphere. <i>Genome Announcements</i> , 2015, 3, .	0.8	1
13	Proteomic analysis of <i>Cucumis sativus</i> cotyledons after glucohexaose treatment as a part of ROS accumulation related resistance mechanism. <i>Proteome Science</i> , 2014, 12, 34.	1.7	5
14	Photosynthesis Regulation by Glucohexaose Through Redox Changes in <i>Cucumis sativus</i> . <i>Journal of Plant Growth Regulation</i> , 2014, 33, 571-578.	5.1	2
15	Proteome-level investigation of <i>Cucumis sativus</i> -derived resistance to <i>Sphaerotheca fuliginea</i> . <i>Acta Physiologiae Plantarum</i> , 2014, 36, 1781-1791.	2.1	12