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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	iTAK: A Program for Genome-wide Prediction andÂClassification of Plant Transcription Factors, Transcriptional Regulators, and Protein Kinases. Molecular Plant, 2016, 9, 1667-1670.	8.3	735
2	Transcriptomics-based screen for genes induced by flagellin and repressed by pathogen effectors identifies a cell wall-associated kinase involved in plant immunity. Genome Biology, 2013, 14, R139.	9.6	137
3	UV-C treatment affects the expression and activity of defense genes in strawberry fruit (Fragaria×ananassa, Duch.). Postharvest Biology and Technology, 2011, 59, 94-102.	6.0	116
4	UV-C irradiation delays strawberry fruit softening and modifies the expression of genes involved in cell wall degradation. Postharvest Biology and Technology, 2009, 51, 141-148.	6.0	113
5	Use of RNA-seq data to identify and validate RT-qPCR reference genes for studying the tomato-Pseudomonas pathosystem. Scientific Reports, 2017, 7, 44905.	3.3	85
6	Transcriptomic analysis reveals tomato genes whose expression is induced specifically during effector-triggered immunity and identifies the Epk1 protein kinase which is required for the host response to three bacterial effector proteins. Genome Biology, 2014, 15, 492.	8.8	75
7	Tomato Wall-Associated Kinase SlWak1 Depends on Fls2/Fls3 to Promote Apoplastic Immune Responses to <i>Pseudomonas syringae</i> . Plant Physiology, 2020, 183, 1869-1882.	4.8	52
8	Cloning of FaPAL6 gene from strawberry fruit and characterization of its expression and enzymatic activity in two cultivars with different anthocyanin accumulation. Plant Science, 2011, 181, 111-118.	3.6	34
9	Transcriptome-based identification and validation of reference genes for plant-bacteria interaction studies using Nicotiana benthamiana. Scientific Reports, 2019, 9, 1632.	3.3	34
10	Heat treatments and expansin gene expression in strawberry fruit. Scientia Horticulturae, 2011, 130, 775-780.	3.6	26
11	The Tomato Kinase Pti1 Contributes to Production of Reactive Oxygen Species in Response to Two Flagellin-Derived Peptides and Promotes Resistance to <i>Pseudomonas syringae</i> Infection. Molecular Plant-Microbe Interactions, 2017, 30, 725-738.	2.6	22
12	WRKY22 and WRKY25 transcription factors are positive regulators of defense responses in Nicotiana benthamiana. Plant Molecular Biology, 2021, 105, 65-82.	3.9	19
13	A novel method of transcriptome interpretation reveals a quantitative suppressive effect on tomato immune signaling by two domains in a single pathogen effector protein. BMC Genomics, 2016, 17, 229.	2.8	9
14	Genome-wide analysis uncovers tomato leaf IncRNAs transcriptionally active upon Pseudomonas syringae pv. tomato challenge. Scientific Reports, 2021, 11, 24523.	3.3	8
15	Nicotiana benthamiana, A Popular Model for Genome Evolution and Plant–Pathogen Interactions. Compendium of Plant Genomes, 2020, , 231-247.	0.5	6