Cunwu Zuo

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9607974/publications.pdf

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16	204	1163117 8	1199594 12
papers	citations	h-index	g-index

16	16	16	223
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Effects of Shading on the Synthesis of Volatile Organic Compounds in â€˜Marselanâ€™ Grape Berries (Vitis) Tj ETQq1 1 0.784314 rgBT	5.1	9
2	Cyclic nucleotide gated channel genes (CNGCs) in Rosaceae: genome-wide annotation, evolution and the roles on Valsa canker resistance. Plant Cell Reports, 2021, 40, 2369-2382.	5.6	10
3	Genome-Wide Analysis of the Apple (Malus domestica) Cysteine-Rich Receptor-Like Kinase (CRK) Family: Annotation, Genomic Organization, and Expression Profiles in Response to Fungal Infection. Plant Molecular Biology Reporter, 2020, 38, 14-24.	1.8	20
4	RNA Sequencing Reveals That Both Abiotic and Biotic Stress-Responsive Genes are Induced during Expression of Steroidal Glycoalkaloid in Potato Tuber Subjected to Light Exposure. Genes, 2019, 10, 920.	2.4	8
5	Genome-Wide Identification and Expression Analysis of GA2ox, GA3ox, and GA20ox Are Related to Gibberellin Oxidase Genes in Grape (Vitis Vinifera L.). Genes, 2019, 10, 680.	2.4	44
6	Genome-wide annotation and expression responses to biotic stresses of the WALL-ASSOCIATED KINASE - RECEPTOR-LIKE KINASE (WAK-RLK) gene family in Apple (Malus domestica). European Journal of Plant Pathology, 2019, 153, 771-785.	1.7	20
7	The mechanism of color fading in sunburned apple peel. Acta Physiologiae Plantarum, 2019, 41, 1.	2.1	7
8	Genome-wide annotation and expression responses to biotic stresses of the WALL-ASSOCIATED KINASE - RECEPTOR-LIKE KINASE (WAK-RLK) gene family in Apple (Malus domestica). , 2019, 153, 771.		1
9	Transcriptome analysis revealed glucose application affects plant hormone signal transduction pathway in â€œRed Globeâ€•grape plantlets. Plant Growth Regulation, 2018, 84, 45-56.	3.4	18
10	Genome-Wide Identification and Expression Analysis of the CrRLK1L Gene Family in Apple (Malus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.8	18
11	RNA sequencing analysis provides new insights into dynamic molecular responses to Valsa mali pathogenicity in apple â€˜Changfu No. 2â€™. Tree Genetics and Genomes, 2018, 14, 1.	1.6	6
12	Significant and unique changes in phosphorylation levels of four phosphoproteins in two apple rootstock genotypes under drought stress. Molecular Genetics and Genomics, 2017, 292, 1307-1322.	2.1	13
13	RNA Sequencing Reveals that Endoplasmic Reticulum Stress and Disruption of Membrane Integrity Underlie Dimethyl Trisulfide Toxicity against Fusarium oxysporum f. sp. cubense Tropical Race 4. Frontiers in Microbiology, 2017, 8, 1365.	3.5	25
14	Transcriptomic Analysis Revealed Hormone-Related and Receptor-Like Kinase Genes Involved in Wound Healing of â€˜Duliâ€™ and its Resistance to Valsa Pyri. Plant Molecular Biology Reporter, 0, , 1.	1.8	1
15	Transcriptome Profile in a Susceptible Pear â€˜Zaosuâ€™ (Pyrus bretschneideri Rehd.)â€“Valsa pyri Interaction. Journal of Plant Growth Regulation, 0, , 1.	5.1	0
16	Transcriptomic analysis reveals that cell wall- and hypersensitive response (HR)-related genes are involved in the responses of apple to Valsa mali. Plant Biotechnology Reports, 0, , .	1.5	4