

Houbin Chen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1484214/publications.pdf>

Version: 2024-02-01

29
papers

577
citations

623734

14
h-index

642732

23
g-index

30
all docs

30
docs citations

30
times ranked

546
citing authors

#	ARTICLE	IF	CITATIONS
1	Two divergent haplotypes from a highly heterozygous lychee genome suggest independent domestication events for early and late-maturing cultivars. <i>Nature Genetics</i> , 2022, 54, 73-83.	21.4	88
2	Changes in Homogalacturonan Metabolism in Banana Peel during Fruit Development and Ripening. <i>International Journal of Molecular Sciences</i> , 2022, 23, 243.	4.1	5
3	Identification of Chilling Accumulation-Associated Genes for Litchi Flowering by Transcriptome-Based Genome-Wide Association Studies. <i>Frontiers in Plant Science</i> , 2022, 13, 819188.	3.6	1
4	Genome-Wide Identification of Banana Csl Gene Family and Their Different Responses to Low Temperature between Chilling-Sensitive and Tolerant Cultivars. <i>Plants</i> , 2021, 10, 122.	3.5	12
5	Metabolomics Analysis of Litchi Leaves during Floral Induction Reveals Metabolic Improvement by Stem Girdling. <i>Molecules</i> , 2021, 26, 4048.	3.8	5
6	Genome-Wide Identification and Expression Analysis of MADS-Box Family Genes in Litchi (<i>Litchi</i>) Tj ETQq0 0 0 rgBT /Qverlock 10 Tf 50 5	3.5	13
7	Development of molecular markers based on the promoter difference of LcFT1 to discriminate easy- and difficult-flowering litchi germplasm resources and its application in crossbreeding. <i>BMC Plant Biology</i> , 2021, 21, 539.	3.6	6
8	Acceleration of Carbon Fixation in Chilling-Sensitive Banana under Mild and Moderate Chilling Stresses. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9326.	4.1	1
9	Genome-Wide Transcriptomic Analysis Reveals a Regulatory Network of Oxidative Stress-Induced Flowering Signals Produced in Litchi Leaves. <i>Genes</i> , 2020, 11, 324.	2.4	9
10	Genome-wide analyses of banana fasciclin-like AGP genes and their differential expression under low-temperature stress in chilling sensitive and tolerant cultivars. <i>Plant Cell Reports</i> , 2020, 39, 693-708.	5.6	17
11	Identification of Genes Involved in Low Temperature-Induced Senescence of Panicle Leaf in Litchi chinensis. <i>Genes</i> , 2019, 10, 111.	2.4	2
12	Genome-wide transcriptome analysis reveals the molecular mechanism of high temperature-induced floral abortion in Litchi chinensis. <i>BMC Genomics</i> , 2019, 20, 127.	2.8	15
13	Functional analysis of a homologue of the FLORICAULA/LEAFY gene in litchi (<i>Litchi chinensis</i> Sonn.) revealing its significance in early flowering process. <i>Genes and Genomics</i> , 2018, 40, 1259-1267.	1.4	10
14	Litchi Flowering is Regulated by Expression of Short Vegetative Phase Genes. <i>Journal of the American Society for Horticultural Science</i> , 2018, 143, 101-109.	1.0	4
15	Comparative Digital Gene Expression Analysis of Tissue-Cultured Plantlets of Highly Resistant and Susceptible Banana Cultivars in Response to <i>Fusarium oxysporum</i> . <i>International Journal of Molecular Sciences</i> , 2018, 19, 350.	4.1	24
16	Reactive oxygen species and nitric oxide induce senescence of rudimentary leaves and the expression profiles of the related genes in Litchi chinensis. <i>Horticulture Research</i> , 2018, 5, 23.	6.3	21
17	LcMCII-1 is involved in the ROS-dependent senescence of the rudimentary leaves of Litchi chinensis. <i>Plant Cell Reports</i> , 2017, 36, 89-102.	5.6	11
18	Expression and distribution of extensins and AGPs in susceptible and resistant banana cultivars in response to wounding and <i>Fusarium oxysporum</i> . <i>Scientific Reports</i> , 2017, 7, 42400.	3.3	30

#	ARTICLE	IF	CITATIONS
19	RNA-seq analysis of apical meristem reveals integrative regulatory network of ROS and chilling potentially related to flowering in Litchi chinensis. <i>Scientific Reports</i> , 2017, 7, 10183.	3.3	15
20	Transcriptome profiling of litchi leaves in response to low temperature reveals candidate regulatory genes and key metabolic events during floral induction. <i>BMC Genomics</i> , 2017, 18, 363.	2.8	13
21	Integrative effect of drought and low temperature on litchi (<i>Litchi chinensis</i> Sonn.) floral initiation revealed by dynamic genome-wide transcriptome analysis. <i>Scientific Reports</i> , 2016, 6, 32005.	3.3	27
22	Variable content and distribution of arabinogalactan proteins in banana (<i>Musa</i> spp.) under low temperature stress. <i>Frontiers in Plant Science</i> , 2015, 6, 353.	3.6	26
23	Promoter difference of LcFT1 is a leading cause of natural variation of flowering timing in different litchi cultivars (<i>Litchi chinensis</i> Sonn.). <i>Plant Science</i> , 2015, 241, 128-137.	3.6	31
24	De novo transcriptome assembly for rudimentary leaves in Litchi chinensis Sonn. and identification of differentially expressed genes in response to reactive oxygen species. <i>BMC Genomics</i> , 2014, 15, 805.	2.8	65
25	A systematic comparison of embryogenic and non-embryogenic cells of banana (<i>Musa</i> spp. AAA): Ultrastructural, biochemical and cell wall component analyses. <i>Scientia Horticulturae</i> , 2013, 159, 178-185.	3.6	9
26	Wound-induced pectin methylesterases enhance banana (<i>Musa</i> spp. AAA) susceptibility to <i>Fusarium oxysporum</i> f. sp. <i>cubense</i> . <i>Journal of Experimental Botany</i> , 2013, 64, 2219-2229.	4.8	33
27	Histological changes and differences in activities of some antioxidant enzymes and hydrogen peroxide content during somatic embryogenesis of <i>Musa</i> AAA cv. Yueyoukang 1. <i>Scientia Horticulturae</i> , 2012, 144, 87-92.	3.6	22
28	Ultrastructural changes and the distribution of arabinogalactan proteins during somatic embryogenesis of banana (<i>Musa</i> spp. AAA cv. "Yueyoukang 1"™). <i>Physiologia Plantarum</i> , 2011, 142, 372-389.	5.2	43
29	Rudimentary leaf abortion with the development of panicle in litchi: Changes in ultrastructure, antioxidant enzymes and phytohormones. <i>Scientia Horticulturae</i> , 2008, 117, 288-296.	3.6	18