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An improved fruit transcriptome and the identification of the candidate genes involved in fruit abscission induced by carbohydrate stress in litchi

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37	Cloning, Functional Characterization, and Catalytic Mechanism of a Bergaptol O-Methyltransferase from Peucedanum praeruptorum Dunn. <i>Frontiers in Plant Science</i> , 2016 , 7, 722	6.2	14
36	Validation of Reference Genes for RT-qPCR Studies of Gene Expression in Preharvest and Postharvest Longan Fruits under Different Experimental Conditions. <i>Frontiers in Plant Science</i> , 2016 , 7, 780	6.2	23
35	Identification and molecular characterization of an IDA-like gene from litchi, LcIDL1, whose ectopic expression promotes floral organ abscission in Arabidopsis. <i>Scientific Reports</i> , 2016 , 6, 37135	4.9	32
34	Transcriptional changes during ovule development in two genotypes of litchi (Litchi chinensis Sonn.) with contrast in seed size. <i>Scientific Reports</i> , 2016 , 6, 36304	4.9	7
33	Advances in Genetic Transformation of Litchi. 2017 , 421-436		1
32	Litchi Fruit Set, Development, and Maturation. 2017 , 1-30		4
31	Abiotic Stress Management in Fruit Crop Litchi chinensis. 2017 , 243-263		O
30	Genome-Wide Identification of Histone Modifiers and Their Expression Patterns during Fruit Abscission in Litchi. <i>Frontiers in Plant Science</i> , 2017 , 8, 639	6.2	24
29	Genetics and Breeding of Fruit Crops in the Sapindaceae Family: Lychee (Litchi chinensis Sonn.) and Longan (Dimocarpus longan Lour.). 2018 , 953-973		1
28	Molecular and related approaches to litchi improvement Ihistorical perspective and future trends. Journal of Horticultural Science and Biotechnology, 2019 , 94, 693-702	1.9	3
27	Molecular and Hormonal Aspects of Drought-Triggered Flower Shedding in Yellow Lupine. International Journal of Molecular Sciences, 2019, 20,	6.3	10
26	Screening the Reference Genes for Quantitative Gene Expression by RT-qPCR During SE Initial Dedifferentiation in Four Cultivars that Have Different SE Capability. <i>Genes</i> , 2019 , 10,	4.2	5
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23	Involvement of HD-ZIP I transcription factors LcHB2 and LcHB3 in fruitlet abscission by promoting transcription of genes related to the biosynthesis of ethylene and ABA in litchi. <i>Tree Physiology</i> , 2019 , 39, 1600-1613	4.2	14
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20	LcEIL2/3 are involved in fruitlet abscission via activating genes related to ethylene biosynthesis and cell wall remodeling in litchi. <i>Plant Journal</i> , 2020 , 103, 1338-1350	6.9	7
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18	Comparative RNA sequencing based transcriptome profiling of regular bearing and alternate bearing mango (Mangifera indica L.) varieties reveals novel insights into the regulatory mechanisms underlying alternate bearing. <i>Biotechnology Letters</i> , 2020 , 42, 1035-1050	3	8
17	Molecular Events Involved in Fruitlet Abscission in Litchi. <i>Plants</i> , 2020 , 9,	4.5	8
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