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Genome-wide digital transcript analysis of putative fruitlet abscission related genes regulated by ethephon in litchi

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#	Paper	IF	Citations
48	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
47	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
46	Ethylene-induced transcriptional and hormonal responses at the onset of sugarcane ripening. <i>Scientific Reports</i> , 2017 , 7, 43364	4.9	30
45	Distinct organ-specific and temporal expression profiles of auxin-related genes during mango fruitlet drop. <i>Plant Physiology and Biochemistry</i> , 2017 , 115, 439-448	5.4	4
44	Biotechnological Interventions in Litchi (Litchi chinensis Sonn.) for the Improvement of Fruit Quality and Postharvest Storage. 2017 , 101-136		1
43	Litchi Fruit Set, Development, and Maturation. 2017 , 1-30		4
42	Abiotic Stress Management in Fruit Crop Litchi chinensis. 2017 , 243-263		O
41	Strategies to Retard Postharvest Pericarp Browning in Lychee Fruit. 2017, 265-279		
40	Ethephon induced oxidative stress in the olive leaf abscission zone enables development of a selective abscission compound. <i>BMC Plant Biology</i> , 2017 , 17, 87	5.3	19
39	Cell Wall Remodeling in Abscission Zone Cells during Ethylene-Promoted Fruit Abscission in Citrus. <i>Frontiers in Plant Science</i> , 2017 , 8, 126	6.2	42
38	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
37	Genetics and Breeding of Fruit Crops in the Sapindaceae Family: Lychee (Litchi chinensis Sonn.) and Longan (Dimocarpus longan Lour.). 2018 , 953-973		1
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35	The Tomato Hybrid Proline-rich Protein regulates the abscission zone competence to respond to ethylene signals. <i>Horticulture Research</i> , 2018 , 5, 28	7.7	10
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33	Metabolomic and transcriptomic profiling of three types of litchi pericarps reveals that changes in the hormone balance constitute the molecular basis of the fruit cracking susceptibility of Litchi chinensis cv. Baitangying. <i>Molecular Biology Reports</i> , 2019 , 46, 5295-5308	2.8	5
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31	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
30	The HD-Zip transcription factor LcHB2 regulates litchi fruit abscission through the activation of two cellulase genes. <i>Journal of Experimental Botany</i> , 2019 , 70, 5189-5203	7	14
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28	Re-evaluation of the ethylene-dependent and -independent pathways in the regulation of floral and organ abscission. <i>Journal of Experimental Botany</i> , 2019 , 70, 1461-1467	7	39
27	Transcriptome and Hormone Analyses Revealed Insights into Hormonal and Vesicle Trafficking Regulation among Fruit Tissues in Late Development. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	4
26	Spatio-temporal immunolocalization of extensin protein and hemicellulose polysaccharides during olive fruit abscission. <i>Planta</i> , 2020 , 252, 32	4.7	1
25	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
24	Cell Wall Composition and Ultrastructural Immunolocalization of Pectin and Arabinogalactan Protein during Olea europaea L. Fruit Abscission. <i>Plant and Cell Physiology</i> , 2020 , 61, 814-825	4.9	3
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21	KNOX protein KNAT1 regulates fruitlet abscission in litchi by repressing ethylene biosynthetic genes. <i>Journal of Experimental Botany</i> , 2020 , 71, 4069-4082	7	11
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