

Laurent Torregrosa

List of Publications by Year in descending order

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33
papers

2,924
citations

304743

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414414

32
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docs citations

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times ranked

2937
citing authors

#	ARTICLE	IF	CITATIONS
1	Ectopic Expression of VvMybPA2 Promotes Proanthocyanidin Biosynthesis in Grapevine and Suggests Additional Targets in the Pathway. <i>Plant Physiology</i> , 2009, 149, 1028-1041.	4.8	354
2	Grapevine MATE-Type Proteins Act as Vacuolar H ⁺ -Dependent Acylated Anthocyanin Transporters. <i>Plant Physiology</i> , 2009, 150, 402-415.	4.8	305
3	<i>In vivo</i> grapevine anthocyanin transport involves vesicle-mediated trafficking and the contribution of anthoMATE transporters and GST. <i>Plant Journal</i> , 2011, 67, 960-970.	5.7	222
4	Ectopic expression of VvmybA1 in grapevine activates a narrow set of genes involved in anthocyanin synthesis and transport. <i>Plant Molecular Biology</i> , 2009, 69, 633-648.	3.9	202
5	Temperature desynchronizes sugar and organic acid metabolism in ripening grapevine fruits and remodels their transcriptome. <i>BMC Plant Biology</i> , 2016, 16, 164.	3.6	192
6	Day and night heat stress trigger different transcriptomic responses in green and ripening grapevine (<i>Vitis vinifera</i>) fruit. <i>BMC Plant Biology</i> , 2014, 14, 108.	3.6	170
7	Genetic dissection of a TIR-NB-LRR locus from the wild North American grapevine species <i>Muscadinia rotundifolia</i> identifies paralogous genes conferring resistance to major fungal and oomycete pathogens in cultivated grapevine. <i>Plant Journal</i> , 2013, 76, 661-674.	5.7	152
8	A negative MYB regulator of proanthocyanidin accumulation, identified through expression quantitative locus mapping in the grape berry. <i>New Phytologist</i> , 2014, 201, 795-809.	7.3	144
9	A grapevine Shaker inward K ⁺ channel activated by the calcineurin B-like calcium sensor 14-3-3 protein kinase CIPK23 network is expressed in grape berries under drought stress conditions. <i>Plant Journal</i> , 2010, 61, 58-69.	5.7	135
10	Transposon-induced gene activation as a mechanism generating cluster shape somatic variation in grapevine. <i>Plant Journal</i> , 2010, 61, 545-557.	5.7	116
11	Impact of agronomic practices on grape aroma composition: a review. <i>Journal of the Science of Food and Agriculture</i> , 2019, 99, 975-985.	3.5	111
12	Identification of grapevine MLO gene candidates involved in susceptibility to powdery mildew. <i>Functional Plant Biology</i> , 2008, 35, 1255.	2.1	101
13	The grape microvine - a model system for rapid forward and reverse genetics of grapevines. <i>Plant Journal</i> , 2010, 62, no-no.	5.7	85
14	Involvement of ethylene signalling in a non-climacteric fruit: new elements regarding the regulation of ADH expression in grapevine. <i>Journal of Experimental Botany</i> , 2004, 55, 2235-2240.	4.8	84
15	Identification of genes associated with flesh morphogenesis during grapevine fruit development. <i>Plant Molecular Biology</i> , 2007, 63, 307-323.	3.9	78
16	Effects of genetic manipulation of alcohol dehydrogenase levels on the response to stress and the synthesis of secondary metabolites in grapevine leaves. <i>Journal of Experimental Botany</i> , 2006, 57, 91-99.	4.8	66
17	Identification of stable QTLs for vegetative and reproductive traits in the microvine (<i>Vitis vinifera</i> L.) using the 18K Infinium chip. <i>BMC Plant Biology</i> , 2015, 15, 205.	3.6	65
18	A Grapevine Gene Encoding a Guard Cell K ⁺ Channel Displays Developmental Regulation in the Grapevine Berry. <i>Plant Physiology</i> , 2002, 128, 564-577.	4.8	53

#	ARTICLE	IF	CITATIONS
19	Vitis vinifera L. Fruit Diversity to Breed Varieties Anticipating Climate Changes. <i>Frontiers in Plant Science</i> , 2018, 9, 455.	3.6	51
20	Manipulation of VvAdh to investigate its function in grape berry development. <i>Plant Science</i> , 2008, 174, 149-155.	3.6	36
21	Developmental, molecular and genetic studies on grapevine response to temperature open breeding strategies for adaptation to warming. <i>Oeno One</i> , 2017, 51, 155-165.	1.4	32
22	The kinetics of grape ripening revisited through berry density sorting. <i>Oeno One</i> , 2019, 53, .	1.4	30
23	First quantitative assessment of growth, sugar accumulation and malate breakdown in a single ripening berry. <i>Oeno One</i> , 2020, 54, 1077-1092.	1.4	28
24	Developmental, molecular and genetic studies on grapevine response to temperature open breeding strategies for adaptation to warming. <i>Oeno One</i> , 2017, 51, 155.	1.4	19
25	Transcripts switched off at the stop of phloem unloading highlight the energy efficiency of sugar import in the ripening <i>V. vinifera</i> fruit. <i>Horticulture Research</i> , 2021, 8, 193.	6.3	15
26	Vitis vinifera L. Diversity for Cations and Acidity Is Suitable for Breeding Fruits Coping With Climate Warming. <i>Frontiers in Plant Science</i> , 2020, 11, 01175.	3.6	14
27	Transcriptome analyses suggest that changes in fungal endophyte lifestyle could be involved in grapevine bud necrosis. <i>Scientific Reports</i> , 2020, 10, 9514.	3.3	14
28	The reduction of plant sink/source does not systematically improve the metabolic composition of Vitis vinifera white fruit. <i>Food Chemistry</i> , 2021, 345, 128825.	8.2	11
29	The <i>Microvine</i> , a plant model to study the effect of vine shoot extract on the accumulation of glycosylated aroma precursors in grapes. <i>Journal of the Science of Food and Agriculture</i> , 2018, 98, 3031-3040.	3.5	10
30	The application of ozonated water rearranges the Vitis vinifera L. leaf and berry transcriptomes eliciting defence and antioxidant responses. <i>Scientific Reports</i> , 2021, 11, 8114.	3.3	9
31	The shoot system architecture of Vitis vinifera ssp. sativa. <i>Scientia Horticulturae</i> , 2021, 288, 110404.	3.6	7
32	The Microvine: A Versatile Plant Model to Boost Grapevine Studies in Physiology and Genetics. , 2019, , .		5
33	Effect of the plant sink/source balance on the metabolic content of the Vitis vinifera L. red grape. <i>European Journal of Agronomy</i> , 2021, 122, 126168.	4.1	4