

Etti Or

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

1,095
citations

516710

16
h-index

794594

19
g-index

20
all docs

20
docs citations

20
times ranked

875
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of potential post-ethylene events in the signaling cascade induced by stimuli of bud dormancy release in grapevine. <i>Plant Journal</i> , 2020, 104, 1251-1268.	5.7	11
2	Distinct gibberellin functions during and after grapevine bud dormancy release. <i>Journal of Experimental Botany</i> , 2018, 69, 1635-1648.	4.8	64
3	Transient induction of a subset of ethylene biosynthesis genes is potentially involved in regulation of grapevine bud dormancy release. <i>Plant Molecular Biology</i> , 2018, 98, 507-523.	3.9	18
4	Abscisic acid catabolism enhances dormancy release of grapevine buds. <i>Plant, Cell and Environment</i> , 2018, 41, 2490-2503.	5.7	52
5	ctsGE" clustering subgroups of expression data. <i>Bioinformatics</i> , 2017, 33, 2053-2055.	4.1	2
6	Abnormal Endogenous Repression of GA Signaling in a Seedless Table Grape Cultivar with High Berry Growth Response to GA Application. <i>Frontiers in Plant Science</i> , 2017, 8, 850.	3.6	28
7	Abscisic acid (ABA) regulates grape bud dormancy, and dormancy release stimuli may act through modification of ABA metabolism. <i>Journal of Experimental Botany</i> , 2015, 66, 1527-1542.	4.8	174
8	Functional characterization and developmental expression profiling of gibberellin signalling components in <i>Vitis vinifera</i> . <i>Journal of Experimental Botany</i> , 2015, 66, 1463-1476.	4.8	36
9	Cytokinin-induced WtFL1A expression may be involved in the control of grapevine fruitfulness. <i>Planta</i> , 2012, 235, 181-192.	3.2	19
10	A Method for Isolating Total RNA from Mature Buds and Other Woody Tissues of <i>Vitis Vinifera</i> . , 2010, , 301-307.		5
11	Transcript profiling in <i>Vitis riparia</i> during chilling requirement fulfillment reveals coordination of gene expression patterns with optimized bud break. <i>Functional and Integrative Genomics</i> , 2009, 9, 81-96.	3.5	99
12	Gene-expression profiling of grape bud response to two alternative dormancy-release stimuli expose possible links between impaired mitochondrial activity, hypoxia, ethylene-ABA interplay and cell enlargement. <i>Plant Molecular Biology</i> , 2009, 71, 403-423.	3.9	131
13	On the mechanism of dormancy release in grapevine buds: a comparative study between hydrogen cyanamide and sodium azide. <i>Plant Growth Regulation</i> , 2009, 59, 145-152.	3.4	58
14	Similar mechanisms might be triggered by alternative external stimuli that induce dormancy release in grape buds. <i>Planta</i> , 2008, 228, 79-88.	3.2	99
15	Involvement of calcium signalling in dormancy release of grape buds. <i>Journal of Experimental Botany</i> , 2007, 58, 3249-3262.	4.8	58
16	Digital expression profiling of a grape-bud EST collection leads to new insight into molecular events during grape-bud dormancy release. <i>Plant Science</i> , 2007, 173, 446-457.	3.6	44
17	Dormancy in grape buds: isolation and characterization of catalase cDNA and analysis of its expression following chemical induction of bud dormancy release. <i>Plant Science</i> , 2002, 162, 121-130.	3.6	66
18	The transduction of the signal for grape bud dormancy breaking induced by hydrogen cyanamide may involve the SNF-like protein kinase GDBRPK. <i>Plant Molecular Biology</i> , 2000, 43, 483-494.	3.9	83

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19	Fermentative metabolism in grape berries: isolation and characterization of pyruvate decarboxylase cDNA and analysis of its expression throughout berry development. <i>Plant Science</i> , 2000, 156, 151-158.	3.6	21
20	Isolation of mitochondrial malate dehydrogenase and phosphoenolpyruvate carboxylase cDNA clones from grape berries and analysis of their expression pattern throughout berry development. <i>Journal of Plant Physiology</i> , 2000, 157, 527-534.	3.5	27