Daniel Gonzalez-Ibeas

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9607567/publications.pdf

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20 papers	1,950 citations	687363 13 h-index	794594 19 g-index
20 all docs	20 docs citations	20 times ranked	2592 citing authors

#	Article	IF	Citations
1	Transcriptomics of Fruit Ripening in Citrus. , 2021, , 602-613.		1
2	Virus-Infected Melon Plants Emit Volatiles that Induce Gene Deregulation in Neighboring Healthy Plants. Phytopathology, 2021, 111, 862-869.	2.2	5
3	Shaping the biology of citrus: I. Genomic determinants of evolution. Plant Genome, 2021, 14, e20104.	2.8	4
4	Shaping the biology of citrus: II. Genomic determinants of domestication. Plant Genome, 2021, 14, e20133.	2.8	2
5	Differential expression of IDA (INFLORESCENCE DEFICIENT IN ABSCISSION)-like genes in Nicotiana benthamiana during corolla abscission, stem growth and water stress. BMC Plant Biology, 2020, 20, 34.	3.6	10
6	Comparative Transcriptomics Among Four White Pine Species. G3: Genes, Genomes, Genetics, 2018, 8, 1461-1474.	1.8	35
7	The Douglas-Fir Genome Sequence Reveals Specialization of the Photosynthetic Apparatus in Pinaceae. G3: Genes, Genomes, Genetics, 2017, 7, 3157-3167.	1.8	103
8	Assessing the Gene Content of the Megagenome: Sugar Pine (Pinus lambertiana). G3: Genes, Genomes, Genetics, 2016, 6, 3787-3802.	1.8	51
9	Transcriptomic profile of leaf tissue from the leguminous tree, Millettia pinnata. Tree Genetics and Genomes, 2016, 12, 1.	1.6	11
10	Sequence of the Sugar Pine Megagenome. Genetics, 2016, 204, 1613-1626.	2.9	169
11	The walnut (Juglans regia) genome sequence reveals diversity in genes coding for the biosynthesis of nonâ€structural polyphenols. Plant Journal, 2016, 87, 507-532.	5.7	233
12	Microarray Analysis Shows That Recessive Resistance to Watermelon mosaic virus in Melon Is Associated with the Induction of Defense Response Genes. Molecular Plant-Microbe Interactions, 2012, 25, 107-118.	2.6	25
13	A Cost-effective Double-Stranded cDNA Synthesis for Plant Microarrays. Plant Molecular Biology Reporter, 2012, 30, 1276-1282.	1.8	1
14	The genome of melon (Cucumis melo L.). Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 11872-11877.	7.1	654
15	Analysis of the melon (Cucumis melo) small RNAome by high-throughput pyrosequencing. BMC Genomics, 2011, 12, 393.	2.8	58
16	An oligo-based microarray offers novel transcriptomic approaches for the analysis of pathogen resistance and fruit quality traits in melon (Cucumis melo L.). BMC Genomics, 2009, 10, 467.	2.8	61
17	A set of EST-SNPs for map saturation and cultivar identification in melon. BMC Plant Biology, 2009, 9, 90.	3.6	90
18	Deep-sequencing of plant viral small RNAs reveals effective and widespread targeting of viral genomes. Virology, 2009, 392, 203-214.	2.4	274

#	Article	IF	Citations
19	Mechanism of plant eIF4Eâ€mediated resistance against a Carmovirus (Tombusviridae): capâ€independent translation of a viral RNA controlled inâ€ƒcis by an (a)virulence determinant. Plant Journal, 2008, 56, 716-727.	5.7	76
20	MELOGEN: an EST database for melon functional genomics. BMC Genomics, 2007, 8, 306.	2.8	87