Keithanne Mockaitis

List of Publications by Year in descending order

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Version: 2024-02-01

567281 839539 1,503 18 15 18 citations h-index g-index papers 19 19 19 2365 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Auxin Receptors and Plant Development: A New Signaling Paradigm. Annual Review of Cell and Developmental Biology, 2008, 24, 55-80.	9.4	547
2	Auxin induces mitogenic activated protein kinase (MAPK) activation in roots of Arabidopsis seedlings. Plant Journal, 2000, 24, 785-796.	5.7	185
3	Camelina seed transcriptome: a tool for meal and oil improvement and translational research. Plant Biotechnology Journal, 2013, 11, 759-769.	8.3	166
4	Oil biosynthesis in a basal angiosperm: transcriptome analysis of Persea Americana mesocarp. BMC Plant Biology, 2015, 15, 203.	3.6	96
5	Toward production of jet fuel functionality in oilseeds: identification of FatB acyl-acyl carrier protein thioesterases and evaluation of combinatorial expression strategies in <i>Camelina</i> seeds. Journal of Experimental Botany, 2015, 66, 4251-4265.	4.8	80
6	Population genomic analyses of the chocolate tree, Theobroma cacao L., provide insights into its domestication process. Communications Biology, 2018, $1,167$.	4.4	73
7	Deep sequencing of the Mexican avocado transcriptome, an ancient angiosperm with a high content of fatty acids. BMC Genomics, 2015, 16, 599.	2.8	69
8	Making a chocolate chip: development and evaluation of a 6K SNP array forTheobroma cacao. DNA Research, 2015, 22, 279-291.	3.4	46
9	A Specialized Diacylglycerol Acyltransferase Contributes to the Extreme Medium-Chain Fatty Acid Content of <i>Cuphea</i> Seed Oil. Plant Physiology, 2017, 174, 97-109.	4.8	44
10	Stilbenoid prenyltransferases define key steps in the diversification of peanut phytoalexins. Journal of Biological Chemistry, 2018, 293, 28-46.	3.4	36
11	The Zinc-Finger Protein SOP1 Is Required for a Subset of the Nuclear Exosome Functions in Arabidopsis. PLoS Genetics, 2016, 12, e1005817.	3.5	36
12	A novel Ca2+-binding protein that can rapidly transduce auxin responses during root growth. PLoS Biology, 2019, 17, e3000085.	5.6	35
13	Structurally divergent lysophosphatidic acid acyltransferases with high selectivity for saturated medium chain fatty acids from <i>Cuphea</i> seeds. Plant Journal, 2015, 84, 1021-1033.	5.7	32
14	Integrating transcriptional controls for plant cell expansion. Genome Biology, 2004, 5, 245.	9.6	20
15	A Stilbenoid-Specific Prenyltransferase Utilizes Dimethylallyl Pyrophosphate from the Plastidic Terpenoid Pathway. Plant Physiology, 2016, 171, 2483-2498.	4.8	18
16	A Longâ€Read Transcriptome Assembly of Cotton (Gossypium hirsutum L.) and Intraspecific Single Nucleotide Polymorphism Discovery. Plant Genome, 2015, 8, eplantgenome2014.10.0068.	2.8	12
17	FEATnotator: A tool for integrated annotation of sequence features and variation, facilitating interpretation in genomics experiments. Methods, 2015, 79-80, 11-17.	3.8	7
18	Development and Evaluation of Quality Metrics for Bioinformatics Analysis of Viral Insertion Site Data Generated Using High Throughput Sequencing. Biomedicines, 2014, 2, 195-210.	3.2	1