

Suzana Tiemi Ivamoto-Suzuki

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

195
citations

1163117

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docs citations

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times ranked

255
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study reveals candidate genes influencing lipids and diterpenes contents in <i>Coffea arabica</i> L. <i>Scientific Reports</i> , 2018, 8, 465.	3.3	53
2	Transcriptome Analysis of Leaves, Flowers and Fruits Perisperm of <i>Coffea arabica</i> L. Reveals the Differential Expression of Genes Involved in Raffinose Biosynthesis. <i>PLoS ONE</i> , 2017, 12, e0169595.	2.5	35
3	An integrated analysis of mRNA and sRNA transcriptional profiles in <i>Coffea arabica</i> L. roots: insights on nitrogen starvation responses. <i>Functional and Integrative Genomics</i> , 2019, 19, 151-169.	3.5	28
4	Diterpenes biochemical profile and transcriptional analysis of cytochrome P450s genes in leaves, roots, flowers, and during <i>Coffea arabica</i> L. fruit development. <i>Plant Physiology and Biochemistry</i> , 2017, 111, 340-347.	5.8	19
5	Capsicum-Colletotrichum interaction: Identification of resistance sources and quantification of secondary metabolites in unripe and ripe fruits in response to anthracnose infection. <i>Scientia Horticulturae</i> , 2019, 246, 469-477.	3.6	15
6	Genetic Progress of Seed Yield and Nitrogen Use Efficiency of Brazilian carioca Common Bean Cultivars Using Bayesian Approaches. <i>Frontiers in Plant Science</i> , 2020, 11, 1168.	3.6	13
7	Capsidiol-related genes are highly expressed in response to <i>Colletotrichum scovillei</i> during <i>Capsicum annuum</i> fruit development stages. <i>Scientific Reports</i> , 2020, 10, 12048.	3.3	10
8	Transcriptional patterns of <i>Coffea arabica</i> L. nitrate reductase, glutamine and asparagine synthetase genes are modulated under nitrogen suppression and coffee leaf rust. <i>PeerJ</i> , 2020, 8, e8320.	2.0	8
9	Low-Copy Genes in Terpenoid Metabolism: The Evolution and Expression of MVK and DXR Genes in Angiosperms. <i>Plants</i> , 2020, 9, 525.	3.5	6
10	Identification of the transcriptionally active cytochrome P450 repertoire in <i>Coffea arabica</i> . <i>Genetics and Molecular Research</i> , 2015, 14, 2399-2412.	0.2	4
11	<i>Coffea arabica</i> L. genes from isoprenoid metabolic pathways are more expressed in full sun cultivation systems than in agroforestry systems. <i>Plant Gene</i> , 2021, 26, 100287.	2.3	2
12	Identificação e Caracterização de Microsatélites de <i>Coffea arabica</i> a partir de dados de sequenciamento de RNA e de BACs. <i>BBR - Biochemistry and Biotechnology Reports</i> , 2013, 2, 186.	0.0	1
13	Adaptability and stability analysis of new popcorn simple hybrids evaluated using additive main effects and multiplicative interaction Bayesian approaches. <i>Bragantia</i> , 0, 81, .	1.3	1