Suzana Tiemi Ivamoto-Suzuki

List of Publications by Year
in descending order

13	195	1163117 8	1199594 12
papers	citations	h-index	g-index

13	13	13	255
all docs	docs citations	times ranked	citing authors

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