Tiago Benedito Dos Santos

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Physiological Responses to Drought, Salinity, and Heat Stress in Plants: A Review. Stresses, 2022, 2, 113-135.	4.8	149
2	Small heat shock protein (Hsp20) gene family in Phaseolus vulgaris L.: Genome-wide identification, evolutionary and expression analysis. Plant Gene, 2022, 31, 100370.	2.3	2
3	Genome-Wide Identification, Evolution, and Expression Profile of Aquaporin Genes in Coffea canephora in Response to Water Deficit. Plant Molecular Biology Reporter, 2021, 39, 146-162.	1.8	4
4	The urea transporter DUR3 is differentially regulated by abiotic and biotic stresses in coffee plants. Physiology and Molecular Biology of Plants, 2021, 27, 203-212.	3.1	3
5	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
6	Urochloa brizantha cv. Marandu presents a better response to in vitro salt stress than other commercial cultivars. Colloquium Agrariae, 2021, 17, 74-82.	0.2	1
7	Validation of reference genes for real-time quantitative PCR in Brachiaria grass under salt stress. Plant Gene, 2021, 27, 100319.	2.3	3
8	An in silicodata miningof theammonium transporter gene familyin Ananas comosusL Colloquium Agrariae, 2021, 16, 10-24.	0.2	1
9	Involvement of the galactinol synthase gene in abiotic and biotic stress responses: A review on current knowledge. Plant Gene, 2020, 24, 100258.	2.3	10
10	Identification, evolutionary and expression analysis of the galactinol synthase (GolS) genes in Panicum virgatum L. and Panicum hallii : An in silico approach. Plant Gene, 2020, 24, 100262.	2.3	0
11	Genome-wide in silico analysis of SOD genes in common bean (Phaseolus vulgaris L.). Journal of Crop Science and Biotechnology, 2020, 23, 241-251.	1.5	3
12	Transcriptional patterns of <i>Coffea arabica</i> L. nitrate reductase, glutamine and asparagine synthetase genes are modulated under nitrogen suppression and coffee leaf rust. PeerJ, 2020, 8, e8320.	2.0	8
13	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
14	Regulation of α-expansins genes in Arabidopsis thaliana seeds during post-osmopriming germination. Physiology and Molecular Biology of Plants, 2019, 25, 511-522.	3.1	10
15	A GENOME-WIDE ANALYSIS OF THE GALACTINOL SYNTHASEGENE FAMILY IN BANANA (Musa acuminata). Colloquium Agrariae, 2018, 14, 01-11.	0.2	3
16	IN SILICO ANALYSIS OF THE Dof TRANSCRIPTION FACTOR FAMILY IN Coffea canephora. Colloquium Agrariae, 2018, 14, 99-111.	0.2	1
17	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
18	Genome-wide identification, classification and transcriptional analysis of nitrate and ammonium transporters in Coffea. Genetics and Molecular Biology, 2017, 40, 346-359.	1.3	10

#	Article	IF	CITATIONS
19	Changes in growth, oxidative metabolism and essential oil composition of lemon balm (Melissa) Tj ETQq1 1 0.784	4314 rgBT	/Qyerlock 1(
20	Galactinol synthase transcriptional profile in two genotypes of Coffea canephora with contrasting tolerance to drought. Genetics and Molecular Biology, 2015, 38, 182-190.	1.3	40
21	Salt stress alters the cell wall polysaccharides and anatomy of coffee (Coffea arabica L.) leaf cells. Carbohydrate Polymers, 2014, 112, 686-694.	10.2	46
22	Heat stress causes alterations in the cell-wall polymers and anatomy of coffee leaves (Coffea arabica) Tj ETQq0 0	0 rgBT /O 10.2	verlock 10 Tf
23	Nitrogen Starvation, Salt and Heat Stress in Coffee (Coffea arabica L.): Identification and Validation of New Genes for qPCR Normalization. Molecular Biotechnology, 2013, 53, 315-325.	2.4	42
24	FISH using a gag-like fragment probe reveals a common Ty3-gypsy-like retrotransposon in genome of Coffea species. Genome, 2012, 55, 825-833.	2.0	9
25	Gene expression and enzymatic activity of pectin methylesterase during fruit development and ripening in Coffea arabica L Genetics and Molecular Research, 2012, 11, 3186-3197.	0.2	9
26	Expression patterns of three αâ€expansin isoforms in <i>Coffea arabica</i> during fruit development. Plant Biology, 2011, 13, 462-471.	3.8	20
27	Expression of three galactinol synthase isoforms in Coffea arabica L. and accumulation of raffinose and stachyose in response to abiotic stresses. Plant Physiology and Biochemistry, 2011, 49, 441-448.	5.8	108

28	Molybdenum (Mo) transporter genes in Panicoideae species: a genome-wide evolution study. Journal of Crop Science and Biotechnology, 0, , 1.	1.5	0
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