## Michael S Brito

List of Publications by Year in descending order

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686830 525886 26 847 13 27 citations h-index g-index papers 27 27 27 1229 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Selection and validation of reference genes by RT-qPCR under photoperiodic induction of flowering in sugarcane (Saccharum spp.). Scientific Reports, 2021, 11, 4589.	1.6	7
2	The sugarcane ShMYB78 transcription factor activates suberin biosynthesis in Nicotiana benthamiana. Plant Molecular Biology, 2020, 104, 411-427.	2.0	15
3	Sugarcane mosaic virus mediated changes in cytosine methylation pattern and differentially transcribed fragments in resistance-contrasting sugarcane genotypes. PLoS ONE, 2020, 15, e0241493.	1.1	8
4	Reference genes for gene expression studies targeting sugarcane infected with Sugarcane mosaic virus (SCMV). BMC Research Notes, 2019, 12, 149.	0.6	8
5	Deposition of lignin in four species of Saccharum. Scientific Reports, 2019, 9, 5877.	1.6	41
6	Biomass Accumulation and Cell Wall Structure of Rice Plants Overexpressing a Dirigent-Jacalin of Sugarcane (ShDJ) Under Varying Conditions of Water Availability. Frontiers in Plant Science, 2019, 10, 65.	1.7	12
7	Overexpression of ScMYBAS1 alternative splicing transcripts differentially impacts biomass accumulation and drought tolerance in rice transgenic plants. PLoS ONE, 2018, 13, e0207534.	1.1	21
8	Ectopic expression of sugarcane SHINE changes cell wall and improves biomass in rice. Biomass and Bioenergy, 2018, 119, 322-334.	2.9	11
9	A novel cysteine-rich peptide regulates cell expansion in the tobacco pistil and influences its final size. Plant Science, 2018, 277, 55-67.	1.7	3
10	Identification, classification and transcriptional profiles of dirigent domain-containing proteins in sugarcane. Molecular Genetics and Genomics, 2017, 292, 1323-1340.	1.0	14
11	Reference genes for normalization of qPCR assays in sugarcane plants under water deficit. Plant Methods, 2017, 13, 28.	1.9	40
12	Building the wall: recent advances in understanding lignin metabolism in grasses. Acta Physiologiae Plantarum, 2016, 38, 1.	1.0	29
13	Characterization of PIP2 aquaporins in Saccharum hybrids. Plant Gene, 2016, 5, 31-37.	1.4	10
14	Pollination triggers female gametophyte development in immature Nicotiana tabacum flowers. Frontiers in Plant Science, 2015, 6, 561.	1.7	13
15	Expression Profile of Sugarcane Transcription Factor Genes Involved in Lignin Biosynthesis. Tropical Plant Biology, 2015, 8, 19-30.	1.0	13
16	Water Stress Alters Lignin Content and Related Gene Expression in Two Sugarcane Genotypes. Journal of Agricultural and Food Chemistry, 2015, 63, 4708-4720.	2.4	42
17	Large-Scale Transcriptome Analysis of Two Sugarcane Genotypes Contrasting for Lignin Content. PLoS ONE, 2015, 10, e0134909.	1.1	51
18	Protocol for the Generation of a Transcription Factor Open Reading Frame Collection (TFome). Bio-protocol, 2015, 5, .	0.2	3

#	Article	IF	CITATIONS
19	Characterization, isolation and cloning of sugarcane genes related to drought stress. BMC Proceedings, $2014,8,.$	1.8	6
20	In situ localization of mRNA of resembling the dirigent protein in sugarcane stems. BMC Proceedings, 2014, $8$ , .	1.8	1
21	The Maize <scp>TF</scp> ome – development of a transcription factor open reading frame collection for functional genomics. Plant Journal, 2014, 80, 356-366.	2.8	55
22	Validation of reference genes from Eucalyptus spp. under different stress conditions. BMC Research Notes, 2012, 5, 634.	0.6	26
23	Stigma/style cell cycle inhibitor 1 (SCI1), a tissueâ€specific cell cycle regulator that controls upper pistil development. New Phytologist, 2011, 190, 882-895.	3.5	17
24	Analysis of the <i>Nicotiana tabacum</i> Stigma/Style Transcriptome Reveals Gene Expression Differences between Wet and Dry Stigma Species  Â. Plant Physiology, 2009, 149, 1211-1230.	2.3	65
25	Comparative Analyses of the Complete Genome Sequences of Pierce's Disease and Citrus Variegated Chlorosis Strains of Xylella fastidiosa. Journal of Bacteriology, 2003, 185, 1018-1026.	1.0	307
26	Dissecting the sugarcane expressed sequence tag (SUCEST) database: unraveling flower-specific genes. Genetics and Molecular Biology, 2001, 24, 77-84.	0.6	6