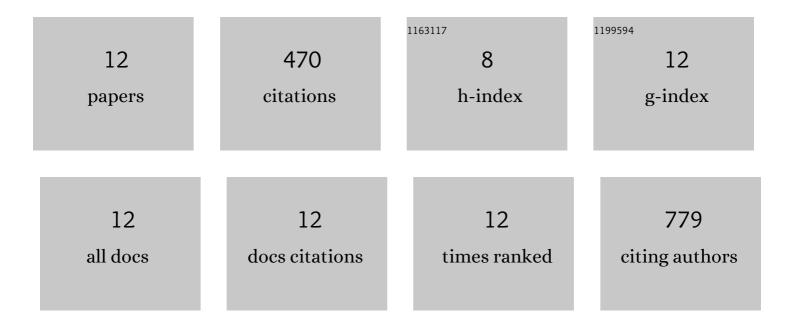
Sarah Muniz Nardeli

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identification and evaluation of new reference genes in Gossypium hirsutumfor accurate normalization of real-time quantitative RT-PCR data. BMC Plant Biology, 2010, 10, 49.	3.6	283
2	Analysis of the arabidopsis REM gene family predicts functions during flower development. Annals of Botany, 2014, 114, 1507-1515.	2.9	55
3	Genome-wide analysis of the MADS-box gene family in polyploid cotton (Gossypium hirsutum) and in its diploid parental species (Gossypium arboreum and Gossypium raimondii). Plant Physiology and Biochemistry, 2018, 127, 169-184.	5.8	30
4	lsolation and functional characterization of a cotton ubiquitination-related promoter and 5'UTR that drives high levels of expression in root and flower tissues. BMC Biotechnology, 2011, 11, 115.	3.3	18
5	Insights into the role of alternative splicing in plant temperature response. Journal of Experimental Botany, 2021, , .	4.8	17
6	Transcriptional responses of Arabidopsis thaliana to oil contamination. Environmental and Experimental Botany, 2016, 127, 63-72.	4.2	13
7	Isolation and Characterization of Three New Promoters from Gossypium hirsutum that Show High Activity in Reproductive Tissues. Plant Molecular Biology Reporter, 2014, 32, 630-643.	1.8	12
8	Functional characterization of AGAMOUS-subfamily members from cotton during reproductive development and in response to plant hormones. Plant Reproduction, 2017, 30, 19-39.	2.2	12
9	Overexpression of the CaHB12 transcription factor in cotton (Gossypium hirsutum) improves drought tolerance. Plant Physiology and Biochemistry, 2021, 165, 80-93.	5.8	11
10	Early Heat Shock Protein Response and Selection of Reference Genes in Arabidopsis thaliana Seedlings Subjected to Marine Fuel Contamination. Water, Air, and Soil Pollution, 2017, 228, 1.	2.4	9
11	Evaluation of Reference Genes for RTâ€qPCR Normalization in Cowpea under Drought Stress during Biological Nitrogen Fixation. Crop Science, 2015, 55, 1660-1672.	1.8	6
12	Promoter isolation and characterization of GhAO-like1, a Gossypium hirsutum gene similar to multicopper oxidases that is highly expressed in reproductive organs. Genome, 2016, 59, 23-36.	2.0	4