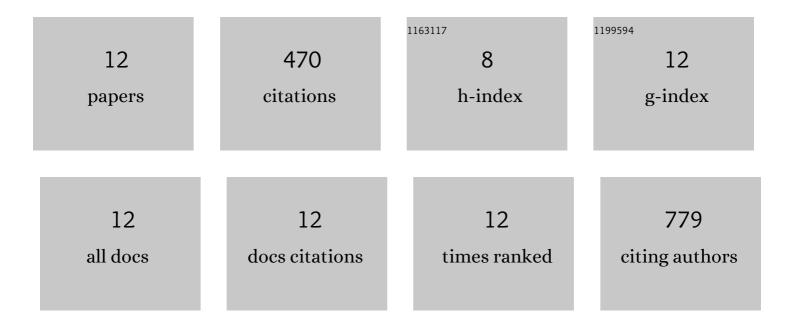
## Sarah Muniz Nardeli

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1848973/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Insights into the role of alternative splicing in plant temperature response. Journal of Experimental Botany, 2021, , .	4.8	17
2	Overexpression of the CaHB12 transcription factor in cotton (Gossypium hirsutum) improves drought tolerance. Plant Physiology and Biochemistry, 2021, 165, 80-93.	5.8	11
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	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
5	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
6	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
7	Transcriptional responses of Arabidopsis thaliana to oil contamination. Environmental and Experimental Botany, 2016, 127, 63-72.	4.2	13
8	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
9	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
10	Analysis of the arabidopsis REM gene family predicts functions during flower development. Annals of Botany, 2014, 114, 1507-1515.	2.9	55
11	Isolation 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
12	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