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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2796042/publications.pdf>

Version: 2024-02-01

12
papers

247
citations

1478505

6
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

427
citing authors

#	ARTICLE	IF	CITATIONS
1	RNA-Seq analysis of the wild barley (<i>H. spontaneum</i>) leaf transcriptome under salt stress. <i>Comptes Rendus - Biologies</i> , 2015, 338, 285-297.	0.2	76
2	Whole Mitochondrial and Plastid Genome SNP Analysis of Nine Date Palm Cultivars Reveals Plastid Heteroplasmy and Close Phylogenetic Relationships among Cultivars. <i>PLoS ONE</i> , 2014, 9, e94158.	2.5	58
3	Metabolomic Response of <i>Calotropis procera</i> Growing in the Desert to Changes in Water Availability. <i>PLoS ONE</i> , 2014, 9, e87895.	2.5	25
4	Characterization of ten date palm (<i>Phoenix dactylifera</i> L.) cultivars from Saudi Arabia using AFLP and ISSR markers. <i>Comptes Rendus - Biologies</i> , 2014, 337, 6-18.	0.2	25
5	Multifunctional activities of ERF109 as affected by salt stress in <i>Arabidopsis</i> . <i>Scientific Reports</i> , 2018, 8, 6403.	3.3	24
6	Control of glycerol biosynthesis under high salt stress in <i>Arabidopsis</i> . <i>Functional Plant Biology</i> , 2014, 41, 87.	2.1	13
7	Corrected sequence of the wheat plastid genome. <i>Comptes Rendus - Biologies</i> , 2014, 337, 499-502.	0.2	6
8	Structural identification of putative USPs in <i>Catharanthus roseus</i> . <i>Comptes Rendus - Biologies</i> , 2015, 338, 643-649.	0.2	6
9	Suppression of PCD-related genes affects salt tolerance in <i>Arabidopsis</i> . <i>Comptes Rendus - Biologies</i> , 2016, 339, 105-114.	0.2	6
10	Stepwise response of MeJA-induced genes and pathways in leaves of <i>C. roseus</i> . <i>Comptes Rendus - Biologies</i> , 2018, 341, 411-420.	0.2	4
11	Transcription factors regulating <i>uspA</i> genes in <i>Catharanthus roseus</i> . <i>Comptes Rendus - Biologies</i> , 2017, 340, 1-6.	0.2	3
12	Detection of phytochrome-like genes from <i>Rhazya stricta</i> (Apocynaceae) using de novo genome assembly. <i>Comptes Rendus - Biologies</i> , 2013, 336, 521-529.	0.2	1