## Danni Yang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7557040/publications.pdf

Version: 2024-02-01

10	115	7	10
papers	citations	h-index	g-index
10	10	10	106
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Cold Response Transcriptome Analysis of the Alternative Splicing Events Induced by the Cold Stress in D. catenatum. International Journal of Molecular Sciences, 2022, 23, 981.	4.1	8
2	Genome-Wide Identification of OSC Gene Family and Potential Function in the Synthesis of Ursane- and Oleanane-Type Triterpene in Momordica charantia. International Journal of Molecular Sciences, 2022, 23, 196.	4.1	4
3	Functional Characterization of the Stipa purpureaÂP5CS Gene under Drought Stress Conditions. International Journal of Molecular Sciences, 2021, 22, 9599.	4.1	23
4	Genome-Wide Analysis of the TCP Transcription Factor Genes in Dendrobium catenatum Lindl International Journal of Molecular Sciences, 2021, 22, 10269.	4.1	14
5	Comparative transcriptome analysis reveals ecological adaption of cold tolerance in northward invasion of Alternanthera philoxeroides. BMC Genomics, 2020, 21, 532.	2.8	10
6	BrrlCE1.1 is associated with putrescine synthesis through regulation of the arginine decarboxylase gene in freezing tolerance of turnip (Brassica rapa var. rapa). BMC Plant Biology, 2020, 20, 504.	3.6	13
7	Uncovering the role of a positive selection site of wax ester synthase/diacylglycerol acyltransferase in two closely related Stipa species in wax ester synthesis under drought stress. Journal of Experimental Botany, 2020, 71, 4159-4170.	4.8	7
8	A Splice Variant of <i>BrrWSD1</i> in Turnip ( <i>Brassica rapa</i> var. rapa) and Its Possible Role in Wax Ester Synthesis under Drought Stress. Journal of Agricultural and Food Chemistry, 2019, 67, 11077-11088.	5.2	9
9	Genome-wide analysis indicates diverse physiological roles of the turnip (Brassica rapa var. rapa) oligopeptide transporters gene family. Plant Diversity, 2018, 40, 57-67.	3.7	9
10	Molecular cloning and characterization of the glutathione reductase gene from Stipa purpurea. Biochemical and Biophysical Research Communications, 2018, 495, 1851-1857.	2.1	18