Xinlong Xiao

List of Publications by Year in descending order

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		1478505	1281871	
10	241	6	11	
papers	citations	h-index	g-index	
11	11	11	408	
11	11	11	400	
all docs	docs citations	times ranked	citing authors	
11 all docs	11 docs citations	11 times ranked	408 citing authors	

#	Article	IF	CITATIONS
1	An Agrobacterium-mediated transient expression method contributes to functional analysis of a transcription factor and potential application of gene editing in Chenopodium quinoa. Plant Cell Reports, 2022, 41, 1975-1985.	5.6	6
2	Mechanism of phosphate sensing and signaling revealed by rice SPX1-PHR2 complex structure. Nature Communications, 2021, 12, 7040.	12.8	37
3	Regulation of Ammonium Cellular Levels is An Important Adaptive Trait for the Euhalophytic Behavior of Salicornia europaea. Plants, 2020, 9, 257.	3.5	9
4	A group of SUVH methylâ€DNA binding proteins regulate expression of the DNA demethylase ROS1 in <i>Arabidopsis</i> . Journal of Integrative Plant Biology, 2019, 61, 110-119.	8.5	44
5	Large-scale de novo transcriptome analysis reveals specific gene expression and novel simple sequence repeats markers in salinized roots of the euhalophyte Salicornia europaea. Acta Physiologiae Plantarum, 2018, 40, 1.	2.1	2
6	A method for the further assembly of targeted unigenes in a transcriptome after assembly by Trinity. Frontiers in Plant Science, 2015, 6, 843.	3.6	6
7	Systematic assessment of reference genes for RT-qPCR across plant species under salt stress and drought stress. Acta Physiologiae Plantarum, 2015, 37, 1.	2.1	6
8	Root transcriptome analysis on the grape genotypes with contrast translocation pattern of excess manganese from root to shoot. Plant and Soil, 2015, 387, 49-67.	3.7	11
9	Validation of suitable reference genes for gene expression analysis in the halophyte Salicornia europaea by real-time quantitative PCR. Frontiers in Plant Science, 2014, 5, 788.	3.6	81
10	Global Transcriptome Profiling of Salicornia europaea L. Shoots under NaCl Treatment. PLoS ONE, 2013, 8, e65877.	2.5	38