Zegang Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptome-wide high-throughput deep m6A-seq reveals unique differential m6A methylation patterns between three organs in Arabidopsis thaliana. Genome Biology, 2015, 16, 272.	8.8	145
2	High-throughput m6A-seq reveals RNA m6A methylation patterns in the chloroplast and mitochondria transcriptomes of Arabidopsis thaliana. PLoS ONE, 2017, 12, e0185612.	2.5	43
3	Iron and callose homeostatic regulation in rice roots under low phosphorus. BMC Plant Biology, 2018, 18, 326.	3.6	10
4	Mechanism of Low Phosphorus Inducing the Main Root Lengthening of Rice. Journal of Plant Growth Regulation, 2021, 40, 1032-1043.	5.1	10
5	Expression of sulfur uptake assimilation-related genes in response to cadmium, bensulfuron-methyl and their co-contamination in rice roots. Journal of Environmental Sciences, 2014, 26, 650-661.	6.1	4
6	Regulation of OsLPR1 gene on the formation of rice root iron plaque under low phosphorus. Plant Biotechnology Reports, 0, , 1.	1.5	0