Fuhua Fan

List of Publications by Year in descending order

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ΕΠΗΠΑ ΕΛΝ

#	Article	IF	CITATIONS
1	Transcriptome-Wide Identification and Expression Profiles of Masson Pine WRKY Transcription Factors in Response to Low Phosphorus Stress. Plant Molecular Biology Reporter, 2021, 39, 1-9.	1.8	12
2	Genetic diversity and population structure of masson pine (Pinus massoniana Lamb.) superior clones in South China as revealed by EST-SSR markers. Genetic Resources and Crop Evolution, 2021, 68, 1987-2002.	1.6	8
3	Exogenous Brassinosteroid Facilitates Xylem Development in Pinus massoniana Seedlings. International Journal of Molecular Sciences, 2021, 22, 7615.	4.1	14
4	Transcriptome-Wide Identification and Expression Profiling of SPX Domain-Containing Members in Responses to Phosphorus Deprivation of Pinus massoniana. Forests, 2021, 12, 1627.	2.1	2
5	Transcriptome-wide identification and expression profiling of Pinus massoniana MYB transcription factors responding to phosphorus deficiency. Journal of Forestry Research, 2020, 31, 909-919.	3.6	15
6	Proteomic analyses provide new insights into the responses of <i>Pinus massoniana</i> seedlings to phosphorus deficiency. Proteomics, 2016, 16, 504-515.	2.2	22
7	The Temporal Transcriptomic Response of Pinus massoniana Seedlings to Phosphorus Deficiency. PLoS ONE, 2014, 9, e105068.	2.5	32
8	LTR-retrotransposon activation, IRAP marker development and its potential in genetic diversity assessment of masson pine (Pinus massoniana). Tree Genetics and Genomes, 2014, 10, 213-222.	1.6	27
9	Isolation, identification, and characterization of genomic LTR retrotransposon sequences from masson pine (Pinus massoniana). Tree Genetics and Genomes, 2013, 9, 1237-1246.	1.6	9
10	Integrated mRNA and miRNA Expression Analyses of Pinus massoniana Roots and Shoots in Long-Term Response to Phosphate Deficiency. Journal of Plant Growth Regulation, 0, , 1.	5.1	5