

Xuejing Wen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6280473/publications.pdf>

Version: 2024-02-01

8
papers

152
citations

1478505

6
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1588992

8
g-index

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all docs

8
docs citations

8
times ranked

141
citing authors

#	ARTICLE	IF	CITATIONS
1	An Improved Syringe Agroinfiltration Protocol to Enhance Transformation Efficiency by Combinative Use of 5-Azacytidine, Ascorbate Acid and Tween-20. <i>Plants</i> , 2017, 6, 9.	3.5	58
2	<i>Betula platyphylla</i> BpHOX2 transcription factor binds to different cis-acting elements and confers osmotic tolerance. <i>Journal of Integrative Plant Biology</i> , 2020, 62, 1762-1779.	8.5	23
3	Identification and characterization of cadmium stress-related LncRNAs from <i>Betula platyphylla</i> . <i>Plant Science</i> , 2020, 299, 110601.	3.6	19
4	PacBio full-length transcriptome of wild apple (<i>Malus sieversii</i>) provides insights into canker disease dynamic response. <i>BMC Genomics</i> , 2021, 22, 52.	2.8	19
5	Transcriptome profiling of <i>Malus sieversii</i> under freezing stress after being cold-acclimated. <i>BMC Genomics</i> , 2021, 22, 681.	2.8	18
6	Reverse Chromatin Immunoprecipitation (R-ChIP) enables investigation of the upstream regulators of plant genes. <i>Communications Biology</i> , 2020, 3, 770.	4.4	10
7	Full-Length Transcriptome-Wide Characteristic and Functional Identification of WRKY Family in <i>Malus sieversii</i> during the Valsa Canker Disease Response. <i>Forests</i> , 2021, 12, 790.	2.1	3
8	Genome-Wide Characterization of HSP90 Gene Family in <i>Malus sieversii</i> and Their Potential Roles in Response to <i>Valsa mali</i> Infection. <i>Forests</i> , 2021, 12, 1232.	2.1	2