

Chuanen Zhou

List of Publications by Year in descending order

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36
papers

1,171
citations

516710

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33
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docs citations

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#	ARTICLE	IF	CITATIONS
1	Overexpression of miR156 in switchgrass (<i>Panicum virgatum</i> L.) results in various morphological alterations and leads to improved biomass production. <i>Plant Biotechnology Journal</i> , 2012, 10, 443-452.	8.3	293
2	From Model to Crop: Functional Analysis of a <i>STAY-GREEN</i> Gene in the Model Legume <i>Medicago truncatula</i> and Effective Use of the Gene for Alfalfa Improvement. <i>Plant Physiology</i> , 2011, 157, 1483-1496.	4.8	124
3	Rhizobial Infection Is Associated with the Development of Peripheral Vasculature in Nodules of <i>Medicago truncatula</i> . <i>Plant Physiology</i> , 2013, 162, 107-115.	4.8	92
4	Developmental Analysis of a <i>Medicago truncatula</i> <i>smooth leaf margin1</i> Mutant Reveals Context-Dependent Effects on Compound Leaf Development. <i>Plant Cell</i> , 2011, 23, 2106-2124.	6.6	82
5	The <i>Trans</i> -Acting Short Interfering RNA3 Pathway and NO APICAL MERISTEM Antagonistically Regulate Leaf Margin Development and Lateral Organ Separation, as Revealed by Analysis of an <i>argonaute7</i> / <i>lobed leaflet1</i> Mutant in <i>Medicago truncatula</i> . <i>Plant Cell</i> , 2014, 25, 4845-4862.	6.6	64
6	A class II KNOX gene, <i>KNOX4</i> , controls seed physical dormancy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6997-7002.	7.1	55
7	STM/BP-Like KNOXI Is Uncoupled from ARP in the Regulation of Compound Leaf Development in <i>Medicago truncatula</i> . <i>Plant Cell</i> , 2014, 26, 1464-1479.	6.6	41
8	Identification and characterization of <i>petiolule</i> -like <i>pulvinus</i> mutants with abolished nyctinastic leaf movement in the model legume <i>Medicago truncatula</i> . <i>New Phytologist</i> , 2012, 196, 92-100.	7.3	38
9	From genes to networks: The genetic control of leaf development. <i>Journal of Integrative Plant Biology</i> , 2021, 63, 1181-1196.	8.5	36
10	Genome-Wide Identification of TCP Family Transcription Factors in <i>Medicago truncatula</i> Reveals Significant Roles of miR319-Targeted TCPs in Nodule Development. <i>Frontiers in Plant Science</i> , 2018, 9, 774.	3.6	29
11	Alfalfa (<i>Medicago sativa</i> L.). <i>Methods in Molecular Biology</i> , 2015, 1223, 213-221.	0.9	27
12	The nodulation and nyctinastic leaf movement is orchestrated by clock gene <i>LHY</i> in <i>Medicago truncatula</i> . <i>Journal of Integrative Plant Biology</i> , 2020, 62, 1880-1895.	8.5	26
13	Efficient Generation of CRISPR/Cas9-Mediated Homozygous/Biallelic <i>Medicago truncatula</i> Mutants Using a Hairy Root System. <i>Frontiers in Plant Science</i> , 2020, 11, 294.	3.6	25
14	Genome-wide characterization of SPL family in <i>Medicago truncatula</i> reveals the novel roles of miR156/SPL module in spiky pod development. <i>BMC Genomics</i> , 2019, 20, 552.	2.8	21
15	Transforming compound leaf patterning by manipulating <i>REVOLUTA</i> in <i>Medicago truncatula</i> . <i>Plant Journal</i> , 2019, 100, 562-571.	5.7	20
16	<i>AGLF</i> provides C-function in floral organ identity through transcriptional regulation of <i>AGAMOUS</i> in <i>Medicago truncatula</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 5176-5181.	7.1	20
17	<i>HEADLESS</i> Regulates Auxin Response and Compound Leaf Morphogenesis in <i>Medicago truncatula</i> . <i>Frontiers in Plant Science</i> , 2019, 10, 1024.	3.6	19
18	Ginsenoside Rb1 in asymmetric somatic hybrid calli of <i>Daucus carota</i> with <i>Panax quinquefolius</i> . <i>Plant Cell Reports</i> , 2009, 28, 627-638.	5.6	17

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19	Genetic characterization of asymmetric somatic hybrids between <i>Bupleurum scorzonerifolium</i> Willd and <i>Triticum aestivum</i> L.: potential application to the study of the wheat genome. <i>Planta</i> , 2006, 223, 714-724.	3.2	15
20	Analysis of remote asymmetric somatic hybrids between common wheat and <i>Arabidopsis thaliana</i> . <i>Plant Cell Reports</i> , 2007, 26, 1233-1241.	5.6	14
21	Functional characterization of PETIOLULE-LIKE PULVINUS (PLP) gene in abscission zone development in <i>Medicago truncatula</i> and its application to genetic improvement of alfalfa. <i>Plant Biotechnology Journal</i> , 2021, 19, 351-364.	8.3	13
22	LATE MERISTEM IDENTITY1 regulates leaf margin development via the auxin transporter gene <i>SMOOTH LEAF MARGIN1</i> . <i>Plant Physiology</i> , 2021, 187, 218-235.	4.8	13
23	Integrated regulation of periclinal cell division by transcriptional module of BZR1-SHR in <i>Arabidopsis</i> roots. <i>New Phytologist</i> , 2022, 233, 795-808.	7.3	13
24	Regeneration of asymmetric somatic hybrid plants from the fusion of two types of wheat with Russian wildrye. <i>Plant Cell Reports</i> , 2004, 23, 461-467.	5.6	10
25	Systematic Analysis of Gibberellin Pathway Components in <i>Medicago truncatula</i> Reveals the Potential Application of Gibberellin in Biomass Improvement. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7180.	4.1	10
26	Potential but limited redundant roles of MtPIN4, MtPIN5 and MtPIN10/SLM1 in the development of <i>Medicago truncatula</i> . <i>Plant Signaling and Behavior</i> , 2011, 6, 1834-1836.	2.4	8
27	Interaction between the MtDELLA-MtGAF1 Complex and MtARF3 Mediates Transcriptional Control of MtGA3ox1 to Elaborate Leaf Margin Formation in <i>Medicago truncatula</i> . <i>Plant and Cell Physiology</i> , 2021, 62, 321-333.	3.1	8
28	Brassinosteroid homeostasis is critical for the functionality of the <i>Medicago truncatula</i> pulvinus. <i>Plant Physiology</i> , 2021, 185, 1745-1763.	4.8	8
29	MtBZR1 Plays an Important Role in Nodule Development in <i>Medicago truncatula</i> . <i>International Journal of Molecular Sciences</i> , 2019, 20, 2941.	4.1	7
30	Construction of Whole Genome Radiation Hybrid Panels and Map of Chromosome 5A of Wheat Using Asymmetric Somatic Hybridization. <i>PLoS ONE</i> , 2012, 7, e40214.	2.5	5
31	The Conserved and Specific Roles of the LUX ARRHYTHMO in Circadian Clock and Nodulation. <i>International Journal of Molecular Sciences</i> , 2022, 23, 3473.	4.1	5
32	<i>AGAMOUS-LIKE FLOWER</i> regulates flower and compound leaf development through different regulatory mechanisms in <i>Medicago truncatula</i> . <i>Plant Signaling and Behavior</i> , 2019, 14, 1612683.	2.4	4
33	Developmental Analysis of the GATA Factor HANABA TARANU Mutants in <i>Medicago truncatula</i> Reveals Their Roles in Nodule Formation. <i>Frontiers in Plant Science</i> , 2021, 12, 616776.	3.6	4
34	MtPIN1 and MtPIN3 Play Dual Roles in Regulation of Shade Avoidance Response under Different Environments in <i>Medicago truncatula</i> . <i>International Journal of Molecular Sciences</i> , 2020, 21, 8742.	4.1	3
35	Phospho-Mutant Activity Assays Provide Evidence for the Negative Regulation of Transcriptional Regulator PRE1 by Phosphorylation. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9183.	4.1	1
36	Developmental Analysis of Compound Leaf Development in <i>Arachis hypogaea</i> . <i>Frontiers in Plant Science</i> , 2022, 13, 749809.	3.6	1