Jun Xiao

List of Publications by Year in descending order

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471509 794594 2,208 21 17 19 citations h-index g-index papers 24 24 24 2959 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	H2A.Z contributes to trithorax activity at the AGAMOUS locus. Molecular Plant, 2022, 15, 207-210.	8.3	2
2	Genome-edited powdery mildew resistance in wheat without growth penalties. Nature, 2022, 602, 455-460.	27.8	181
3	Chromatin remodeling complexes regulate genome architecture in Arabidopsis. Plant Cell, 2022, 34, 2638-2651.	6.6	24
4	LEAFY is a pioneer transcription factor and licenses cell reprogramming to floral fate. Nature Communications, 2021, 12, 626.	12.8	68
5	Dissecting the plant chromatin interactome using mass spectrometry. Trends in Biotechnology, 2021, ,	9.3	O
6	PRC2 activity, recruitment, and silencing: a comparative perspective. Trends in Plant Science, 2021, 26, 1186-1198.	8.8	42
7	The vernalization-induced long non-coding RNA VAS functions with the transcription factor TaRF2b to promote TaVRN1 expression for flowering in hexaploid wheat. Molecular Plant, 2021, 14, 1525-1538.	8.3	42
8	Integration of Transcriptional Repression and Polycomb-Mediated Silencing of <i>WUSCHEL</i> in Floral Meristems. Plant Cell, 2019, 31, 1488-1505.	6.6	77
9	The Protein Modifications of <i>O</i> -GlcNAcylation and Phosphorylation Mediate Vernalization Response for Flowering in Winter Wheat. Plant Physiology, 2019, 180, 1436-1449.	4.8	34
10	<i>Arabidopsis O</i> â€Glc <scp>NA</scp> c transferase <scp>SEC</scp> activates histone methyltransferase <scp>ATX</scp> 1 to regulate flowering. EMBO Journal, 2018, 37, .	7.8	47
11	Cis and trans determinants of epigenetic silencing by Polycomb repressive complex 2 in Arabidopsis. Nature Genetics, 2017, 49, 1546-1552.	21.4	226
12	Systematic discovery of novel eukaryotic transcriptional regulators using sequence homology independent prediction. BMC Genomics, 2017, 18, 480.	2.8	12
13	Developmental transitions: integrating environmental cues with hormonal signaling in the chromatin landscape in plants. Genome Biology, 2017, 18, 88.	8.8	47
14	Tug of war: adding and removing histone lysine methylation in Arabidopsis. Current Opinion in Plant Biology, 2016, 34, 41-53.	7.1	121
15	AtJAC1 Regulates Nuclear Accumulation of GRP7, Influencing RNA Processing of FLC Antisense Transcripts and Flowering Time in Arabidopsis. Plant Physiology, 2015, 169, pp.00801.2015.	4.8	29
16	Auxin-regulated chromatin switch directs acquisition of flower primordium founder fate. ELife, 2015, 4, e09269.	6.0	187
17	COLD1 Confers Chilling Tolerance in Rice. Cell, 2015, 160, 1209-1221.	28.9	724
18	Polycomb repression in the regulation of growth and development in Arabidopsis. Current Opinion in Plant Biology, 2015, 23, 15-24.	7.1	153

#	Article	IF	CITATIONS
19	O-GlcNAc-mediated interaction between VER2 and TaGRP2 elicits TaVRN1 mRNA accumulation during vernalization in winter wheat. Nature Communications, 2014, 5, 4572.	12.8	108
20	Requirement of histone acetyltransferases HAM1 and HAM2 for epigenetic modification of FLC in regulating flowering in Arabidopsis. Journal of Plant Physiology, 2013, 170, 444-451.	3.5	66
21	A Semi-fragile Watermarking Tolerant of Laplacian Sharpening. , 2008, , .		10