## Yuanzhong Jiang

List of Publications by Year in descending order

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516710 677142 23 864 16 22 citations g-index h-index papers 23 23 23 847 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Genome-wide identification and characterization of the Populus WRKY transcription factor family and analysis of their expression in response to biotic and abiotic stresses. Journal of Experimental Botany, 2014, 65, 6629-6644.	4.8	186
2	WRKY33 interacts with WRKY12 protein to upâ€regulate <i>RAP2</i> . <i>2</i> during submergence induced hypoxia response in <i>Arabidopsis thaliana</i> . New Phytologist, 2021, 229, 106-125.	7.3	71
3	Hybrid speciation via inheritance of alternate alleles of parental isolating genes. Molecular Plant, 2021, 14, 208-222.	8.3	68
4	PtrWRKY19, a novel WRKY transcription factor, contributes to the regulation of pith secondary wall formation in Populus trichocarpa. Scientific Reports, 2016, 6, 18643.	3.3	65
5	Isolation and characterization of a subgroup Ila WRKY transcription factor PtrWRKY40 from <i>Populus trichocarpa</i> . Tree Physiology, 2015, 35, 1129-1139.	3.1	55
6	The WRKY transcription factors PtrWRKY18 and PtrWRKY35 promote Melampsora resistance in Populus. Tree Physiology, 2017, 37, 665-675.	3.1	49
7	The PalWRKY77 transcription factor negatively regulates salt tolerance and abscisic acid signaling in <i>Populus</i> . Plant Journal, 2021, 105, 1258-1273.	5.7	49
8	Transcriptional landscape of highly lignified poplar stems at single-cell resolution. Genome Biology, 2021, 22, 319.	8.8	47
9	The poplar R2R3 MYB transcription factor PtrMYB94 coordinates with abscisic acid signaling to improve drought tolerance in plants. Tree Physiology, 2020, 40, 46-59.	3.1	35
10	The ubiquitin E3 ligase SR1 modulates the submergence response by degrading phosphorylated WRKY33 in <i>Arabidopsis</i> . Plant Cell, 2021, 33, 1771-1789.	6.6	34
11	Overexpression of Poplar PtrWRKY89 in Transgenic Arabidopsis Leads to a Reduction of Disease Resistance by Regulating Defense-Related Genes in Salicylate- and Jasmonate-Dependent Signaling. PLoS ONE, 2016, 11, e0149137.	2.5	33
12	Constitutive expression of the poplar WRKY transcription factor PtoWRKY60 enhances resistance to Dothiorella gregaria Sacc. in transgenic plants. Tree Physiology, 2014, 34, 1118-1129.	3.1	26
13	The Uâ€box E3 ubiquitin ligase PalPUB79 positively regulates ABAâ€dependent drought tolerance via ubiquitination of PalWRKY77 in <i>Populus</i> ). Plant Biotechnology Journal, 2021, 19, 2561-2575.	8.3	26
14	Heterologous gene silencing induced by tobacco rattle virus (TRV) is efficient for pursuing functional genomics studies in woody plants. Plant Cell, Tissue and Organ Culture, 2014, 116, 163-174.	2.3	24
15	The PalERF109 transcription factor positively regulates salt tolerance via PalHKT1;2 in Populus alba var. pyramidalis. Tree Physiology, 2020, 40, 717-730.	3.1	22
16	CHYR1 ubiquitinates the phosphorylated WRKY70 for degradation to balance immunity in <i>Arabidopsis thaliana </i> New Phytologist, 2021, 230, 1095-1109.	7.3	22
17	Allelic shift in cis-elements of the transcription factor <i>RAP2.12</i> underlies adaptation associated with humidity in <i>Arabidopsis thaliana</i> Science Advances, 2022, 8, eabn8281.	10.3	15
18	One AP2/ERF Transcription Factor Positively Regulates Pi Uptake and Drought Tolerance in Poplar. International Journal of Molecular Sciences, 2022, 23, 5241.	4.1	10

#	Article	lF	CITATIONS
19	PtoNF-YC9-SRMT-PtoRD26 module regulates the high saline tolerance of a triploid poplar. Genome Biology, 2022, 23, .	8.8	10
20	Heterologous Expression of Poplar WRKY18/35 Paralogs in Arabidopsis Reveals Their Antagonistic Regulation on Pathogen Resistance and Abiotic Stress Tolerance via Variable Hormonal Pathways. International Journal of Molecular Sciences, 2020, 21, 5440.	4.1	9
21	Genome-Wide Analysis of the Homeobox Gene Family and Identification of Drought-Responsive Members in Populus trichocarpa. Plants, 2021, 10, 2284.	3.5	4
22	Molecular signatures of parallel adaptive divergence causing reproductive isolation and speciation across two genera. Innovation(China), 2022, 3, 100247.	9.1	4
23	The complete chloroplast genome sequence of <i>Olmediella betschleriana</i> and its phylogenetic analysis. Mitochondrial DNA Part B: Resources, 2019, 4, 521-522.	0.4	0