

# Wan-Chen Li

## List of Publications by Year in descending order

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33  
papers

726  
citations

471509

17  
h-index

552781

26  
g-index

33  
all docs

33  
docs citations

33  
times ranked

724  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cloning and characterization of BES1/BZR1 transcription factor genes in maize. <i>Plant Growth Regulation</i> , 2018, 86, 235-249.	3.4	62
2	Maize ZmBES1/BZR1-5 Decreases ABA Sensitivity and Confers Tolerance to Osmotic Stress in Transgenic Arabidopsis. <i>International Journal of Molecular Sciences</i> , 2020, 21, 996.	4.1	53
3	Interaction network of core ABA signaling components in maize. <i>Plant Molecular Biology</i> , 2018, 96, 245-263.	3.9	51
4	Differential Expression of MicroRNAs in Response to Drought Stress in Maize. <i>Journal of Integrative Agriculture</i> , 2013, 12, 1414-1422.	3.5	50
5	Maize transcription factor ZmBES1/BZR1-5 positively regulates kernel size. <i>Journal of Experimental Botany</i> , 2021, 72, 1714-1726.	4.8	46
6	Quantitative Trait Loci for Resistance to Banded Leaf and Sheath Blight in Maize. <i>Crop Science</i> , 2006, 46, 1039-1045.	1.8	42
7	Cloning and Characterization of Functional Trehalose-6-Phosphate Synthase Gene in Maize. <i>Journal of Plant Biology</i> , 2010, 53, 134-141.	2.1	36
8	Improvement of resistance to maize dwarf mosaic virus mediated by transgenic RNA interference. <i>Journal of Biotechnology</i> , 2011, 153, 181-187.	3.8	35
9	A betaine aldehyde dehydrogenase gene from <i>Ammopiptanthus nanus</i> enhances tolerance of Arabidopsis to high salt and drought stresses. <i>Plant Growth Regulation</i> , 2017, 83, 265-276.	3.4	32
10	Expression Profile of Maize MicroRNAs Corresponding to Their Target Genes Under Drought Stress. <i>Biochemical Genetics</i> , 2014, 52, 474-493.	1.7	26
11	Introgression of Perennial Teosinte Genome into Maize and Identification of Genomic In Situ Hybridization and Microsatellite Markers. <i>Crop Science</i> , 2005, 45, 717-721.	1.8	24
12	RNA Interference-Based Transgenic Maize Resistant to Maize Dwarf Mosaic Virus. <i>Journal of Plant Biology</i> , 2010, 53, 297-305.	2.1	24
13	Interaction between abscisic acid receptor PYL3 and protein phosphatase type 2C in response to ABA signaling in maize. <i>Gene</i> , 2014, 549, 179-185.	2.2	24
14	Heterologous expression of betaine aldehyde dehydrogenase gene from <i>Ammopiptanthus nanus</i> confers high salt and heat tolerance to <i>Escherichia coli</i> . <i>Gene</i> , 2014, 549, 77-84.	2.2	24
15	Mutation loci and intragenic selection marker of the granule-bound starch synthase gene in waxy maize. <i>Molecular Breeding</i> , 2007, 20, 93-102.	2.1	20
16	Evaluation and Quantitative Inheritance of Several Drought-Relative Traits in Maize. <i>Agricultural Sciences in China</i> , 2008, 7, 280-290.	0.6	20
17	Heterologous expression of antifreeze protein gene AnAFP from <i>Ammopiptanthus nanus</i> enhances cold tolerance in <i>Escherichia coli</i> and tobacco. <i>Gene</i> , 2014, 539, 132-140.	2.2	20
18	ZmPP2C26 Alternative Splicing Variants Negatively Regulate Drought Tolerance in Maize. <i>Frontiers in Plant Science</i> , 2022, 13, 851531.	3.6	19

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19	Differential Expression of Serine/Threonine Protein Phosphatase Type-2C Under Drought Stress in Maize. <i>Plant Molecular Biology Reporter</i> , 2009, 27, 29-37.	1.8	16
20	RNA interference-mediated resistance to maize dwarf mosaic virus. <i>Plant Cell, Tissue and Organ Culture</i> , 2013, 113, 571-578.	2.3	16
21	Isolation and identification of a vegetative organ-specific promoter from maize. <i>Physiology and Molecular Biology of Plants</i> , 2019, 25, 277-287.	3.1	14
22	Combinatorial interaction of two adjacent cis-active promoter regions mediates the synergistic induction of Bt2 gene by sucrose and ABA in maize endosperm. <i>Plant Science</i> , 2018, 274, 332-340.	3.6	14
23	Maize ZmBES1/BZR1-3 and -9 Transcription Factors Negatively Regulate Drought Tolerance in Transgenic Arabidopsis. <i>International Journal of Molecular Sciences</i> , 2022, 23, 6025.	4.1	11
24	Ectopic expression of antifreeze protein gene from <i>Ammopiptanthus nanus</i> confers chilling tolerance in maize. <i>Crop Journal</i> , 2021, 9, 924-933.	5.2	10
25	Differential Gene Expression in Response to Drought Stress in Maize Seedling. <i>Agricultural Sciences in China</i> , 2009, 8, 767-776.	0.6	9
26	Antifreeze protein from <i>Ammopiptanthus nanus</i> functions in temperature-stress through domain A. <i>Scientific Reports</i> , 2021, 11, 8458.	3.3	6
27	Cloning and truncation modification of trehalose-6-phosphate synthase gene from <i>Selaginella pulvinata</i> . <i>Gene</i> , 2013, 512, 414-421.	2.2	4
28	Functional polymorphism among members of abscisic acid receptor family (ZmPYL) in maize. <i>Journal of Integrative Agriculture</i> , 2020, 19, 2165-2176.	3.5	4
29	Genome-wide analysis of BES1/BZR1 transcription factors and their responses to osmotic stress in <i>Ammopiptanthus nanus</i> . <i>Journal of Forest Research</i> , 2021, 26, 127-135.	1.4	4
30	Characterization of phenylalanine ammonia-lyase genes facilitating flavonoid biosynthesis from two species of medicinal plant <i>Anoectochilus</i> . <i>PeerJ</i> , 0, 10, e13614.	2.0	4
31	Overexpression of vacuolar H <sup>+</sup> pyrophosphatase (H <sup>+</sup> PPase) gene from <i>Ammopiptanthus nanus</i> enhances drought tolerance in maize. <i>Journal of Agronomy and Crop Science</i> , 0, , .	3.5	3
32	Zinc Transporter ZmLAZ1-4 Modulates Zinc Homeostasis on Plasma and Vacuolar Membrane in Maize. <i>Frontiers in Plant Science</i> , 2022, 13, 881055.	3.6	2
33	Genome-Wide Identification and Expression Analyses of AnSnRK2 Gene Family under Osmotic Stress in <i>Ammopiptanthus nanus</i> . <i>Plants</i> , 2021, 10, 882.	3.5	1