

Shou-Yi Chen

List of Publications by Year in descending order

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97
papers

11,047
citations

30047

54
h-index

34964

98
g-index

100
all docs

100
docs citations

100
times ranked

9864
citing authors

#	ARTICLE	IF	CITATIONS
1	AtNAC2, a transcription factor downstream of ethylene and auxin signaling pathways, is involved in salt stress response and lateral root development. <i>Plant Journal</i> , 2005, 44, 903-916.	2.8	634
2	Soybean WRKY-type transcription factor genes, <i>GmWRKY13</i> , <i>GmWRKY21</i> , and <i>GmWRKY54</i> , confer differential tolerance to abiotic stresses in transgenic <i>Arabidopsis</i> plants. <i>Plant Biotechnology Journal</i> , 2008, 6, 486-503.	4.1	582
3	De novo assembly of soybean wild relatives for pan-genome analysis of diversity and agronomic traits. <i>Nature Biotechnology</i> , 2014, 32, 1045-1052.	9.4	535
4	Melatonin enhances plant growth and abiotic stress tolerance in soybean plants. <i>Journal of Experimental Botany</i> , 2015, 66, 695-707.	2.4	493
5	Modulation of Ethylene Responses Affects Plant Salt-Stress Responses. <i>Plant Physiology</i> , 2007, 143, 707-719.	2.3	474
6	Soybean NAC transcription factors promote abiotic stress tolerance and lateral root formation in transgenic plants. <i>Plant Journal</i> , 2011, 68, 302-313.	2.8	471
7	Wheat <i>WRKY</i> genes <i>TaWRKY2</i> and <i>TaWRKY19</i> regulate abiotic stress tolerance in transgenic <i>Arabidopsis</i> plants. <i>Plant, Cell and Environment</i> , 2012, 35, 1156-1170.	2.8	377
8	Soybean <i>GmbZIP44</i> , <i>GmbZIP62</i> and <i>GmbZIP78</i> genes function as negative regulator of ABA signaling and confer salt and freezing tolerance in transgenic <i>Arabidopsis</i> . <i>Planta</i> , 2008, 228, 225-240.	1.6	350
9	Receptor-like kinase <i>OsSIK1</i> improves drought and salt stress tolerance in rice (<i>Oryza sativa</i>) plants. <i>Plant Journal</i> , 2010, 62, 316-329.	2.8	335
10	Identification of miRNAs and their target genes in developing soybean seeds by deep sequencing. <i>BMC Plant Biology</i> , 2011, 11, 5.	1.6	287
11	Melatonin delays leaf senescence and enhances salt stress tolerance in rice. <i>Journal of Pineal Research</i> , 2015, 59, 91-101.	3.4	272
12	The Role of Ethylene in Plants Under Salinity Stress. <i>Frontiers in Plant Science</i> , 2015, 6, 1059.	1.7	246
13	The soybean Dof-type transcription factor genes, <i>GmDof4</i> and <i>GmDof11</i> , enhance lipid content in the seeds of transgenic <i>Arabidopsis</i> plants. <i>Plant Journal</i> , 2007, 52, 716-729.	2.8	217
14	The Ethylene Receptor <i>ETR2</i> Delays Floral Transition and Affects Starch Accumulation in Rice. <i>Plant Cell</i> , 2009, 21, 1473-1494.	3.1	205
15	Soybean <i>GmMYB76</i> , <i>GmMYB92</i> , and <i>GmMYB177</i> genes confer stress tolerance in transgenic <i>Arabidopsis</i> plants. <i>Cell Research</i> , 2008, 18, 1047-1060.	5.7	204
16	<i>GmWRKY27</i> interacts with <i>GmMYB174</i> to reduce expression of <i>GmNAC29</i> for stress tolerance in soybean plants. <i>Plant Journal</i> , 2015, 83, 224-236.	2.8	199
17	Ethylene Signaling in Rice and <i>Arabidopsis</i> : Conserved and Diverged Aspects. <i>Molecular Plant</i> , 2015, 8, 495-505.	3.9	171
18	<i>MAOHUZI6/ETHYLENE INSENSITIVE3-LIKE1</i> and <i>ETHYLENE INSENSITIVE3-LIKE2</i> Regulate Ethylene Response of Roots and Coleoptiles and Negatively Affect Salt Tolerance in Rice. <i>Plant Physiology</i> , 2015, 169, 148-165.	2.3	163

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19	Soybean DRE-binding transcription factors that are responsive to abiotic stresses. <i>Theoretical and Applied Genetics</i> , 2005, 110, 1355-1362.	1.8	156
20	A PP2C-1 Allele Underlying a Quantitative Trait Locus Enhances Soybean 100-Seed Weight. <i>Molecular Plant</i> , 2017, 10, 670-684.	3.9	144
21	Genome-Wide Analysis of DNA Methylation in Soybean. <i>Molecular Plant</i> , 2013, 6, 1961-1974.	3.9	143
22	Plant NAC-type transcription factor proteins contain a NARD domain for repression of transcriptional activation. <i>Planta</i> , 2010, 232, 1033-1043.	1.6	135
23	An R2R3-type transcription factor gene <i>AtMYB59</i> regulates root growth and cell cycle progression in <i>Arabidopsis</i> . <i>Cell Research</i> , 2009, 19, 1291-1304.	5.7	131
24	Soybean Trihelix Transcription Factors <i>GmGT-2A</i> and <i>GmGT-2B</i> Improve Plant Tolerance to Abiotic Stresses in Transgenic <i>Arabidopsis</i> . <i>PLoS ONE</i> , 2009, 4, e6898.	1.1	120
25	Identification of Rice Ethylene-Response Mutants and Characterization of <i>MHZ7/OsEIN2</i> in Distinct Ethylene Response and Yield Trait Regulation. <i>Molecular Plant</i> , 2013, 6, 1830-1848.	3.9	117
26	The transcriptomic signature of developing soybean seeds reveals the genetic basis of seed trait adaptation during domestication. <i>Plant Journal</i> , 2016, 86, 530-544.	2.8	113
27	An S-Domain Receptor-Like Kinase, <i>OsSIK2</i> , Confers Abiotic Stress Tolerance and Delays Dark-Induced Leaf Senescence in Rice. <i>Plant Physiology</i> , 2013, 163, 1752-1765.	2.3	110
28	Ethylene-Inhibited Jasmonic Acid Biosynthesis Promotes Mesocotyl/Coleoptile Elongation of Etiolated Rice Seedlings. <i>Plant Cell</i> , 2017, 29, 1053-1072.	3.1	109
29	Ethylene Responses in Rice Roots and Coleoptiles Are Differentially Regulated by a Carotenoid Isomerase-Mediated Abscisic Acid Pathway. <i>Plant Cell</i> , 2015, 27, 1061-1081.	3.1	107
30	A rice transcription factor <i>OsBHLH1</i> is involved in cold stress response. <i>Theoretical and Applied Genetics</i> , 2003, 107, 1402-1409.	1.8	106
31	Ethylene-Induced Inhibition of Root Growth Requires Abscisic Acid Function in Rice (<i>Oryza sativa</i> L.) Seedlings. <i>PLoS Genetics</i> , 2014, 10, e1004701.	1.5	103
32	Expression of tobacco ethylene receptor <i>NTHK1</i> alters plant responses to salt stress. <i>Plant, Cell and Environment</i> , 2006, 29, 1210-1219.	2.8	99
33	Ethylene signaling regulates salt stress response. <i>Plant Signaling and Behavior</i> , 2008, 3, 761-763.	1.2	98
34	Characterization of a DRE-binding transcription factor from a halophyte <i>Atriplex hortensis</i> . <i>Theoretical and Applied Genetics</i> , 2003, 107, 155-161.	1.8	94
35	Soybean <i>GmPHD</i> -Type Transcription Regulators Improve Stress Tolerance in Transgenic <i>Arabidopsis</i> Plants. <i>PLoS ONE</i> , 2009, 4, e7209.	1.1	93
36	A class B heat shock factor selected for during soybean domestication contributes to salt tolerance by promoting flavonoid biosynthesis. <i>New Phytologist</i> , 2020, 225, 268-283.	3.5	92

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37	Serine/threonine kinase activity in the putative histidine kinase-like ethylene receptor NTHK1 from tobacco. <i>Plant Journal</i> , 2003, 33, 385-393.	2.8	91
38	Ethylene signaling in rice and <i>Arabidopsis</i> : New regulators and mechanisms. <i>Journal of Integrative Plant Biology</i> , 2021, 63, 102-125.	4.1	91
39	EIN2 regulates salt stress response and interacts with a MA3 domain-containing protein ECIP1 in <i>Arabidopsis</i> . <i>Plant, Cell and Environment</i> , 2011, 34, 1678-1692.	2.8	90
40	OsGLU1, A Putative Membrane-bound Endo-1,4- α -D-glucanase from Rice, Affects Plant Internode Elongation. <i>Plant Molecular Biology</i> , 2006, 60, 137-151.	2.0	89
41	GmWRKY54 improves drought tolerance through activating genes in abscisic acid and Ca ²⁺ signaling pathways in transgenic soybean. <i>Plant Journal</i> , 2019, 100, 384-398.	2.8	87
42	A Putative Plasma Membrane Cation/proton Antiporter from Soybean Confers Salt Tolerance in <i>Arabidopsis</i> . <i>Plant Molecular Biology</i> , 2005, 59, 809-820.	2.0	86
43	QTL mapping of phosphorus deficiency tolerance in soybean (<i>Glycine max</i> L. Merr.). <i>Euphytica</i> , 2005, 142, 137-142.	0.6	84
44	Characterization of soybean genomic features by analysis of its expressed sequence tags. <i>Theoretical and Applied Genetics</i> , 2004, 108, 903-913.	1.8	83
45	Soybean GmMYB73 promotes lipid accumulation in transgenic plants. <i>BMC Plant Biology</i> , 2014, 14, 73.	1.6	83
46	Soybean GmbZIP123 gene enhances lipid content in the seeds of transgenic <i>Arabidopsis</i> plants. <i>Journal of Experimental Botany</i> , 2013, 64, 4329-4341.	2.4	81
47	Soybean miR172a Improves Salt Tolerance and Can Function as a Long-Distance Signal. <i>Molecular Plant</i> , 2016, 9, 1337-1340.	3.9	74
48	Role of Soybean GmbZIP132 under Abscisic Acid and Salt Stresses. <i>Journal of Integrative Plant Biology</i> , 2008, 50, 221-230.	4.1	73
49	Selection for a Zinc-Finger Protein Contributes to Seed Oil Increase during Soybean Domestication. <i>Plant Physiology</i> , 2017, 173, 2208-2224.	2.3	73
50	AhCMO, regulated by stresses in <i>Atriplex hortensis</i> , can improve drought tolerance in transgenic tobacco. <i>Theoretical and Applied Genetics</i> , 2002, 105, 815-821.	1.8	72
51	An AP2/EREBP-type transcription-factor gene from rice is cold-inducible and encodes a nuclear-localized protein. <i>Theoretical and Applied Genetics</i> , 2003, 107, 972-979.	1.8	66
52	E3 ubiquitin ligase SOR1 regulates ethylene response in rice root by modulating stability of Aux/IAA protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 4513-4518.	3.3	66
53	The <i>A</i> -like homeodomain finger protein <i>AL</i> 5 suppresses multiple negative factors to confer abiotic stress tolerance in <i>Arabidopsis</i> . <i>Plant Journal</i> , 2015, 81, 871-883.	2.8	60
54	DREB1C from <i>Medicago truncatula</i> enhances freezing tolerance in transgenic <i>M. truncatula</i> and China Rose (<i>Rosa chinensis</i> Jacq.). <i>Plant Growth Regulation</i> , 2010, 60, 199-211.	1.8	59

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55	Evidence for Serine/Threonine and Histidine Kinase Activity in the Tobacco Ethylene Receptor Protein NTHK2. <i>Plant Physiology</i> , 2004, 136, 2971-2981.	2.3	58
56	Effects of Tobacco Ethylene Receptor Mutations on Receptor Kinase Activity, Plant Growth and Stress Responses. <i>Plant and Cell Physiology</i> , 2009, 50, 1636-1650.	1.5	53
57	Isolation and Characterization of a Salt- and Drought-inducible Gene for S-adenosylmethionine Decarboxylase from Wheat (<i>Triticum aestivum</i> L.). <i>Journal of Plant Physiology</i> , 2000, 156, 386-393.	1.6	49
58	Cloning and comparative analysis of the gene encoding diacylglycerol acyltransferase from wild type and cultivated soybean. <i>Theoretical and Applied Genetics</i> , 2006, 112, 1086-1097.	1.8	49
59	A new AOX homologous gene OsIM1 from rice (<i>Oryza sativa</i> L.) with an alternative splicing mechanism under salt stress. <i>Theoretical and Applied Genetics</i> , 2003, 107, 326-331.	1.8	48
60	The transcription factor AtDOF4.2 regulates shoot branching and seed coat formation in <i>Arabidopsis</i> . <i>Biochemical Journal</i> , 2013, 449, 373-388.	1.7	48
61	Diverse Roles of Ethylene in Regulating Agronomic Traits in Rice. <i>Frontiers in Plant Science</i> , 2017, 8, 1676.	1.7	47
62	Roles of ethylene receptor NTHK1 domains in plant growth, stress response and protein phosphorylation. <i>FEBS Letters</i> , 2006, 580, 1239-1250.	1.3	46
63	Trihelix transcription factor GT-4 mediates salt tolerance via interaction with TEM2 in <i>Arabidopsis</i> . <i>BMC Plant Biology</i> , 2014, 14, 339.	1.6	46
64	A Histone Code Reader and a Transcriptional Activator Interact to Regulate Genes for Salt Tolerance. <i>Plant Physiology</i> , 2017, 175, 1304-1320.	2.3	45
65	Enhancement of salt tolerance in alfalfa transformed with the gene encoding for betaine aldehyde dehydrogenase. <i>Euphytica</i> , 2011, 178, 363-372.	0.6	41
66	A transcriptional regulatory module controls lipid accumulation in soybean. <i>New Phytologist</i> , 2021, 231, 661-678.	3.5	38
67	Nuclear factor Y subunit GmNFYA competes with GmHDA13 for interaction with GmFVE to positively regulate salt tolerance in soybean. <i>Plant Biotechnology Journal</i> , 2021, 19, 2362-2379.	4.1	38
68	Membrane protein MHZ3 stabilizes OsEIN2 in rice by interacting with its Nramp-like domain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 2520-2525.	3.3	37
69	Histidine kinase MHZ1/OsHK1 interacts with ethylene receptors to regulate root growth in rice. <i>Nature Communications</i> , 2020, 11, 518.	5.8	37
70	The GDSL Lipase MHZ11 Modulates Ethylene Signaling in Rice Roots. <i>Plant Cell</i> , 2020, 32, 1626-1643.	3.1	36
71	Isolation and characterization of a full-length resistance gene homolog from soybean. <i>Theoretical and Applied Genetics</i> , 2003, 106, 786-793.	1.8	35
72	Tobacco Translationally Controlled Tumor Protein Interacts with Ethylene Receptor Tobacco Histidine Kinase1 and Enhances Plant Growth through Promotion of Cell Proliferation. <i>Plant Physiology</i> , 2015, 169, 96-114.	2.3	35

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73	OsDREB4 Genes in Rice Encode AP2-Containing Proteins that Bind Specifically to the Dehydration-Responsive Element. <i>Journal of Integrative Plant Biology</i> , 2005, 47, 467-476.	4.1	34
74	Soybean GmDREBL Increases Lipid Content in Seeds of Transgenic Arabidopsis. <i>Scientific Reports</i> , 2016, 6, 34307.	1.6	34
75	NIMA-related kinase NEK6 affects plant growth and stress response in Arabidopsis. <i>Plant Journal</i> , 2011, 68, 830-843.	2.8	31
76	Tobacco Ankyrin Protein NEIP2 Interacts with Ethylene Receptor NTHK1 and Regulates Plant Growth and Stress Responses. <i>Plant and Cell Physiology</i> , 2015, 56, 803-818.	1.5	31
77	Spatial Expression and Characterization of a Putative Ethylene Receptor Protein NTHK1 in Tobacco. <i>Plant and Cell Physiology</i> , 2002, 43, 810-815.	1.5	30
78	An Alfin-like gene from <i>Atriplex hortensis</i> enhances salt and drought tolerance and abscisic acid response in transgenic Arabidopsis. <i>Scientific Reports</i> , 2018, 8, 2707.	1.6	30
79	Genomic characterization of the S-adenosylmethionine decarboxylase genes from soybean. <i>Theoretical and Applied Genetics</i> , 2004, 108, 842-850.	1.8	26
80	Leveraging <i>Atriplex hortensis</i> choline monooxygenase to improve chilling tolerance in cotton. <i>Environmental and Experimental Botany</i> , 2019, 162, 364-373.	2.0	26
81	Cloning and characterization of an HDZip I gene GmHZ1 from soybean. <i>Planta</i> , 2005, 221, 831-843.	1.6	25
82	Soybean NIMA-Related Kinase1 Promotes Plant Growth and Improves Salt and Cold Tolerance. <i>Plant and Cell Physiology</i> , 2017, 58, 1268-1278.	1.5	22
83	Isolation and characterization of a Pti1 homologue from soybean. <i>Journal of Experimental Botany</i> , 2004, 55, 535-537.	2.4	21
84	Two New Group 3 LEA Genes of Wheat and Their Functional Analysis in Yeast. <i>Journal of Integrative Plant Biology</i> , 2005, 47, 1372-1381.	4.1	21
85	Analysis of expressed receptor-like kinases (RLKs) in soybean. <i>Journal of Genetics and Genomics</i> , 2009, 36, 611-619.	1.7	20
86	The OsEIL1-OsERF115 target gene regulatory module controls grain size and weight in rice. <i>Plant Biotechnology Journal</i> , 2022, 20, 1470-1486.	4.1	20
87	Activation of a DRE-binding transcription factor from <i>Medicago truncatula</i> by deleting a Ser/Thr-rich region. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 2009, 45, 1-11.	0.9	17
88	Characterization of a novel cell cycle-related gene from Arabidopsis. <i>Journal of Experimental Botany</i> , 2005, 56, 807-816.	2.4	16
89	The Putative Ser/Thr Protein Kinase Gene GmAAPK from Soybean is Regulated by Abiotic Stress. <i>Journal of Integrative Plant Biology</i> , 2006, 48, 327-333.	4.1	16
90	Ethylene Biosynthesis, Signaling, and Crosstalk with Other Hormones in Rice. <i>Small Methods</i> , 2020, 4, 1900278.	4.6	16

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91	Development of Soybean EST-SSR Markers and Their Use to Assess Genetic Diversity in the Subgenus Soja. <i>Agricultural Sciences in China</i> , 2010, 9, 1423-1429.	0.6	14
92	Roles of Ethylene in Plant Growth and Responses to Stresses. , 2014, , 81-118.		11
93	Rapid Construction of a Plant RNA Interference Expression Vector for Hairpin RNA-Mediated Targeting Using a PCR-Based Method. <i>DNA and Cell Biology</i> , 2009, 28, 605-613.	0.9	5
94	RNA Extraction and Preparation in Rice (<i>Oryza sativa</i>). <i>Current Protocols in Plant Biology</i> , 2016, 1, 411-418.	2.8	5
95	The continuous accumulation of Na ⁺ in detached leaf sections is associated with over-expression of NTHK1 and salt tolerance in poplar plants. <i>Functional Plant Biology</i> , 2011, 38, 236.	1.1	4
96	Establishment of a transgenic system in fast-growing black locust (<i>Robinia pseudoacacia</i> L.). <i>Forestry Studies in China</i> , 2008, 10, 243-252.	0.4	3
97	Simple Methods for Screening and Statistical Analysis of Leaf Epidermal Cells in Dicotyledonous Plants. <i>Bio-protocol</i> , 2016, 6, .	0.2	3