## Siyi Guo

## List of Publications by Year in descending order

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		109321	106344
67	5,461	35	65
papers	5,461 citations	h-index	g-index
68	68	68	6659
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Abscisic acid dynamics, signaling, and functions in plants. Journal of Integrative Plant Biology, 2020, 62, 25-54.	8.5	771
2	An Arabidopsis Glutathione Peroxidase Functions as Both a Redox Transducer and a Scavenger in Abscisic Acid and Drought Stress Responses. Plant Cell, 2006, 18, 2749-2766.	6.6	466
3	Reactive oxygen species signaling and stomatal movement in plant responses to drought stress and pathogen attack. Journal of Integrative Plant Biology, 2018, 60, 805-826.	8 <b>.</b> 5	397
4	Nitric oxide negatively regulates abscisic acid signaling in guard cells by S-nitrosylation of OST1.  Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 613-618.	7.1	318
5	The interaction between OsMADS57 and OsTB1 modulates rice tillering via DWARF14. Nature Communications, 2013, 4, 1566.	12.8	266
6	Timing Mechanism Dependent on Cell Division Is Invoked by Polycomb Eviction in Plant Stem Cells. Science, 2014, 343, 1248559.	12.6	197
7	OsmiR396d-Regulated OsGRFs Function in Floral Organogenesis in Rice through Binding to Their Targets <i>OsJMJ706</i> and <i>OsCR4</i> Plant Physiology, 2014, 165, 160-174.	4.8	172
8	A RAF-SnRK2 kinase cascade mediates early osmotic stress signaling in higher plants. Nature Communications, 2020, 11, 613.	12.8	147
9	The SOS2-SCaBP8 Complex Generates and Fine-Tunes an AtANN4-Dependent Calcium Signature under Salt Stress. Developmental Cell, 2019, 48, 697-709.e5.	7.0	133
10	Dynamics of Brassinosteroid Response Modulated by Negative Regulator LIC in Rice. PLoS Genetics, 2012, 8, e1002686.	<b>3.</b> 5	130
11	OsmiR396d Affects Gibberellin and Brassinosteroid Signaling to Regulate Plant Architecture in Rice. Plant Physiology, 2018, 176, 946-959.	4.8	127
12	A Receptor-Like Kinase Mediates Ammonium Homeostasis and Is Important for the Polar Growth of Root Hairs in <i>Arabidopsis</i> <li>Plant Cell, 2014, 26, 1497-1511.</li>	6.6	124
13	Overexpression of stressâ€inducible <scp>OsBURP16</scp> , the <i>β</i> subunit of polygalacturonase 1, decreases pectin content and cell adhesion and increases abiotic stress sensitivity in rice. Plant, Cell and Environment, 2014, 37, 1144-1158.	5.7	122
14	AIK1, A Mitogen-Activated Protein Kinase, Modulates Abscisic Acid Responses through the MKK5-MPK6 Kinase Cascade. Plant Physiology, 2017, 173, 1391-1408.	4.8	117
15	The Arabidopsis MYB transcription factor, MYB111 modulates salt responses by regulating flavonoid biosynthesis. Environmental and Experimental Botany, 2019, 166, 103807.	4.2	117
16	Overexpression of a homopeptide repeat-containing bHLH protein gene (OrbHLH001) from Dongxiang Wild Rice confers freezing and salt tolerance in transgenic Arabidopsis. Plant Cell Reports, 2010, 29, 977-986.	<b>5.</b> 6	111
17	BRASSINOSTEROID-INSENSITIVE2 Negatively Regulates the Stability of Transcription Factor ICE1 in Response to Cold Stress in Arabidopsis. Plant Cell, 2019, 31, tpc.00058.2019.	6.6	110
18	MiR156 regulates anthocyanin biosynthesis through SPL targets and other microRNAs in poplar. Horticulture Research, 2020, 7, 118.	6.3	90

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19	Abscisic Acid as an Internal Integrator of Multiple Physiological Processes Modulates Leaf Senescence Onset in Arabidopsis thaliana. Frontiers in Plant Science, 2016, 7, 181.	3.6	89
20	Proteomics and Phosphoproteomics of Heat Stress-Responsive Mechanisms in Spinach. Frontiers in Plant Science, 2018, 9, 800.	3.6	79
21	The Cyclophilin CYP20-2 Modulates the Conformation of BRASSINAZOLE-RESISTANT1, Which Binds the Promoter of FLOWERING LOCUS D to Regulate Flowering in Arabidopsis. Plant Cell, 2013, 25, 2504-2521.	6.6	78
22	ABI5 modulates seed germination via feedback regulation of the expression of the <i>PYR/PYL/RCAR</i> ABA receptor genes. New Phytologist, 2020, 228, 596-608.	7.3	78
23	A Membrane-Bound NAC-Like Transcription Factor OsNTL5 Represses the Flowering in Oryza sativa. Frontiers in Plant Science, 2018, 9, 555.	3.6	77
24	Biosynthesis of DHGA12 and its roles in Arabidopsis seedling establishment. Nature Communications, 2019, 10, 1768.	12.8	72
25	BZU2/ZmMUTE controls symmetrical division of guard mother cell and specifies neighbor cell fate in maize. PLoS Genetics, 2019, 15, e1008377.	3.5	64
26	An amplification-selection model for quantified rhizosphere microbiota assembly. Science Bulletin, 2020, 65, 983-986.	9.0	64
27	Flavonoids improve drought tolerance of maize seedlings by regulating the homeostasis of reactive oxygen species. Plant and Soil, 2021, 461, 389-405.	3.7	64
28	Trehaloseâ€6â€phosphate phosphatase E modulates ABAâ€controlled root growth and stomatal movement in <i>Arabidopsis</i> . Journal of Integrative Plant Biology, 2020, 62, 1518-1534.	8.5	58
29	A vacuole localized β-glucosidase contributes to drought tolerance in Arabidopsis. Science Bulletin, 2011, 56, 3538-3546.	1.7	55
30	Modulation of Guard Cell Turgor and Drought Tolerance by a Peroxisomal Acetate–Malate Shunt. Molecular Plant, 2018, 11, 1278-1291.	8.3	53
31	Reduced expression of a gene encoding a Golgi localized monosaccharide transporter (OsGMST1) confers hypersensitivity to salt in rice (Oryza sativa). Journal of Experimental Botany, 2011, 62, 4595-4604.	4.8	42
32	Beyond Light: Insights Into the Role of Constitutively Photomorphogenic1 in Plant Hormonal Signaling. Frontiers in Plant Science, 2019, 10, 557.	3.6	42
33	Physiological and comparative proteomic analyses of saline-alkali NaHCO3-responses in leaves of halophyte Puccinellia tenuiflora. Plant and Soil, 2019, 437, 137-158.	3.7	41
34	The RING E3 ligase CLG1 targets GS3 for degradation via the endosome pathway to determine grain size in rice. Molecular Plant, 2021, 14, 1699-1713.	8.3	41
35	Mutation of 4-coumarate: coenzyme A ligase 1 gene affects lignin biosynthesis and increases the cell wall digestibility in maize brown midrib5 mutants. Biotechnology for Biofuels, 2019, 12, 82.	6.2	40
36	Antagonistic Interaction between Auxin and SA Signaling Pathways Regulates Bacterial Infection through Lateral Root in Arabidopsis. Cell Reports, 2020, 32, 108060.	6.4	38

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37	Analysis of Global Expression Profiles of Arabidopsis Genes Under Abscisic Acid and H2O2 Applications. Journal of Integrative Plant Biology, 2006, 48, 62-74.	8.5	36
38	Co-ordination of Flower Development Through Epigenetic Regulation in Two Model Species: Rice and Arabidopsis. Plant and Cell Physiology, 2015, 56, 830-842.	3.1	35
39	COP1 promotes ABAâ€induced stomatal closure by modulating the abundance of ABI/HAB and AHG3 phosphatases. New Phytologist, 2021, 229, 2035-2049.	7.3	32
40	PIFs coordinate shade avoidance by inhibiting auxin repressor <i>ARF18</i> and metabolic regulator <i>QQS</i> . New Phytologist, 2020, 228, 609-621.	7.3	29
41	Plant Chloroplast Stress Response: Insights from Thiol Redox Proteomics. Antioxidants and Redox Signaling, 2020, 33, 35-57.	5.4	29
42	From mouse to mouseâ€ear cress: Nanomaterials as vehicles in plant biotechnology. Exploration, 2021, 1, 9-20.	11.0	27
43	Efficient Generation of CRISPR/Cas9-Mediated Homozygous/Biallelic Medicago truncatula Mutants Using a Hairy Root System. Frontiers in Plant Science, 2020, 11, 294.	3.6	25
44	Heat-Responsive Proteomics of a Heat-Sensitive Spinach Variety. International Journal of Molecular Sciences, 2019, 20, 3872.	4.1	23
45	Nod factor receptor complex phosphorylates GmGEF2 to stimulate ROP signaling during nodulation. Current Biology, 2021, 31, 3538-3550.e5.	3.9	22
46	Allele-defined genome reveals biallelic differentiation during cassava evolution. Molecular Plant, 2021, 14, 851-854.	8.3	20
47	NaCl-responsive ROS scavenging and energy supply in alkaligrass callus revealed from proteomic analysis. BMC Genomics, 2019, 20, 990.	2.8	19
48	AtHB7/12 Regulate Root Growth in Response to Aluminum Stress. International Journal of Molecular Sciences, 2020, 21, 4080.	4.1	19
49	Proteomic discovery of H2O2 response in roots and functional characterization of PutGLP gene from alkaligrass. Planta, 2018, 248, 1079-1099.	3.2	18
50	ABC1K10a, an atypical kinase, functions in plant salt stress tolerance. BMC Plant Biology, 2020, 20, 270.	3 <b>.</b> 6	15
51	The efficacy of anti-VEGF antibody-modified liposomes loaded with paeonol in the prevention and treatment of hypertrophic scars. Drug Development and Industrial Pharmacy, 2019, 45, 439-455.	2.0	14
52	Behaviour of cell penetrating peptide TAT-modified liposomes loaded with salvianolic acid B on the migration, proliferation, and survival of human skin fibroblasts. Journal of Liposome Research, 2020, 30, 93-106.	3.3	14
53	Screening of abiotic stressâ€responsive cotton genes using a cotton fullâ€length cDNA overexpressing <i>Arabidopsis</i> library. Journal of Integrative Plant Biology, 2020, 62, 998-1016.	8.5	12
54	The UBP14-CDKB1;1-CDKG2 cascade controls endoreduplication and cell growth in Arabidopsis. Plant Cell, 2022, 34, 1308-1325.	6.6	12

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55	Na2CO3-responsive Photosynthetic and ROS Scavenging Mechanisms in Chloroplasts of Alkaligrass Revealed by Phosphoproteomics. Genomics, Proteomics and Bioinformatics, 2020, 18, 271-288.	6.9	10
56	Large-area gold nanohole arrays fabricated by one-step method for surface plasmon resonance biochemical sensing. Science China Life Sciences, 2018, 61, 476-482.	4.9	8
57	GhWRKY46 from upland cotton positively regulates the drought and salt stress responses in plant. Environmental and Experimental Botany, 2021, 186, 104438.	4.2	8
58	The maize single-nucleus transcriptome comprehensively describes signaling networks governing movement and development of grass stomata. Plant Cell, 2022, , .	6.6	8
59	Selection and Validation of Reference Genes for RT-qPCR Analysis in Spinacia oleracea under Abiotic Stress. BioMed Research International, 2021, 2021, 1-12.	1.9	6
60	Overexpression of AHL9 accelerates leaf senescence in Arabidopsis thaliana. BMC Plant Biology, 2022, 22, 248.	3.6	6
61	Characterization of Two New brown midrib1 Mutations From an EMS-Mutagenic Maize Population for Lignocellulosic Biomass Utilization. Frontiers in Plant Science, 2020, 11, 594798.	3.6	5
62	Genome-wide identification and expression analysis reveals spinach brassinosteroid-signaling kinase (BSK) gene family functions in temperature stress response. BMC Genomics, 2022, 23, .	2.8	5
63	SICKLE modulates lateral root development by promoting degradation of lariat intronic RNA. Plant Physiology, 0, , .	4.8	4
64	A cytogenetic analysis of male meiosis in Asparagus officinalis. Bioscience, Biotechnology and Biochemistry, 2019, 83, 666-674.	1.3	3
65	A set of sampling, preparation, and staining techniques for studying meiosis in cucumber. Plant Science, 2022, 319, 111245.	3.6	3
66	Tanshinone IIA down-regulated p-Smad3 signaling to inhibit TGF-β1-mediated fibroblast proliferation via lncRNA-HSRL/SNX9. International Journal of Biochemistry and Cell Biology, 2020, 129, 105863.	2.8	2
67	Determination of UDP-Glucose and UDP-Galactose in Maize by Hydrophilic Interaction Liquid Chromatography and Tandem Mass Spectrometry. Journal of Analytical Methods in Chemistry, 2022, 2022, 1-6.	1.6	0