

# Yinbo Gan

## List of Publications by Year in descending order

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55  
papers

2,958  
citations

126907

33  
h-index

175258

52  
g-index

56  
all docs

56  
docs citations

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times ranked

3077  
citing authors

#	ARTICLE	IF	CITATIONS
1	Seed priming with zinc oxide nanoparticles downplayed ultrastructural damage and improved photosynthetic apparatus in maize under cobalt stress. <i>Journal of Hazardous Materials</i> , 2022, 423, 127021.	12.4	122
2	Exploring the Adaptive Responses of Plants to Abiotic Stresses Using Transcriptome Data. <i>Agriculture (Switzerland)</i> , 2022, 12, 211.	3.1	22
3	C2H2 Zinc Finger Proteins Response to Abiotic Stress in Plants. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2730.	4.1	41
4	Dichromate-induced ethylene biosynthesis, perception, and signaling regulate the variance in root growth inhibition among Shaheen basmati and basmati-385 rice varieties. <i>Environmental Science and Pollution Research</i> , 2021, 28, 38016-38025.	5.3	5
5	BnaA02.NIP6;1a encodes a boron transporter required for plant development under boron deficiency in <i>Brassica napus</i> . <i>Plant Physiology and Biochemistry</i> , 2021, 161, 36-45.	5.8	8
6	A rice transcription factor, <i>OsMADS57</i> , positively regulates high salinity tolerance in transgenic <i>Arabidopsis thaliana</i> and <i>Oryza sativa</i> plants. <i>Physiologia Plantarum</i> , 2021, 173, 1120-1135.	5.2	20
7	RNA-Seq-Based Profiling of pl Mutant Reveals Transcriptional Regulation of Anthocyanin Biosynthesis in Rice ( <i>Oryza sativa</i> L.). <i>International Journal of Molecular Sciences</i> , 2021, 22, 9787.	4.1	4
8	Ethylene participates in zinc oxide nanoparticles induced biochemical, molecular and ultrastructural changes in rice seedlings. <i>Ecotoxicology and Environmental Safety</i> , 2021, 226, 112844.	6.0	27
9	Overexpression of MADS-box transcription factor <i>OsMADS25</i> enhances salt stress tolerance in Rice and <i>Arabidopsis</i> . <i>Plant Growth Regulation</i> , 2020, 90, 163-171.	3.4	21
10	Ethylene mediates CuO NP-induced ultrastructural changes and oxidative stress in <i>Arabidopsis thaliana</i> leaves. <i>Environmental Science: Nano</i> , 2020, 7, 938-953.	4.3	24
11	Nitrate regulation of lateral root and root hair development in plants. <i>Journal of Experimental Botany</i> , 2020, 71, 4405-4414.	4.8	45
12	Zinc finger protein 5 (ZFP5) associates with ethylene signaling to regulate the phosphate and potassium deficiency-induced root hair development in <i>Arabidopsis</i> . <i>Plant Molecular Biology</i> , 2020, 102, 143-158.	3.9	39
13	Zinc Finger Protein 1 (ZFP1) Is Involved in Trichome Initiation in <i>Arabidopsis thaliana</i> . <i>Agriculture (Switzerland)</i> , 2020, 10, 645.	3.1	9
14	The WRKY6 transcription factor affects seed oil accumulation and alters fatty acid compositions in <i>Arabidopsis thaliana</i> . <i>Physiologia Plantarum</i> , 2020, 169, 612-624.	5.2	35
15	Chromium-Induced Reactive Oxygen Species Accumulation by Altering the Enzymatic Antioxidant System and Associated Cytotoxic, Genotoxic, Ultrastructural, and Photosynthetic Changes in Plants. <i>International Journal of Molecular Sciences</i> , 2020, 21, 728.	4.1	157
16	The new insight of auxin functions: transition from seed dormancy to germination and floral opening in plants. <i>Plant Growth Regulation</i> , 2020, 91, 169-174.	3.4	25
17	Involvement of ethylene signaling in zinc oxide nanoparticle-mediated biochemical changes in <i>Arabidopsis thaliana</i> leaves. <i>Environmental Science: Nano</i> , 2019, 6, 341-355.	4.3	50
18	Ethylene mediates dichromate-induced oxidative stress and regulation of the enzymatic antioxidant system-related transcriptome in <i>Arabidopsis thaliana</i> . <i>Environmental and Experimental Botany</i> , 2019, 161, 166-179.	4.2	50

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19	Ethylene mediates dichromate-induced inhibition of primary root growth by altering <i>AUX1</i> expression and auxin accumulation in <i>Arabidopsis thaliana</i> . <i>Plant, Cell and Environment</i> , 2018, 41, 1453-1467.	5.7	46
20	PIL5 represses floral transition in <i>Arabidopsis</i> under long day conditions. <i>Biochemical and Biophysical Research Communications</i> , 2018, 499, 513-518.	2.1	11
21	Involvement of histone acetylation and deacetylation in regulating auxin responses and associated phenotypic changes in plants. <i>Plant Cell Reports</i> , 2018, 37, 51-59.	5.6	14
22	NbGIS regulates glandular trichome initiation through GA signaling in tobacco. <i>Plant Molecular Biology</i> , 2018, 98, 153-167.	3.9	29
23	SPATULA regulates floral transition and photomorphogenesis in a PHYTOCHROME B-dependent manner in <i>Arabidopsis</i> . <i>Biochemical and Biophysical Research Communications</i> , 2018, 503, 2380-2385.	2.1	5
24	A model for the ethylene-mediated auxin distribution under Cr(VI) stress in <i>Arabidopsis thaliana</i> . <i>Plant Signaling and Behavior</i> , 2018, 13, 1-2.	2.4	1
25	OsFTIP7 determines auxin-mediated anther dehiscence in rice. <i>Nature Plants</i> , 2018, 4, 495-504.	9.3	63
26	The SPATULA transcription factor regulates seed oil content by controlling seed specific genes in <i>Arabidopsis thaliana</i> . <i>Plant Growth Regulation</i> , 2017, 82, 111-121.	3.4	14
27	OsFTIP1-Mediated Regulation of Florigen Transport in Rice Is Negatively Regulated by the Ubiquitin-Like Domain Kinase OsUbrK134. <i>Plant Cell</i> , 2017, 29, 491-507.	6.6	55
28	Biochemical responses and ultrastructural changes in ethylene insensitive mutants of <i>Arabidopsis thaliana</i> subjected to bisphenol A exposure. <i>Ecotoxicology and Environmental Safety</i> , 2017, 144, 62-71.	6.0	39
29	AtGIS, a C2H2 zinc-finger transcription factor from <i>Arabidopsis</i> regulates glandular trichome development through GA signaling in tobacco. <i>Biochemical and Biophysical Research Communications</i> , 2017, 483, 209-215.	2.1	40
30	Multiple phytohormones promote root hair elongation by regulating a similar set of genes in the root epidermis in <i>Arabidopsis</i> . <i>Journal of Experimental Botany</i> , 2016, 67, 6363-6372.	4.8	78
31	Toxicological effects of bisphenol A on growth and antioxidant defense system in <i>Oryza sativa</i> as revealed by ultrastructure analysis. <i>Ecotoxicology and Environmental Safety</i> , 2016, 124, 277-284.	6.0	62
32	Linkage Mapping of Stem Saccharification Digestibility in Rice. <i>PLoS ONE</i> , 2016, 11, e0159117.	2.5	6
33	The <i>Arabidopsis</i> Gene zinc finger protein 3 (ZFP3) Is Involved in Salt Stress and Osmotic Stress Response. <i>PLoS ONE</i> , 2016, 11, e0168367.	2.5	53
34	<i>GLABROUS INFLORESCENCE STEMS3</i> ( <i>GIS3</i> ) regulates trichome initiation and development in <i>Arabidopsis</i> . <i>New Phytologist</i> , 2015, 206, 220-230.	7.3	90
35	MADS-box Transcription Factor OsMADS25 Regulates Root Development through Affection of Nitrate Accumulation in Rice. <i>PLoS ONE</i> , 2015, 10, e0135196.	2.5	81
36	AtEXP2 Is Involved in Seed Germination and Abiotic Stress Response in <i>Arabidopsis</i> . <i>PLoS ONE</i> , 2014, 9, e85208.	2.5	93

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37	The Effects of Fluctuations in the Nutrient Supply on the Expression of Five Members of the AGL17 Clade of MADS-Box Genes in Rice. PLoS ONE, 2014, 9, e105597.	2.5	30
38	Involvement of C2H2 zinc finger proteins in the regulation of epidermal cell fate determination in <i>Arabidopsis</i> . Journal of Integrative Plant Biology, 2014, 56, 1112-1117.	8.5	39
39	<i>ZFP6</i> ( <i>ZFP6</i> ) regulates trichome initiation by integrating gibberellin and cytokinin signaling in <i>Arabidopsis thaliana</i> . New Phytologist, 2013, 198, 699-708.	7.3	144
40	Differential control of seed primary dormancy in <i>Arabidopsis</i> ecotypes by the transcription factor SPATULA. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 10866-10871.	7.1	106
41	<i>ZFP5</i> encodes a functionally equivalent <i>GIS</i> protein to control trichome initiation. Plant Signaling and Behavior, 2012, 7, 28-30.	2.4	32
42	GLABROUS INFLORESCENCE STEMS (GIS) is Required for Trichome Branching Through Gibberellic Acid Signaling in <i>Arabidopsis</i> . Plant and Cell Physiology, 2012, 53, 457-469.	3.1	45
43	Overexpressing the ANR1 MADS-Box Gene in Transgenic Plants Provides New Insights into its Role in the Nitrate Regulation of Root Development. Plant and Cell Physiology, 2012, 53, 1003-1016.	3.1	103
44	A zinc finger protein gene <i>ZFP5</i> integrates phytohormone signaling to control root hair development in <i>Arabidopsis</i> . Plant Journal, 2012, 72, 474-490.	5.7	79
45	The responses of trichome mutants to enhanced ultraviolet-B radiation in <i>Arabidopsis thaliana</i> . Journal of Photochemistry and Photobiology B: Biology, 2012, 113, 29-35.	3.8	83
46	<i>Zinc Finger Protein5</i> is Required for the Control of Trichome Initiation by Acting Upstream of <i>Zinc Finger Protein8</i> in <i>Arabidopsis</i> . Plant Physiology, 2011, 157, 673-682.	4.8	106
47	Expression patterns of nitrate, phosphate, and sulfate transporters in <i>Arabidopsis</i> roots exposed to different nutritional regimes. Botany, 2011, 89, 647-653.	1.0	30
48	Progress on trichome development regulated by phytohormone signaling. Plant Signaling and Behavior, 2011, 6, 1959-1962.	2.4	41
49	A DELLA in Disguise: SPATULA Restrains the Growth of the Developing <i>Arabidopsis</i> Seedling. Plant Cell, 2011, 23, 1337-1351.	6.6	77
50	SPATULA Links Daytime Temperature and Plant Growth Rate. Current Biology, 2010, 20, 1493-1497.	3.9	47
51	The effects of fluctuations in the nutrient supply on the expression of <i>ANR1</i> and 11 other MADS box genes in shoots and roots of <i>Arabidopsis thaliana</i> . Botany, 2010, 88, 1023-1031.	1.0	16
52	Genetic and Molecular Regulation by DELLA Proteins of Trichome Development in <i>Arabidopsis</i> . Plant Physiology, 2007, 145, 1031-1042.	4.8	35
53	Integration of cytokinin and gibberellin signalling by <i>Arabidopsis</i> transcription factors GIS, ZFP8 and GIS2 in the regulation of epidermal cell fate. Development (Cambridge), 2007, 134, 2073-2081.	2.5	178
54	GLABROUS INFLORESCENCE STEMS Modulates the Regulation by Gibberellins of Epidermal Differentiation and Shoot Maturation in <i>Arabidopsis</i> . Plant Cell, 2006, 18, 1383-1395.	6.6	134

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55	Nutritional regulation of ANR1 and other root-expressed MADS-box genes in <i>Arabidopsis thaliana</i> . <i>Planta</i> , 2005, 222, 730-742.	3.2	148