

Gang Li

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

2,018
citations

430874

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501196

28
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docs citations

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times ranked

2650
citing authors

#	ARTICLE	IF	CITATIONS
1	Chromatin and regulatory differentiation between bundle sheath and mesophyll cells in maize. <i>Plant Journal</i> , 2022, 109, 675-692.	5.7	16
2	Dynamic epigenetic modifications in plant sugar signal transduction. <i>Trends in Plant Science</i> , 2022, 27, 379-390.	8.8	24
3	Characterization of regulatory modules controlling leaf angle in maize. <i>Plant Physiology</i> , 2022, 190, 500-515.	4.8	10
4	Molecular and functional dissection of EARLY-FLOWERING 3 (ELF3) and ELF4 in Arabidopsis. <i>Plant Science</i> , 2021, 303, 110786.	3.6	22
5	Arabidopsis FAR-RED ELONGATED HYPOCOTYL3 negatively regulates carbon starvation responses. <i>Plant, Cell and Environment</i> , 2021, 44, 1816-1829.	5.7	11
6	The transcription factor PagLBD3 contributes to the regulation of secondary growth in <i>Populus</i> . <i>Journal of Experimental Botany</i> , 2021, 72, 7092-7106.	4.8	10
7	Light and Abscisic Acid Coordinately Regulate Greening of Seedlings. <i>Plant Physiology</i> , 2020, 183, 1281-1294.	4.8	18
8	Transcription Factors FHY3 and FAR1 Regulate Light-Induced <i>CIRCADIAN CLOCK ASSOCIATED1</i> Gene Expression in Arabidopsis. <i>Plant Cell</i> , 2020, 32, 1464-1478.	6.6	50
9	Arabidopsis FAR-RED ELONGATED HYPOCOTYL3 Integrates Age and Light Signals to Negatively Regulate Leaf Senescence. <i>Plant Cell</i> , 2020, 32, 1574-1588.	6.6	58
10	Arabidopsis NUCLEAR FACTOR Y A8 inhibits the juvenile-to-adult transition by activating transcription of <i>MIR156s</i> . <i>Journal of Experimental Botany</i> , 2020, 71, 4890-4902.	4.8	23
11	Heterologous expression of ELF4 from <i>Chlamydomonas reinhardtii</i> and <i>Physcomitrella patens</i> delays flowering in Arabidopsis thaliana. <i>Plant Systematics and Evolution</i> , 2019, 305, 777-785.	0.9	1
12	<i>FAR-RED ELONGATED HYPOCOTYL3</i> negatively regulates shade avoidance responses in Arabidopsis. <i>Plant, Cell and Environment</i> , 2019, 42, 3280-3292.	5.7	11
13	Auxin-Dependent Cell Elongation During the Shade Avoidance Response. <i>Frontiers in Plant Science</i> , 2019, 10, 914.	3.6	53
14	Molecular mechanisms governing shade responses in maize. <i>Biochemical and Biophysical Research Communications</i> , 2019, 516, 112-119.	2.1	22
15	WRKY18 and WRKY53 Coordinate with HISTONE ACETYLTRANSFERASE1 to Regulate Rapid Responses to Sugar. <i>Plant Physiology</i> , 2019, 180, 2212-2226.	4.8	54
16	Arabidopsis ELF4-like proteins EFL1 and EFL3 influence flowering time. <i>Gene</i> , 2019, 700, 131-138.	2.2	15
17	Genome-wide analysis of the basic Helix-Loop-Helix (bHLH) transcription factor family in maize. <i>BMC Plant Biology</i> , 2018, 18, 235.	3.6	102
18	FAR1-RELATED SEQUENCE (FRS) and FRS-RELATED FACTOR (FRF) Family Proteins in Arabidopsis Growth and Development. <i>Frontiers in Plant Science</i> , 2018, 9, 692.	3.6	130

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19	Regulation of Leaf Angle by Auricle Development in Maize. <i>Molecular Plant</i> , 2017, 10, 516-519.	8.3	33
20	<i>Arabidopsis thaliana</i> FAR-RED ELONGATED HYPOCOTYLS3 (FHY3) and FAR-RED-IMPAIRED RESPONSE1 (FAR1) modulate starch synthesis in response to light and sugar. <i>New Phytologist</i> , 2017, 213, 1682-1696.	7.3	49
21	Functional Characterization of the Maize Phytochrome-Interacting Factors PIF4 and PIF5. <i>Frontiers in Plant Science</i> , 2017, 8, 2273.	3.6	46
22	<i>Arabidopsis</i> FHY3 and FAR1 Regulate Light-Induced myo -Inositol Biosynthesis and Oxidative Stress Responses by Transcriptional Activation of MIPS1. <i>Molecular Plant</i> , 2016, 9, 541-557.	8.3	81
23	The <i>Arabidopsis thaliana</i> Nuclear Factor Y Transcription Factors. <i>Frontiers in Plant Science</i> , 2016, 07, 2045.	3.6	158
24	<i>Arabidopsis</i> FHY3 and HY5 Positively Mediate Induction of <i>COP1</i> Transcription in Response to Photomorphogenic UV-B Light. <i>Plant Cell</i> , 2012, 24, 4590-4606.	6.6	157
25	Phytochrome Signaling Mechanisms. <i>The Arabidopsis Book</i> , 2011, 9, e0148.	0.5	336
26	Coordinated transcriptional regulation underlying the circadian clock in <i>Arabidopsis</i> . <i>Nature Cell Biology</i> , 2011, 13, 616-622.	10.3	245
27	Genome-Wide Binding Site Analysis of FAR-RED ELONGATED HYPOCOTYL3 Reveals Its Novel Function in <i>Arabidopsis</i> Development. <i>Plant Cell</i> , 2011, 23, 2514-2535.	6.6	118
28	<i>Arabidopsis</i> Transcription Factor ELONGATED HYPOCOTYL5 Plays a Role in the Feedback Regulation of Phytochrome A Signaling. <i>Plant Cell</i> , 2010, 22, 3634-3649.	6.6	165