

Genji Qin

List of Publications by Year in descending order

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34
papers

2,662
citations

236925

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times ranked

4443
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#	ARTICLE	IF	CITATIONS
1	Arabidopsis transcription factor TCP4 represses chlorophyll biosynthesis to prevent petal greening. <i>Plant Communications</i> , 2022, 3, 100309.	7.7	16
2	MicroRNA775 regulates intrinsic leaf size and reduces cell wall pectin levels by targeting a galactosyltransferase gene in Arabidopsis. <i>Plant Cell</i> , 2021, 33, 581-602.	6.6	22
3	TCP transcription factors suppress cotyledon trichomes by impeding a cell differentiation-regulating complex. <i>Plant Physiology</i> , 2021, 186, 434-451.	4.8	20
4	The Regulation of CIN-like TCP Transcription Factors. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4498.	4.1	35
5	Arabidopsis Transcription Factor TCP5 Controls Plant Thermomorphogenesis by Positively Regulating PIF4 Activity. <i>IScience</i> , 2019, 15, 611-622.	4.1	82
6	The Transcription Factors TCP4 and PIF3 Antagonistically Regulate Organ-Specific Light Induction of SAUR Genes to Modulate Cotyledon Opening during De-Etiolation in Arabidopsis. <i>Plant Cell</i> , 2019, 31, 1155-1170.	6.6	74
7	The SWI/SNF subunit SWI3B regulates IAMT1 expression via chromatin remodeling in Arabidopsis leaf development. <i>Plant Science</i> , 2018, 271, 127-132.	3.6	10
8	TANDEM ZINC-FINGER/PLUS3 Is a Key Component of Phytochrome A Signaling. <i>Plant Cell</i> , 2018, 30, 835-852.	6.6	49
9	The TIE1 transcriptional repressor controls shoot branching by directly repressing BRANCHED1 in Arabidopsis. <i>PLoS Genetics</i> , 2018, 14, e1007296.	3.5	33
10	The Arabidopsis USL1 controls multiple aspects of development by affecting late endosome morphology. <i>New Phytologist</i> , 2018, 219, 1388-1405.	7.3	7
11	The Arabidopsis RING-Type E3 Ligase TEAR1 Controls Leaf Development by Targeting the TIE1 Transcriptional Repressor for Degradation. <i>Plant Cell</i> , 2017, 29, 243-259.	6.6	33
12	A Novel Imprinted Gene NUWA Controls Mitochondrial Function in Early Seed Development in Arabidopsis. <i>PLoS Genetics</i> , 2017, 13, e1006553.	3.5	40
13	EXB1/WRKY71 transcription factor regulates both shoot branching and responses to abiotic stresses. <i>Plant Signaling and Behavior</i> , 2016, 11, e1150404.	2.4	26
14	CFLAP1 and CFLAP2 Are Two bHLH Transcription Factors Participating in Synergistic Regulation of AtCFL1-Mediated Cuticle Development in Arabidopsis. <i>PLoS Genetics</i> , 2016, 12, e1005744.	3.5	22
15	The alteration in the architecture of a Tâ€DNA insertion rice mutant osmtd1 is caused by upâ€regulation of MicroRNA156f. <i>Journal of Integrative Plant Biology</i> , 2015, 57, 819-829.	8.5	26
16	The WRKY Transcription Factor WRKY71/EXB1 Controls Shoot Branching by Transcriptionally Regulating RAX Genes in Arabidopsis. <i>Plant Cell</i> , 2015, 27, 3112-3127.	6.6	102
17	The molecular mechanism of SPOROCTELESS/NOZZLE in controlling Arabidopsis ovule development. <i>Cell Research</i> , 2015, 25, 121-134.	12.0	93
18	ADP1 Affects Plant Architecture by Regulating Local Auxin Biosynthesis. <i>PLoS Genetics</i> , 2014, 10, e1003954.	3.5	47

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19	The Arabidopsis Mediator subunit MED16 regulates iron homeostasis by associating with EIN3/EIL1 through subunit MED25. <i>Plant Journal</i> , 2014, 77, 838-851.	5.7	120
20	AtMYB14 Regulates Cold Tolerance in Arabidopsis. <i>Plant Molecular Biology Reporter</i> , 2013, 31, 87-97.	1.8	102
21	The TIE1 Transcriptional Repressor Links TCP Transcription Factors with TOPLESS/TOPLESS-RELATED Corepressors and Modulates Leaf Development in Arabidopsis. <i>Plant Cell</i> , 2013, 25, 421-437.	6.6	116
22	Transcriptional Profiling of Rice Early Response to Magnaporthe oryzae Identified OsWRKYs as Important Regulators in Rice Blast Resistance. <i>PLoS ONE</i> , 2013, 8, e59720.	2.5	84
23	Arabidopsis RAP2.2 plays an important role in plant resistance to Botrytis cinerea and ethylene responses. <i>New Phytologist</i> , 2012, 195, 450-460.	7.3	129
24	Arabidopsis AtVPS15 Plays Essential Roles in Pollen Germination Possibly by Interacting with AtVPS34. <i>Journal of Genetics and Genomics</i> , 2012, 39, 81-92.	3.9	29
25	CFL1, a WW Domain Protein, Regulates Cuticle Development by Modulating the Function of HDG1, a Class IV Homeodomain Transcription Factor, in Rice and Arabidopsis. <i>Plant Cell</i> , 2011, 23, 3392-3411.	6.6	148
26	A nuclear-encoded mitochondrial gene AtCIB22 is essential for plant development in Arabidopsis. <i>Journal of Genetics and Genomics</i> , 2010, 37, 667-683.	3.9	15
27	Dof5.6/HCA2, a Dof Transcription Factor Gene, Regulates Interfascicular Cambium Formation and Vascular Tissue Development in Arabidopsis. <i>Plant Cell</i> , 2009, 21, 3518-3534.	6.6	162
28	Targeted Degradation of the Cyclin-Dependent Kinase Inhibitor ICK4/KRP6 by RING-Type E3 Ligases Is Essential for Mitotic Cell Cycle Progression during Arabidopsis Gametogenesis. <i>Plant Cell</i> , 2008, 20, 1538-1554.	6.6	142
29	NPY1, a BTB-NPH3-like protein, plays a critical role in auxin-regulated organogenesis in Arabidopsis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 18825-18829.	7.1	125
30	Disruption of phytoene desaturase gene results in albino and dwarf phenotypes in Arabidopsis by impairing chlorophyll, carotenoid, and gibberellin biosynthesis. <i>Cell Research</i> , 2007, 17, 471-482.	12.0	313
31	Arabidopsis AtBECLIN 1/AtAtg6/AtVps30 is essential for pollen germination and plant development. <i>Cell Research</i> , 2007, 17, 249-263.	12.0	107
32	GAMT2 Encodes a Methyltransferase of Gibberellic Acid That is Involved in Seed Maturation and Germination in Arabidopsis. <i>Journal of Integrative Plant Biology</i> , 2007, 49, 368-381.	8.5	14
33	An Indole-3-Acetic Acid Carboxyl Methyltransferase Regulates Arabidopsis Leaf Development. <i>Plant Cell</i> , 2005, 17, 2693-2704.	6.6	260
34	Obtaining and analysis of flanking sequences from T-DNA transformants of Arabidopsis. <i>Plant Science</i> , 2003, 165, 941-949.	3.6	54