

Tai-Ping Sun

List of Publications by Year in descending order

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papers

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41344

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times ranked

9410
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel nucleocytoplasmic protein O-fucosylation by SPINDLY regulates diverse developmental processes in plants. <i>Current Opinion in Structural Biology</i> , 2021, 68, 113-121.	5.7	26
2	Nuclear Localized O-Fucosyltransferase SPY Facilitates PRR5 Proteolysis to Fine-Tune the Pace of Arabidopsis Circadian Clock. <i>Molecular Plant</i> , 2020, 13, 446-458.	8.3	41
3	Plasmonic Nanobiosensing: from in situ plant monitoring to cancer diagnostics at the point of care. <i>JPhys Photonics</i> , 2020, 2, 034012.	4.6	3
4	Plasmonic Nanoprobles for in Vivo Multimodal Sensing and Bioimaging of MicroRNA within Plants. <i>ACS Applied Materials & Interfaces</i> , 2019, 11, 7743-7754.	8.0	42
5	NCP activates chloroplast transcription by controlling phytochrome-dependent dual nuclear and plastidial switches. <i>Nature Communications</i> , 2019, 10, 2630.	12.8	38
6	Inverse Molecular Sentinel-Integrated Fiberoptic Sensor for Direct and <i>in Situ</i> Detection of miRNA Targets. <i>Analytical Chemistry</i> , 2019, 91, 6345-6352.	6.5	31
7	The Interaction between DELLA and ARF/IAA Mediates Crosstalk between Gibberellin and Auxin Signaling to Control Fruit Initiation in Tomato. <i>Plant Cell</i> , 2018, 30, 1710-1728.	6.6	129
8	Identification and functional study of a mild allele of SIDEELLA gene conferring the potential for improved yield in tomato. <i>Scientific Reports</i> , 2018, 8, 12043.	3.3	37
9	Gibberellin Signaling Requires Chromatin Remodeler PICKLE to Promote Vegetative Growth and Phase Transitions. <i>Plant Physiology</i> , 2017, 173, 1463-1474.	4.8	55
10	Structure of the SHR-SCR heterodimer bound to the BIRD/IDD transcriptional factor JKD. <i>Nature Plants</i> , 2017, 3, 17010.	9.3	65
11	The Arabidopsis O-fucosyltransferase SPINDLY activates nuclear growth repressor DELLA. <i>Nature Chemical Biology</i> , 2017, 13, 479-485.	8.0	130
12	The ERF11 Transcription Factor Promotes Internode Elongation by Activating Gibberellin Biosynthesis and Signaling. <i>Plant Physiology</i> , 2016, 171, 2760-2770.	4.8	80
13	O-GlcNAcylation of master growth repressor DELLA by SECRET AGENT modulates multiple signaling pathways in Arabidopsis. <i>Genes and Development</i> , 2016, 30, 164-176.	5.9	101
14	Functional characterization and developmental expression profiling of gibberellin signalling components in Vitis vinifera. <i>Journal of Experimental Botany</i> , 2015, 66, 1463-1476.	4.8	36
15	Sex and the single fern. <i>Science</i> , 2014, 346, 423-424.	12.6	2
16	Role of the gibberellin receptors GID1 during fruit set in Arabidopsis. <i>Plant Journal</i> , 2014, 79, 1020-1032.	5.7	68
17	Leaf-Induced Gibberellin Signaling Is Essential for Internode Elongation, Cambial Activity, and Fiber Differentiation in Tobacco Stems. <i>Plant Cell</i> , 2012, 24, 66-79.	6.6	117
18	Brassinosteroid, gibberellin and phytochrome impinge on a common transcription module in Arabidopsis. <i>Nature Cell Biology</i> , 2012, 14, 810-817.	10.3	549

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19	Hypocotyl Transcriptome Reveals Auxin Regulation of Growth-Promoting Genes through GA-Dependent and -Independent Pathways. <i>PLoS ONE</i> , 2012, 7, e36210.	2.5	127
20	Plant hormone jasmonate prioritizes defense over growth by interfering with gibberellin signaling cascade. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E1192-200.	7.1	697
21	Rapid and orthogonal logic gating with a gibberellin-induced dimerization system. <i>Nature Chemical Biology</i> , 2012, 8, 465-470.	8.0	183
22	The Molecular Mechanism and Evolution of the GA-“GID1”-DELLA Signaling Module in Plants. <i>Current Biology</i> , 2011, 21, R338-R345.	3.9	464
23	SCARECROW-LIKE 3 promotes gibberellin signaling by antagonizing master growth repressor DELLA in <i>Arabidopsis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 2160-2165.	7.1	302
24	Gibberellin-GID1-DELLA: A Pivotal Regulatory Module for Plant Growth and Development. <i>Plant Physiology</i> , 2010, 154, 567-570.	4.8	314
25	Isolation and Characterization of <i>cul1-7</i> , a Recessive Allele of <i>CULLIN1</i> That Disrupts SCF Function at the C Terminus of CUL1 in <i>Arabidopsis thaliana</i> . <i>Genetics</i> , 2009, 181, 945-963.	2.9	41
26	Gibberellin Perception and Signalling. <i>Seibutsu Butsuri</i> , 2009, 49, 200-201.	0.1	0
27	Gibberellin-induced DELLA recognition by the gibberellin receptor GID1. <i>Nature</i> , 2008, 456, 459-463.	27.8	594
28	Proteolysis-Independent Downregulation of DELLA Repression in <i>Arabidopsis</i> by the Gibberellin Receptor GIBBERELLIN INSENSITIVE DWARF1. <i>Plant Cell</i> , 2008, 20, 2447-2459.	6.6	144
29	Potential Sites of Bioactive Gibberellin Production during Reproductive Growth in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2008, 20, 320-336.	6.6	209
30	Gibberellin Metabolism, Perception and Signaling Pathways in <i>Arabidopsis</i> . <i>The Arabidopsis Book</i> , 2008, 6, e0103.	0.5	207
31	Global Analysis of DELLA Direct Targets in Early Gibberellin Signaling in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2007, 19, 3037-3057.	6.6	572
32	PIL5, a Phytochrome-Interacting bHLH Protein, Regulates Gibberellin Responsiveness by Binding Directly to the GAI and RGA Promoters in <i>Arabidopsis</i> Seeds. <i>Plant Cell</i> , 2007, 19, 1192-1208.	6.6	405
33	Functional Analysis of SPINDLY in Gibberellin Signaling in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2007, 143, 987-1000.	4.8	146
34	Genetic Characterization and Functional Analysis of the GID1 Gibberellin Receptors in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2007, 18, 3399-3414.	6.6	665
35	Regulation of hormone metabolism in <i>Arabidopsis</i> seeds: phytochrome regulation of abscisic acid metabolism and abscisic acid regulation of gibberellin metabolism. <i>Plant Journal</i> , 2006, 48, 354-366.	5.7	403
36	Distinct and overlapping roles of two gibberellin 3-oxidases in <i>Arabidopsis</i> development. <i>Plant Journal</i> , 2006, 45, 804-818.	5.7	282

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37	A DELLAcate balance: the role of gibberellin in plant morphogenesis. <i>Current Opinion in Plant Biology</i> , 2005, 8, 77-85.	7.1	433
38	A Novel Dwarfing Mutation in a Green Revolution Gene from <i>Brassica rapa</i> . <i>Plant Physiology</i> , 2005, 137, 931-938.	4.8	77
39	Update on Gibberellin Signaling. A Tale of the Tall and the Short. <i>Plant Physiology</i> , 2004, 135, 668-676.	4.8	158
40	Arabidopsis CAND1, an Unmodified CUL1-Interacting Protein, Is Involved in Multiple Developmental Pathways Controlled by Ubiquitin/Proteasome-Mediated Protein Degradation. <i>Plant Cell</i> , 2004, 16, 1870-1882.	6.6	135
41	The Arabidopsis F-Box Protein SLEEPY1 Targets Gibberellin Signaling Repressors for Gibberellin-Induced Degradation[W]. <i>Plant Cell</i> , 2004, 16, 1392-1405.	6.6	523
42	DELLA Proteins and Gibberellin-Regulated Seed Germination and Floral Development in Arabidopsis. <i>Plant Physiology</i> , 2004, 135, 1008-1019.	4.8	521
43	MOLECULAR MECHANISM OF GIBBERELLIN SIGNALING IN PLANTS. <i>Annual Review of Plant Biology</i> , 2004, 55, 197-223.	18.7	629
44	The ArabidopsisSLEEPY1Gene Encodes a Putative F-Box Subunit of an SCF E3 Ubiquitin Ligase[W]. <i>Plant Cell</i> , 2003, 15, 1120-1130.	6.6	505
45	Overexpression of AtCPS and AtKS in Arabidopsis Confers Increased ent-Kaurene Production But No Increase in Bioactive Gibberellins. <i>Plant Physiology</i> , 2003, 132, 830-839.	4.8	119
46	Gibberellin Signaling. <i>Plant Cell</i> , 2002, 14, S61-S80.	6.6	870
47	Distinct cell-specific expression patterns of early and late gibberellin biosynthetic genes during Arabidopsis seed germination. <i>Plant Journal</i> , 2002, 28, 443-453.	5.7	156
48	Characterization of cis-regulatory regions responsible for developmental regulation of the gibberellin biosynthetic gene GA1 in Arabidopsis thaliana. <i>Plant Molecular Biology</i> , 2002, 49, 579-589.	3.9	15
49	Repressing a Repressor. <i>Plant Cell</i> , 2001, 13, 1555-1566.	6.6	412
50	Synergistic Derepression of Gibberellin Signaling by Removing RGA and GAI Function in <i>Arabidopsis thaliana</i> . <i>Genetics</i> , 2001, 159, 777-785.	2.9	399
51	Gibberellin signal transduction. <i>Current Opinion in Plant Biology</i> , 2000, 3, 374-380.	7.1	97
52	Gibberellins and the Green Revolution. <i>Trends in Plant Science</i> , 2000, 5, 1-2.	8.8	100
53	The Arabidopsis RGA Gene Encodes a Transcriptional Regulator Repressing the Gibberellin Signal Transduction Pathway. <i>Plant Cell</i> , 1998, 10, 155-169.	6.6	699
54	Phytochrome Regulation and Differential Expression of Gibberellin 3 ^β -Hydroxylase Genes in Germinating Arabidopsis Seeds. <i>Plant Cell</i> , 1998, 10, 2115-2126.	6.6	330

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55	The GA2 Locus of <i>Arabidopsis thaliana</i> Encodes ent-Kaurene Synthase of Gibberellin Biosynthesis. <i>Plant Physiology</i> , 1998, 116, 1271-1278.	4.8	197
56	Phytochrome Regulation and Differential Expression of Gibberellin 3 β -Hydroxylase Genes in Germinating <i>Arabidopsis</i> Seeds. <i>Plant Cell</i> , 1998, 10, 2115.	6.6	185
57	The <i>Arabidopsis</i> RGA Gene Encodes a Transcriptional Regulator Repressing the Gibberellin Signal Transduction Pathway. <i>Plant Cell</i> , 1998, 10, 155.	6.6	43
58	Regulation and cellular localization of ent-kaurene synthesis. <i>Physiologia Plantarum</i> , 1997, 101, 701-708.	5.2	44
59	The LS locus of pea encodes the gibberellin biosynthesis enzyme ent-kaurene synthase A. <i>Plant Journal</i> , 1997, 11, 443-454.	5.7	104
60	Developmental regulation of the gibberellin biosynthetic gene GA1 in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 1997, 12, 9-19.	5.7	210
61	Regulation and cellular localization of ent-kaurene synthesis. <i>Physiologia Plantarum</i> , 1997, 101, 701-708.	5.2	6
62	The New <i>RGA</i> Locus Encodes a Negative Regulator of Gibberellin Response in <i>Arabidopsis thaliana</i> . <i>Genetics</i> , 1997, 146, 1087-1099.	2.9	262
63	The <i>Arabidopsis</i> GA1 Locus Encodes the Cyclase ent-Kaurene Synthetase A of Gibberellin Biosynthesis. <i>Plant Cell</i> , 1994, 6, 1509.	6.6	102
64	Cloning the <i>Arabidopsis</i> GA1 Locus by Genomic Subtraction. <i>Plant Cell</i> , 1992, 4, 119.	6.6	94
65	Cloning <i>Arabidopsis</i> genes by genomic subtraction. , 1992, , 331-341.		1