Xing Wang Deng

List of Publications by Year
in descending order

127 papers	13,847 citations	20817 60 h-index	22166 113 g-index
217 all docs	217 docs citations	217 times ranked	10189 citing authors

#	Article	IF	Citations
1	Targeted destabilization of HY5 during light-regulated development of Arabidopsis. Nature, 2000, 405, 462-466.	27.8	1,227
2	Light-regulated transcriptional networks in higher plants. Nature Reviews Genetics, 2007, 8, 217-230.	16.3	892
3	Analysis of Transcription Factor HY5 Genomic Binding Sites Revealed Its Hierarchical Role in Light Regulation of Development. Plant Cell, 2007, 19, 731-749.	6.6	829
4	The photomorphogenic repressors COP1 and DET1: 20 years later. Trends in Plant Science, 2012, 17, 584-593.	8.8	530
5	Light Control of Arabidopsis Development Entails Coordinated Regulation of Genome Expression and Cellular Pathways. Plant Cell, 2001, 13, 2589-2607.	6.6	498
6	Direct Interaction of Arabidopsis Cryptochromes with COP1 in Light Control Development. Science, 2001, 294, 154-158.	12.6	473
7	The COP1-SPA1 interaction defines a critical step in phytochrome A-mediated regulation of HY5 activity. Genes and Development, 2003, 17, 2642-2647.	5.9	403
8	<i>Arabidopsis</i> noncoding RNA mediates control of photomorphogenesis by red light. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 10359-10364.	7.1	317
9	COP1 â€“ from plant photomorphogenesis to mammalian tumorigenesis. Trends in Cell Biology, 2005, 15, 618-625.	7.9	302
10	Genome-Wide Analysis of DNA Methylation and Gene Expression Changes in Two <i>Arabidopsis</i> Ecotypes and Their Reciprocal Hybrids. Plant Cell, 2012, 24, 875-892.	6.6	297
11	A High-Density SNP Genotyping Array for Rice Biology and Molecular Breeding. Molecular Plant, 2014, 7, 541-553.	8.3	251
12	Coordinated transcriptional regulation underlying the circadian clock in Arabidopsis. Nature Cell Biology, 2011, 13, 616-622.	10.3	245
13	From seed to seed: the role of photoreceptors in Arabidopsis development. Developmental Biology, 2003, 260, 289-297.	2.0	214
14	BBX21, an <i>Arabidopsis</i> B-box protein, directly activates <i>HY5</i> and is targeted by COP1 for 26S proteasome-mediated degradation. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 7655-7660.	7.1	204
15	Biochemical Characterization of<i>Arabidopsis</i>Complexes Containing CONSTITUTIVELY PHOTOMORPHOGENIC1 and SUPPRESSOR OF PHYA Proteins in Light Control of Plant Development. Plant Cell, 2008, 20, 2307-2323.	6.6	202
16	Arabidopsis COP10 forms a complex with DDB1 and DET1 in vivo and enhances the activity of ubiquitin conjugating enzymes. Genes and Development, 2004, 18, 2172-2181.	5.9	186
17	Targeted Degradation of Abscisic Acid Receptors Is Mediated by the Ubiquitin Ligase Substrate Adaptor DDA1 in <i>Arabidopsis</i>. Plant Cell, 2014, 26, 712-728.	6.6	186
18	<i>Arabidopsis</i> CULLIN4-Damaged DNA Binding Protein 1 Interacts with CONSTITUTIVELY PHOTOMORPHOGENIC1-SUPPRESSOR OF PHYA Complexes to Regulate Photomorphogenesis and Flowering Time Â. Plant Cell, 2010, 22, 108-123.	6.6	182

#	Article	IF	Citations
19	The Epigenome and Plant Development. Annual Review of Plant Biology, 2011, 62, 411-435.	18.7	172
20	DELLA-mediated PIF degradation contributes to coordination of light and gibberellin signalling in Arabidopsis. Nature Communications, 2016, 7, 11868.	12.8	172
21	<i>Arabidopsis</i> Transcription Factor ELONGATED HYPOCOTYL5 Plays a Role in the Feedback Regulation of Phytochrome A Signaling Â. Plant Cell, 2010, 22, 3634-3649.	6.6	165
22	MicroRNA408 Is Critical for the <i>HY5-SPL7</i> Gene Network That Mediates the Coordinated Response to Light and Copper Â. Plant Cell, 2015, 26, 4933-4953.	6.6	164
23	A high-quality genome assembly highlights rye genomic characteristics and agronomically important genes. Nature Genetics, 2021, 53, 574-584.	21.4	164
24	Conversion from CUL4-based COP1â€"SPA E3 apparatus to UVR8â€"COP1â€"SPA complexes underlies a distinct biochemical function of COP1 under UV-B. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 16669-16674.	7.1	163
25	Convergence of Light and ABA Signaling on the ABI5 Promoter. PLoS Genetics, 2014, 10, e1004197.	3.5	163
26	ArabidopsisFHY3 defines a key phytochrome A signaling component directly interacting with its homologous partner FAR1. EMBO Journal, 2002, 21, 1339-1349.	7.8	141
27	Beyond repression of photomorphogenesis: role switching of COP/DET/FUS in light signaling. Current Opinion in Plant Biology, 2014, 21, 96-103.	7.1	141
28	HFR1 Sequesters PIF1 to Govern the Transcriptional Network Underlying Light-Initiated Seed Germination in <i>Arabidopsis</i> Â Â Â. Plant Cell, 2013, 25, 3770-3784.	6.6	128
29	<i>Arabidopsis</i> SAURs are critical for differential light regulation of the development of various organs. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6071-6076.	7.1	127
30	Rare earth elements activate endocytosis in plant cells. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 12936-12941.	7.1	120
31	Seedlings Transduce the Depth and Mechanical Pressure of Covering Soil Using COP1 and Ethylene to Regulate EBF1/EBF2 for Soil Emergence. Current Biology, 2016, 26, 139-149.	3.9	120
32	Genome-Wide Binding Site Analysis of FAR-RED ELONGATED HYPOCOTYL3 Reveals Its Novel Function in <i>Arabidopsis</i> Development. Plant Cell, 2011, 23, 2514-2535.	6.6	118
33	Conservation and divergence of transcriptomic and epigenomic variation in maize hybrids. Genome Biology, 2013, 14, R57.	8.8	117
34	The Roles of Photoreceptor Systems and the COP1-Targeted Destabilization of HY5 in Light Control of Arabidopsis Seedling Development. Plant Physiology, 2000, 124, 1520-1524.	4.8	116
35	Arabidopsis DE-ETIOLATED1 Represses Photomorphogenesis by Positively Regulating Phytochrome-Interacting Factors in the Dark. Plant Cell, 2014, 26, 3630-3645.	6.6	116
36	Noncanonical role of <i>Arabidopsis</i> COP1/SPA complex in repressing BIN2-mediated PIF3 phosphorylation and degradation in darkness. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 3539-3544.	7.1	109

#	ARTICLE	IF	CITATIONS
37	B-BOX DOMAIN PROTEIN28 Negatively Regulates Photomorphogenesis by Repressing the Activity of Transcription Factor HY5 and Undergoes COP1-Mediated Degradation. Plant Cell, 2018, 30, 2006-2019.	6.6	105
38	Arabidopsis COP1/SPA1 Complex and FHY1/FHY3 Associate with Distinct Phosphorylated Forms of Phytochrome A in Balancing Light Signaling. Molecular Cell, 2008, 31, 607-613.	9.7	104
39	A Genome-Wide Transcription Analysis Reveals a Close Correlation of Promoter INDEL Polymorphism and Heterotic Gene Expression in Rice Hybrids. Molecular Plant, 2008, 1, 720-731.	8.3	101
40	Basic leucine zipper transcription factor OsbZIP16 positively regulates drought resistance in rice. Plant Science, 2012, 193-194, 8-17.	3.6	98
41	The PP6 Phosphatase Regulates ABI5 Phosphorylation and Abscisic Acid Signaling in<i>Arabidopsis</i>Â Â. Plant Cell, 2013, 25, 517-534.	6.6	98
42	CUL4 forms an E3 ligase with COP1 and SPA to promote light-induced degradation of PIF1. Nature Communications, 2015, 6, 7245.	12.8	97
43	Light-Dependent Degradation of PIF3 by SCFEBF1/2 Promotes a Photomorphogenic Response in Arabidopsis. Current Biology, 2017, 27, 2420-2430.e6.	3.9	95
44	The Photomorphogenic Central Repressor COP1: Conservation and Functional Diversification during Evolution. Plant Communications, 2020, 1, 100044.	7.7	95
45	Ethylene Promotes Hypocotyl Growth and HY5 Degradation by Enhancing the Movement of COP1 to the Nucleus in the Light. PLoS Genetics, 2013, 9, e1004025.	3.5	93
46	Salicylic acid biosynthesis is enhanced and contributes to increased biotrophic pathogen resistance in Arabidopsis hybrids. Nature Communications, 2015, 6, 7309.	12.8	93
47	The Red Light Receptor Phytochrome B Directly Enhances Substrate-E3 Ligase Interactions to Attenuate Ethylene Responses. Developmental Cell, 2016, 39, 597-610.	7.0	91
48	Photoactivated UVR8-COP1 Module Determines Photomorphogenic UV-B Signaling Output in Arabidopsis. PLoS Genetics, 2014, 10, e1004218.	3.5	88
49	Overexpression of the Heterotrimeric G-Protein Î±-Subunit Enhances Phytochrome-Mediated Inhibition of Hypocotyl Elongation in Arabidopsis. Plant Cell, 2001, 13, 1639-1652.	6.6	85
50	Origin and Evolution of Core Components Responsible for Monitoring Light Environment Changes during Plant Terrestrialization. Molecular Plant, 2019, 12, 847-862.	8.3	85
51	A PP6-Type Phosphatase Holoenzyme Directly Regulates PIN Phosphorylation and Auxin Efflux in <i>Arabidopsis</i>. Plant Cell, 2012, 24, 2497-2514.	6.6	84
52	Poaceae-specific<i>MS1</i>encodes a phospholipid-binding protein for male fertility in bread wheat. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 12614-12619.	7.1	83
53	PHYTOCHROME INTERACTING FACTOR1 Enhances the E3 Ligase Activity of CONSTITUTIVE PHOTOMORPHOGENIC1 to Synergistically Repress Photomorphogenesis in <i>Arabidopsis</i> Â Â. Plant Cell, 2014, 26, 1992-2006.	6.6	78
54	The B-Box Domain Protein BBX21 Promotes Photomorphogenesis. Plant Physiology, 2018, 176, 2365-2375.	4.8	78

#	Article	IF	Citations
55	The Transcription Factors TCP4 and PIF3 Antagonistically Regulate Organ-Specific Light Induction of <i>SAUR</i> Genes to Modulate Cotyledon Opening during De-Etiolation in Arabidopsis. Plant Cell, 2019, 31, 1155-1170.	6.6	74
56	<i>Arabidopsis</i> Phytochrome A Directly Targets Numerous Promoters for Individualized Modulation of Genes in a Wide Range of Pathways. Plant Cell, 2014, 26, 1949-1966.	6.6	73
57	Genomic architecture of biomass heterosis in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8101-8106.	7.1	73
58	<i>Arabidopsis</i> DET1 degrades HFR1 but stabilizes PIF1 to precisely regulate seed germination. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 3817-3822.	7.1	69
59	B-Box Containing Proteins BBX30 and BBX31, Acting Downstream of HY5, Negatively Regulate Photomorphogenesis in <i>Arabidopsis</i>. Plant Physiology, 2019, 180, 497-508.	4.8	69
60	Genome-wide regulation of light-controlled seedling morphogenesis by three families of transcription factors. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 6482-6487.	7.1	68
61	Salt Stress and Ethylene Antagonistically Regulate Nucleocytoplasmic Partitioning of COP1 to Control Seed Germination. Plant Physiology, 2016, 170, 2340-2350.	4.8	67
62	COP9 signalosome: Discovery, conservation, activity, and function. Journal of Integrative Plant Biology, 2020, 62, 90-103.	8.5	66
63	Two E3 ligases antagonistically regulate the UV-B response in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 4722-4731.	7.1	61
64	Modulation of BIN2 kinase activity by HY5 controls hypocotyl elongation in the light. Nature Communications, 2020, 11, 1592.	12.8	61
65	CRISPR/Cas9-mediated disruption of TaNP1 genes results in complete male sterility in bread wheat. Journal of Genetics and Genomics, 2020, 47, 263-272.	3.9	58
66	The COP9 Signalosome regulates seed germination by facilitating protein degradation of RGL2 and ABI5. PLoS Genetics, 2018, 14, e1007237.	3.5	55
67	Arabinogalactan proteinâ€“rare earth element complexes activate plant endocytosis. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 14349-14357.	7.1	52
68	HY5 regulates nitrite reductase 1 (NIR1) and ammonium transporter1;2 (AMT1;2) in Arabidopsis seedlings. Plant Science, 2015, 238, 330-339.	3.6	49
69	TANDEM ZINC-FINGER/PLUS3 Is a Key Component of Phytochrome A Signaling. Plant Cell, 2018, 30, 835-852.	6.6	49
70	Diurnal down-regulation of ethylene biosynthesis mediates biomass heterosis. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 5606-5611.	7.1	49
71	Mammalian DET1 Regulates Cul4A Activity and Forms Stable Complexes with E2 Ubiquitin-Conjugating Enzymes. Molecular and Cellular Biology, 2007, 27, 4708-4719.	2.3	46
72	SAUR17 and SAUR50 Differentially Regulate PP2C-D1 during Apical Hook Development and Cotyledon Opening in Arabidopsis. Plant Cell, 2020, 32, 3792-3811.	6.6	46

#	Article	IF	Citations
73	BBX28/BBX29, HY5 and BBX30/31 form a feedback loop to fineâ€tune photomorphogenic development. Plant Journal, 2020, 104, 377-390.	5.7	46
74	Divergent selection and genetic introgression shape the genome landscape of heterosis in hybrid rice. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 4623-4631.	7.1	46
75	Structural organization and interactions of COP1, a light-regulated developmental switch. , 1999, 41, 151-158.		43
76	The RING-Finger E3 Ubiquitin Ligase COP1 SUPPRESSOR1 Negatively Regulates COP1 Abundance in Maintaining COP1 Homeostasis in Dark-Grown <i>Arabidopsis</i> Seedlings Â Â. Plant Cell, 2014, 26, 1981-1991.	6.6	41
77	A Positive Feedback Loop of BBX11â€“BBX21â€“HY5 Promotes Photomorphogenic Development in Arabidopsis. Plant Communications, 2020, 1, 100045.	7.7	39
78	Pedigreeâ€based analysis of derivation of genome segments of an elite rice reveals key regions during its breeding. Plant Biotechnology Journal, 2016, 14, 638-648.	8.3	38
79	Biological pathway expression complementation contributes to biomass heterosis in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	38
80	The Asymmetric Expression of SAUR Genes Mediated by ARF7/19 Promotes the Gravitropism and Phototropism of Plant Hypocotyls. Cell Reports, 2020, 31, 107529.	6.4	35
81	BBX4, a phyB-interacting and modulated regulator, directly interacts with PIF3 to fine tune red light-mediated photomorphogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 26049-26056.	7.1	34
82	A new regulator of seed size control in <i>Arabidopsis</i> identified by a genomeâ€wide association study. New Phytologist, 2019, 222, 895-906.	7.3	34
83	Development of the â€œThird-Generationâ€• Hybrid Rice in China. Genomics, Proteomics and Bioinformatics, 2018, 16, 393-396.	6.9	33
84	ASG2 is a farnesylated DWD protein that acts as ABA negative regulator in <i>Arabidopsis</i>. Plant, Cell and Environment, 2016, 39, 185-198.	5.7	32
85	Improved de novo genome assembly and analysis of the Chinese cucurbit Siraitia grosvenorii, also known as monk fruit or luo-han-guo. GigaScience, 2018, 7, .	6.4	32
86	COLD-REGULATED GENE27 Integrates Signals from Light and the Circadian Clock to Promote Hypocotyl Growth in Arabidopsis. Plant Cell, 2020, 32, 3155-3169.	6.6	32
87	The telomereâ€toâ€telomere gapâ€free genome of four rice parents reveals <scp>SV</scp> and <scp>PAV</scp> patterns in hybrid rice breeding. Plant Biotechnology Journal, 2022, 20, 1642-1644.	8.3	31
88	Phytochrome B Induces Intron Retention and Translational Inhibition of PHYTOCHROME-INTERACTING FACTOR3. Plant Physiology, 2020, 182, 159-166.	4.8	29
89	Light modulates the gravitropic responses through organ-specific PIFs and HY5 regulation of <i>LAZY4</i> expression in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 18840-18848.	7.1	29
90	From hybrid genomes to heterotic trait output: Challenges and opportunities. Current Opinion in Plant Biology, 2022, 66, 102193.	7.1	29

#	Article	IF	Citations
91	Arabidopsis DET1 Represses Photomorphogenesis in Part by Negatively Regulating DELLA Protein Abundance in Darkness. Molecular Plant, 2015, 8, 622-630.	8.3	26
92	<i>Arabidopsis</i> small nucleolar RNA monitors the efficient pre-rRNA processing during ribosome biogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 11967-11972.	7.1	26
93	Red-light is an environmental effector for mutualism between begomovirus and its vector whitefly. PLoS Pathogens, 2021, 17, e1008770.	4.7	26
94	Arabidopsis COP1 SUPPRESSOR 2 Represses COP1 E3 Ubiquitin Ligase Activity through Their Coiled-Coil Domains Association. PLoS Genetics, 2015, 11, e1005747.	3.5	23
95	Phosphorylation and negative regulation of CONSTITUTIVELY PHOTOMORPHOGENIC 1 by PINOID <i>in Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 6617-6622.	7.1	23
96	Hinge region of <i>Arabidopsis</i> phyA plays an important role in regulating phyA function. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E11864-E11873.	7.1	22
97	CBF-phyB-PIF Module Links Light and Low Temperature Signaling. Trends in Plant Science, 2020, 25, 952-954.	8.8	22
98	The role of COP1 in repression of photoperiodic flowering. F1000Research, 2016, 5, 178.	1.6	22
99	Arabidopsis Atypical Kinases ABC1K1 and ABC1K3 Act Oppositely to Cope with Photodamage Under Red Light. Molecular Plant, 2015, 8, 1122-1124.	8.3	20
100	Characterization of a Novel DWD Protein that Participates in Heat Stress Response in Arabidopsis. Molecules and Cells, 2014, 37, 833-840.	2.6	18
101	Genome-wide dissection of heterosis for yield traits in two-line hybrid rice populations. Scientific Reports, 2017, 7, 7635.	3.3	18
102	Light and Abscisic Acid Coordinately Regulate Greening of Seedlings. Plant Physiology, 2020, 183, 1281-1294.	4.8	18
103	A central circadian oscillator confers defense heterosis in hybrids without growth vigor costs. Nature Communications, 2021, 12, 2317.	12.8	18
104	Natural variation of H3K27me3 modification in two <i>Arabidopsis</i> accessions and their hybrid. Journal of Integrative Plant Biology, 2016, 58, 466-474.	8.5	17
105	The PCY-SAG14 phytocyanin module regulated by PIFs and miR408 promotes dark-induced leaf senescence in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	17
106	The photomorphogenic repressors BBX28 and BBX29 integrate light and brassinosteroid signaling to inhibit seedling development in Arabidopsis. Plant Cell, 2022, 34, 2266-2285.	6.6	17
107	BBX11 promotes red light-mediated photomorphogenic development by modulating phyB-PIF4 signaling. ABIOTECH, 2021, 2, 117-130.	3.9	16
108	Photoreceptor partner FHY1 has an independent role in gene modulation and plant development under far-red light. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 11888-11893.	7.1	14

#	Article	IF	Citations
109	<i>Arabidopsis</i> PP6 phosphatases dephosphorylate PIF proteins to repress photomorphogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 20218-20225.	7.1	14
110	The <i>Arabidopsis</i> DREAM complex antagonizes WDR5A to modulate histone H3K4me2/3 deposition for a subset of genome repression. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	14
111	Multiple photomorphogenic repressors work in concert to regulate Arabidopsis seedling development. Plant Signaling and Behavior, 2015, 10, e1011934.	2.4	13
112	Genome-wide study of an elite rice pedigree reveals a complex history of genetic architecture for breeding improvement. Scientific Reports, 2017, 7, 45685.	3.3	13
113	Coordinated photomorphogenic UV-B signaling network captured by mathematical modeling. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 11539-11544.	7.1	12
114	COP1 SUPPRESSOR 4 promotes seedling photomorphogenesis by repressing <i>CCA1</i> and <i>PIF4</i> expression in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 11631-11636.	7.1	12
115	Structural insight into UV-Bâ€"activated UVR8 bound to COP1. Science Advances, 2022, 8, eabn3337.	10.3	12
116	SWELLMAP 2, a phyB-Interacting Splicing Factor, Negatively Regulates Seedling Photomorphogenesis in Arabidopsis. Frontiers in Plant Science, 2022, 13, 836519.	3.6	11
117	Arabidopsis atypical kinase ABC1K1 is involved in red light-mediated development. Plant Cell Reports, 2016, 35, 1213-1220.	5.6	9
118	Gibberellin Signal Transduction in Rice. Journal of Integrative Plant Biology, 2007, 49, 731-741.	8.5	8
119	A missense mutation in WRKY32 converts its function from a positive regulator to a repressor of photomorphogenesis. New Phytologist, 2021, , .	7.3	8
120	Genomic insights on the contribution of introgressions from Xian/Indica to the genetic improvement of Geng/Japonica rice cultivars. Plant Communications, 2022, 3, 100325.	7.7	8
121	Allele-specific DNA methylation analyses associated with siRNAs in Arabidopsis hybrids. Science China Life Sciences, 2014, 57, 519-525.	4.9	7
122	Natural variation in the transcription factor REPLUMLESS contributes to both disease resistance and plant growth in Arabidopsis. Plant Communications, 2022, 3, 100351.	7.7	4
123	Analysis of the Transcriptional Dynamics of Regulatory Genes During Peanut Pod Development Caused by Darkness and Mechanical Stress. Frontiers in Plant Science, 2022, 13, .	3.6	3
124	Single-Molecule Sequencing Assists Genome Assembly Improvement and Structural Variation Inference. Molecular Plant, 2016, 9, 1085-1087.	8.3	2
125	Organization of protein complexes under photomorphogenic UV-B inArabidopsis. Plant Signaling and Behavior, 2013, 8, e27206.	2.4	1
126	Exploring the genetic characteristics of 93-11 and Nipponbare recombination inbred lines based on the GoldenGate SNP assay. Science China Life Sciences, 2016, 59, 700-708.	4.9	0

#	Article	IF	Citations
127	Reply to Jin and Zhu: PINOID-mediated COP1 phosphorylation matters in photomorphogenesis in Arabidopsis. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8136-E8137.	7.1	0