

George Coupland

List of Publications by Year in descending order

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217
papers

34,769
citations

3874

91
h-index

4217

180
g-index

246
all docs

246
docs citations

246
times ranked

18857
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Arabis alpina</i> : A perennial model plant for ecological genomics and life history evolution. <i>Molecular Ecology Resources</i> , 2022, 22, 468-486.	2.2	7
2	A rice single cell transcriptomic atlas defines the developmental trajectories of rice floret and inflorescence meristems. <i>New Phytologist</i> , 2022, 234, 494-512.	3.5	41
3	MicroRNA172 controls inflorescence meristem size through regulation of APETALA2 in Arabidopsis. <i>New Phytologist</i> , 2022, 235, 356-371.	3.5	10
4	Gene regulatory networks controlled by FLOWERING LOCUS C that confer variation in seasonal flowering and life history. <i>Journal of Experimental Botany</i> , 2021, 72, 4-14.	2.4	41
5	Cytokinin-promoted secondary growth and nutrient storage in the perennial stem zone of <i>Arabis alpina</i> . <i>Plant Journal</i> , 2021, 105, 1459-1476.	2.8	5
6	Systematic analyses of the MIR172 family members of Arabidopsis define their distinct roles in regulation of APETALA2 during floral transition. <i>PLoS Biology</i> , 2021, 19, e3001043.	2.6	44
7	Unraveling the role of MADS transcription factor complexes in apple tree dormancy. <i>New Phytologist</i> , 2021, 232, 2071-2088.	3.5	31
8	Transposition and duplication of MADS-domain transcription factor genes in annual and perennial <i>Arabis</i> species modulates flowering. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	6
9	Genetic and Molecular Analysis of Root Hair Development in <i>Arabis alpina</i> . <i>Frontiers in Plant Science</i> , 2021, 12, 767772.	1.7	2
10	Functional Divergence of the Arabidopsis Florigen-Interacting bZIP Transcription Factors FD and FDP. <i>Cell Reports</i> , 2020, 31, 107717.	2.9	49
11	Mutagenesis of a Quintuple Mutant Impaired in Environmental Responses Reveals Roles for <i>CHROMATIN REMODELING4</i> in the Arabidopsis Floral Transition. <i>Plant Cell</i> , 2020, 32, 1479-1500.	3.1	17
12	The sugar transporter SWEET10 acts downstream of FLOWERING LOCUS T during floral transition of <i>Arabidopsis thaliana</i> . <i>BMC Plant Biology</i> , 2020, 20, 53.	1.6	59
13	Regulation of shoot meristem shape by photoperiodic signaling and phytohormones during floral induction of Arabidopsis. <i>ELife</i> , 2020, 9, .	2.8	30
14	A regulatory circuit conferring varied flowering response to cold in annual and perennial plants. <i>Science</i> , 2019, 363, 409-412.	6.0	69
15	FLOWERING LOCUS C Isolation and Characterization: Two Articles That Opened Many Doors. <i>Plant Cell</i> , 2019, 31, 1190-1191.	3.1	8
16	Genetic and molecular analysis of trichome development in <i>Arabis alpina</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12078-12083.	3.3	28
17	Gibberellins Act Downstream of <i>Arabis</i> PERPETUAL FLOWERING1 to Accelerate Floral Induction during Vernalization. <i>Plant Physiology</i> , 2019, 180, 1549-1563.	2.3	17
18	Floral regulators FLC and SOC1 directly regulate expression of the B3-type transcription factor TARGET OF FLC AND SVP 1 at the Arabidopsis shoot apex via antagonistic chromatin modifications. <i>PLoS Genetics</i> , 2019, 15, e1008065.	1.5	48

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19	Ubiquitin carboxyl-terminal hydrolases are required for period maintenance of the circadian clock at high temperature in <i>Arabidopsis</i> . <i>Scientific Reports</i> , 2019, 9, 17030.	1.6	17
20	Linking genes with ecological strategies in <i>Arabidopsis thaliana</i> . <i>Journal of Experimental Botany</i> , 2019, 70, 1141-1151.	2.4	37
21	<i>PERPETUAL FLOWERING2</i> coordinates the vernalization response and perennial flowering in <i>Arabis alpina</i> . <i>Journal of Experimental Botany</i> , 2019, 70, 949-961.	2.4	17
22	Floral homeotic proteins modulate the genetic program for leaf development to suppress trichome formation in flowers. <i>Development (Cambridge)</i> , 2018, 145, .	1.2	21
23	Demography and mating system shape the genome-wide impact of purifying selection in <i>Arabis alpina</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 816-821.	3.3	55
24	Differential effects of light-to-dark transitions on phase setting in circadian expression among clock-controlled genes in <i>Pharbitis nil</i> . <i>Plant Signaling and Behavior</i> , 2018, 13, e1473686.	1.2	4
25	Revised nomenclature and functional overview of the ULP gene family of plant deSUMOylating proteases. <i>Journal of Experimental Botany</i> , 2018, 69, 4505-4509.	2.4	20
26	Competence to Flower: Age-Controlled Sensitivity to Environmental Cues. <i>Plant Physiology</i> , 2017, 173, 36-46.	2.3	100
27	PSEUDO RESPONSE REGULATORS stabilize CONSTANS protein to promote flowering in response to day length. <i>EMBO Journal</i> , 2017, 36, 904-918.	3.5	103
28	Improving and correcting the contiguity of long-read genome assemblies of three plant species using optical mapping and chromosome conformation capture data. <i>Genome Research</i> , 2017, 27, 778-786.	2.4	155
29	Divergence of annual and perennial species in the Brassicaceae and the contribution of cis-acting variation at <i>FLC</i> orthologues. <i>Molecular Ecology</i> , 2017, 26, 3437-3457.	2.0	63
30	The Root Growth-Regulating Brevicompanine Natural Products Modulate the Plant Circadian Clock. <i>ACS Chemical Biology</i> , 2017, 12, 1466-1471.	1.6	9
31	Root-associated fungal microbiota of nonmycorrhizal <i>Arabis alpina</i> and its contribution to plant phosphorus nutrition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E9403-E9412.	3.3	239
32	Two SUMO Proteases SUMO PROTEASE RELATED TO FERTILITY1 and 2 Are Required for Fertility in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2017, 175, 1703-1719.	2.3	31
33	Evolution of the selfing syndrome: Anther orientation and herkogamy together determine reproductive assurance in a self-compatible plant. <i>Evolution; International Journal of Organic Evolution</i> , 2017, 71, 2206-2218.	1.1	44
34	Divergence of regulatory networks governed by the orthologous transcription factors FLC and PEP1 in Brassicaceae species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E11037-E11046.	3.3	50
35	The timing of <i>GIGANTEA</i> expression during day/night cycles varies with the geographical origin of <i>Arabidopsis</i> accessions. <i>Plant Signaling and Behavior</i> , 2017, 12, e1342026.	1.2	16
36	Root microbiota dynamics of perennial <i>Arabis alpina</i> are dependent on soil residence time but independent of flowering time. <i>ISME Journal</i> , 2017, 11, 43-55.	4.4	133

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37	Pinpointing genes underlying annual/perennial transitions with comparative genomics. <i>BMC Genomics</i> , 2016, 17, 921.	1.2	16
38	Photoperiodic and thermosensory pathways interact through <i>CONSTANS</i> to promote flowering at high temperature under short days. <i>Plant Journal</i> , 2016, 86, 426-440.	2.8	100
39	Multi-layered Regulation of <i>SPL15</i> and Cooperation with <i>SOC1</i> Integrate Endogenous Flowering Pathways at the Arabidopsis Shoot Meristem. <i>Developmental Cell</i> , 2016, 37, 254-266.	3.1	174
40	Editorial: Recent Advances in Flowering Time Control. <i>Frontiers in Plant Science</i> , 2016, 7, 2011.	1.7	26
41	Sample Preparation of Arabidopsis thaliana Shoot Apices for Expression Studies of Photoperiod-Induced Genes. <i>Methods in Molecular Biology</i> , 2016, 1398, 81-91.	0.4	0
42	A Luciferase-Based Assay to Test Whether Gene Expression Responses to Environmental Inputs Are Temporally Restricted by the Circadian Clock. <i>Methods in Molecular Biology</i> , 2016, 1398, 93-106.	0.4	0
43	Mechanisms controlling time measurement in plants and their significance in natural populations. , 2016, , 187-208.		0
44	Phosphorylation of <i>CONSTANS</i> and its <i>COP1</i> -dependent degradation during photoperiodic flowering of Arabidopsis. <i>Plant Journal</i> , 2015, 84, 451-463.	2.8	59
45	Genome expansion of Arabis alpina linked with retrotransposition and reduced symmetric DNA methylation. <i>Nature Plants</i> , 2015, 1, 14023.	4.7	156
46	The dynamics of <i>FLOWERING LOCUS T</i> expression encodes long-day information. <i>Plant Journal</i> , 2015, 83, 952-961.	2.8	33
47	The Plant Cell Introduces Breakthrough Reports: A New Forum for Cutting-Edge Plant Research. <i>Plant Cell</i> , 2015, , tpc.15.00862.	3.1	1
48	The <i>GL</i> <i>CDF</i> module of Arabidopsis affects freezing tolerance and growth as well as flowering. <i>Plant Journal</i> , 2015, 81, 695-706.	2.8	104
49	Natural diversity in daily rhythms of gene expression contributes to phenotypic variation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 905-910.	3.3	68
50	Site-directed mutagenesis in Arabidopsis thaliana using dividing tissue-targeted RGEN of the CRISPR/Cas system to generate heritable null alleles. <i>Planta</i> , 2015, 241, 271-284.	1.6	159
51	<i>SWP73</i> Subunits of Arabidopsis <i>SWI/SNF</i> Chromatin Remodeling Complexes Play Distinct Roles in Leaf and Flower Development. <i>Plant Cell</i> , 2015, 27, 1889-1906.	3.1	42
52	The Arabidopsis DNA Polymerase δ Has a Role in the Deposition of Transcriptionally Active Epigenetic Marks, Development and Flowering. <i>PLoS Genetics</i> , 2015, 11, e1004975.	1.5	36
53	Evolution of <i>CONSTANS</i> Regulation and Function after Gene Duplication Produced a Photoperiodic Flowering Switch in the Brassicaceae. <i>Molecular Biology and Evolution</i> , 2015, 32, 2284-2301.	3.5	49
54	Combinatorial activities of <i>SHORT VEGETATIVE PHASE</i> and <i>FLOWERING LOCUS C</i> define distinct modes of flowering regulation in Arabidopsis. <i>Genome Biology</i> , 2015, 16, 31.	3.8	150

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55	Floral induction in <i>Arabidopsis thaliana</i> by FLOWERING LOCUS T requires direct repression of BLADE-ON-PETIOLE genes by homeodomain protein PENNYWISE. <i>Plant Physiology</i> , 2015, 169, pp.00960.2015.	2.3	51
56	Large-scale adaptive differentiation in the alpine perennial herb <i>Androsace alpina</i> . <i>New Phytologist</i> , 2015, 206, 459-470.	3.5	36
57	Deeper Rooting: Pflanzenanzüchtung und die Herausforderungen des Klimawandels. , 2015, , 153-166.		0
58	Elevated Levels of MYB30 in the Phloem Accelerate Flowering in <i>Arabidopsis</i> through the Regulation of FLOWERING LOCUS T. <i>PLoS ONE</i> , 2014, 9, e89799.	1.1	30
59	<i>Arabidopsis</i> florigen FT binds to diurnally oscillating phospholipids that accelerate flowering. <i>Nature Communications</i> , 2014, 5, 3553.	5.8	143
60	SHORT VEGETATIVE PHASE reduces gibberellin biosynthesis at the <i>Arabidopsis</i> shoot apex to regulate the floral transition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E2760-9.	3.3	132
61	Evolutionary conservation of cold-induced antisense RNAs of FLOWERING LOCUS C in <i>Arabidopsis thaliana</i> perennial relatives. <i>Nature Communications</i> , 2014, 5, 4457.	5.8	72
62	Diurnal and circadian expression profiles of glycerolipid biosynthetic genes in <i>Arabidopsis</i> . <i>Plant Signaling and Behavior</i> , 2014, 9, e29715.	1.2	21
63	A tribute to Ko Shimamoto (1949-2013). <i>Journal of Experimental Botany</i> , 2014, 65, 6755-6759.	2.4	0
64	Evening Expression of <i>Arabidopsis GIGANTEA</i> Is Controlled by Combinatorial Interactions among Evolutionarily Conserved Regulatory Motifs. <i>Plant Cell</i> , 2014, 26, 3999-4018.	3.1	17
65	Analysis of TTG1 function in <i>Arabidopsis thaliana</i> . <i>BMC Plant Biology</i> , 2014, 14, 16.	1.6	25
66	Elevated salicylic acid levels conferred by increased expression of <i>ISOGORISMATE SYNTHASE</i> 1 contribute to hyperaccumulation of <i>SUMO</i> 1 conjugates in the <i>Arabidopsis</i> mutant <i>early in short days 4</i> . <i>Plant Journal</i> , 2014, 79, 206-219.	2.8	42
67	<i>miR824</i> Regulated <i>AGAMOUS-LIKE16</i> Contributes to Flowering Time Repression in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 2024-2037.	3.1	112
68	NATURAL VARIATION IN EPIGENETIC GENE REGULATION AND ITS EFFECTS ON PLANT DEVELOPMENTAL TRAITS. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 620-631.	1.1	38
69	Flowering responses to seasonal cues: what's new?. <i>Current Opinion in Plant Biology</i> , 2014, 21, 120-127.	3.5	91
70	The (r)evolution of gene regulatory networks controlling <i>Arabidopsis</i> plant reproduction: a two-decade history. <i>Journal of Experimental Botany</i> , 2014, 65, 4731-4745.	2.4	106
71	Identification of pathways directly regulated by SHORT VEGETATIVE PHASE during vegetative and reproductive development in <i>Arabidopsis</i> . <i>Genome Biology</i> , 2013, 14, R56.	3.8	134
72	DELLA-Interacting SWI3C Core Subunit of Switch/Sucrose Nonfermenting Chromatin Remodeling Complex Modulates Gibberellin Responses and Hormonal Cross Talk in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2013, 163, 305-317.	2.3	98

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73	Identification of <i>Arabidopsis</i> SUMO-interacting proteins that regulate chromatin activity and developmental transitions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 19956-19961.	3.3	66
74	Mechanisms of Age-Dependent Response to Winter Temperature in Perennial Flowering of <i>Arabis alpina</i> . <i>Science</i> , 2013, 340, 1094-1097.	6.0	207
75	Small Ubiquitin-Like Modifier Conjugating Enzyme with Active Site Mutation Acts as Dominant Negative Inhibitor of SUMO Conjugation in <i>Arabidopsis</i> ^F . <i>Journal of Integrative Plant Biology</i> , 2013, 55, 75-82.	4.1	16
76	A Molecular Framework for Auxin-Mediated Initiation of Flower Primordia. <i>Developmental Cell</i> , 2013, 24, 271-282.	3.1	262
77	Mutation identification by direct comparison of whole-genome sequencing data from mutant and wild-type individuals using k-mers. <i>Nature Biotechnology</i> , 2013, 31, 325-330.	9.4	149
78	PEP1 of <i>Arabis alpina</i> Is Encoded by Two Overlapping Genes That Contribute to Natural Genetic Variation in Perennial Flowering. <i>PLoS Genetics</i> , 2012, 8, e1003130.	1.5	69
79	Analysis of the <i>Arabidopsis</i> Shoot Meristem Transcriptome during Floral Transition Identifies Distinct Regulatory Patterns and a Leucine-Rich Repeat Protein That Promotes Flowering. <i>Plant Cell</i> , 2012, 24, 444-462.	3.1	178
80	EARLY FLOWERING4 Recruitment of EARLY FLOWERING3 in the Nucleus Sustains the <i>Arabidopsis</i> Circadian Clock. <i>Plant Cell</i> , 2012, 24, 428-443.	3.1	275
81	The Circadian Clock-Associated Small GTPase LIGHT INSENSITIVE PERIOD1 Suppresses Light-Controlled Endoreplication and Affects Tolerance to Salt Stress in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2012, 161, 278-290.	2.3	8
82	The genetic basis of flowering responses to seasonal cues. <i>Nature Reviews Genetics</i> , 2012, 13, 627-639.	7.7	1,200
83	Spatially distinct regulatory roles for gibberellins in the promotion of flowering of <i>Arabidopsis</i> under long photoperiods. <i>Development (Cambridge)</i> , 2012, 139, 2198-2209.	1.2	193
84	Mutation in <i>TERMINAL FLOWER1</i> Reverses the Photoperiodic Requirement for Flowering in the Wild Strawberry <i>Fragaria vesca</i> . <i>Plant Physiology</i> , 2012, 159, 1043-1054.	2.3	158
85	Functional characterisation of <i>HvCO1</i> , the barley (<i>Hordeum vulgare</i>) flowering time ortholog of <i>CONSTANS</i> . <i>Plant Journal</i> , 2012, 69, 868-880.	2.8	136
86	Prieurianin/endosidin ϵ 1 is an actin-stabilizing small molecule identified from a chemical genetic screen for circadian clock effectors in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 2012, 71, 338-352.	2.8	53
87	Comparing genetic diversity within a crop and its wild progenitor: a case study for barley.., 2012, , 186-192.		0
88	When Vernalization Makes Sense. <i>Science</i> , 2011, 331, 36-37.	6.0	17
89	DOF-binding sites additively contribute to guard cell-specificity of <i>AtMYB60</i> promoter. <i>BMC Plant Biology</i> , 2011, 11, 162.	1.6	65
90	Cytokinin promotes flowering of <i>Arabidopsis</i> via transcriptional activation of the <i>FT</i> paralogue <i>TSF</i> . <i>Plant Journal</i> , 2011, 65, 972-979.	2.8	172

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91	The Arabidopsis <i>SOC1</i> -like genes <i>AGL42</i> , <i>AGL71</i> and <i>AGL72</i> promote flowering in the shoot apical and axillary meristems. <i>Plant Journal</i> , 2011, 67, 1006-1017.	2.8	117
92	Distinct roles for Arabidopsis SUMO protease ESD4 and its closest homolog ELS1. <i>Planta</i> , 2011, 233, 63-73.	1.6	52
93	Nitrate regulates floral induction in Arabidopsis, acting independently of light, gibberellin and autonomous pathways. <i>Planta</i> , 2011, 233, 539-552.	1.6	158
94	Aa <i>TFL1</i> Confers an Age-Dependent Response to Vernalization in Perennial <i>Arabis alpina</i> . <i>Plant Cell</i> , 2011, 23, 1307-1321.	3.1	117
95	Speeding Cis-Trans Regulation Discovery by Phylogenomic Analyses Coupled with Screenings of an Arrayed Library of Arabidopsis Transcription Factors. <i>PLoS ONE</i> , 2011, 6, e21524.	1.1	78
96	Plant development goes like clockwork. <i>Trends in Genetics</i> , 2010, 26, 296-306.	2.9	166
97	SynRg - Biotechnologie zur Steigerung von Ertrag und Ertragsstabilität nachwachsender Rohstoffe. <i>Chemie-Ingenieur-Technik</i> , 2010, 82, 1517-1518.	0.4	0
98	Proteome-wide screens for small ubiquitin-like modifier (SUMO) substrates identify <i>Arabidopsis</i> proteins implicated in diverse biological processes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 17415-17420.	3.3	159
99	Comparative Analysis of Flowering in Annual and Perennial Plants. <i>Current Topics in Developmental Biology</i> , 2010, 91, 323-348.	1.0	130
100	SnapShot: Control of Flowering in Arabidopsis. <i>Cell</i> , 2010, 141, 550-550.e2.	13.5	529
101	<i>cis</i> -Regulatory Elements and Chromatin State Coordinately Control Temporal and Spatial Expression of <i>FLOWERING LOCUS T</i> in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2010, 22, 1425-1440.	3.1	274
102	Effects of Genetic Perturbation on Seasonal Life History Plasticity. <i>Science</i> , 2009, 323, 930-934.	6.0	340
103	Distinct Patterns of Genetic Variation Alter Flowering Responses of Arabidopsis Accessions to Different Daylengths. <i>Plant Physiology</i> , 2009, 152, 177-191.	2.3	26
104	Substrates Related to Chromatin and to RNA-Dependent Processes Are Modified by Arabidopsis SUMO Isoforms That Differ in a Conserved Residue with Influence on Desumoylation. <i>Plant Physiology</i> , 2009, 149, 1529-1540.	2.3	91
105	The <i>Arabidopsis</i> B-Box Zinc Finger Family. <i>Plant Cell</i> , 2009, 21, 3416-3420.	3.1	306
106	<i>Chlamydomonas</i> CONSTANS and the Evolution of Plant Photoperiodic Signaling. <i>Current Biology</i> , 2009, 19, 359-368.	1.8	106
107	Control of perennial flowering and perenniality in <i>Arabis alpina</i> , a relative of <i>Arabidopsis thaliana</i> . <i>Comparative Biochemistry and Physiology Part A, Molecular & Integrative Physiology</i> , 2009, 153, S195-S196.	0.8	2
108	Genetic and spatial interactions between <i>FT</i> , <i>TSF</i> and <i>SVP</i> during the early stages of floral induction in Arabidopsis. <i>Plant Journal</i> , 2009, 60, 614-625.	2.8	194

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109	PEP1 regulates perennial flowering in <i>Arabis alpina</i> . <i>Nature</i> , 2009, 459, 423-427.	13.7	325
110	Possible role of EARLY FLOWERING 3 (ELF3) in clock-dependent floral regulation by SHORT VEGETATIVE PHASE (SVP) in <i>Arabidopsis thaliana</i> . <i>New Phytologist</i> , 2009, 182, 838-850.	3.5	48
111	<i>Arabidopsis</i> DOF Transcription Factors Act Redundantly to Reduce CONSTANS Expression and Are Essential for a Photoperiodic Flowering Response. <i>Developmental Cell</i> , 2009, 17, 75-86.	3.1	493
112	Plant Phase Transitions Make a SPLash. <i>Cell</i> , 2009, 138, 625-627.	13.5	80
113	<i>Arabidopsis</i> COP1 shapes the temporal pattern of CO accumulation conferring a photoperiodic flowering response. <i>EMBO Journal</i> , 2008, 27, 1277-1288.	3.5	424
114	Genome-scale <i>Arabidopsis</i> promoter array identifies targets of the histone acetyltransferase GCN5. <i>Plant Journal</i> , 2008, 56, 493-504.	2.8	120
115	Phloem transport of flowering signals. <i>Current Opinion in Plant Biology</i> , 2008, 11, 687-694.	3.5	71
116	Regulation and Identity of Florigen: FLOWERING LOCUS T Moves Center Stage. <i>Annual Review of Plant Biology</i> , 2008, 59, 573-594.	8.6	889
117	The impact of chromatin regulation on the floral transition. <i>Seminars in Cell and Developmental Biology</i> , 2008, 19, 560-573.	2.3	69
118	Circadian Clock Proteins LHY and CCA1 Regulate SVP Protein Accumulation to Control Flowering in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2008, 20, 2960-2971.	3.1	180
119	A Circadian Rhythm Set by Dusk Determines the Expression of FT Homologs and the Short-Day Photoperiodic Flowering Response in <i>Pharbitis</i> . <i>Plant Cell</i> , 2007, 19, 2988-3000.	3.1	181
120	<i>Arabidopsis</i> TFL2/LHP1 Specifically Associates with Genes Marked by Trimethylation of Histone H3 Lysine 27. <i>PLoS Genetics</i> , 2007, 3, e86.	1.5	537
121	FT Protein Movement Contributes to Long-Distance Signaling in Floral Induction of <i>Arabidopsis</i> . <i>Science</i> , 2007, 316, 1030-1033.	6.0	1,855
122	Isolation of novel gain- and loss-of-function alleles of the circadian clock gene LATE ELONGATED HYPOCOTYL (LHY) in <i>Arabidopsis</i> . <i>Plant Biotechnology</i> , 2007, 24, 457-465.	0.5	2
123	<i>Arabidopsis</i> SPA proteins regulate photoperiodic flowering and interact with the floral inducer CONSTANS to regulate its stability. <i>Development (Cambridge)</i> , 2006, 133, 4608-4608.	1.2	2
124	The CCAAT binding factor can mediate interactions between CONSTANS-like proteins and DNA. <i>Plant Journal</i> , 2006, 46, 462-476.	2.8	247
125	<i>Arabidopsis</i> SPA proteins regulate photoperiodic flowering and interact with the floral inducer CONSTANS to regulate its stability. <i>Development (Cambridge)</i> , 2006, 133, 3213-3222.	1.2	272
126	The quest for florigen: a review of recent progress. <i>Journal of Experimental Botany</i> , 2006, 57, 3395-3403.	2.4	185

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127	CONSTANS and the CCAAT Box Binding Complex Share a Functionally Important Domain and Interact to Regulate Flowering of Arabidopsis. <i>Plant Cell</i> , 2006, 18, 2971-2984.	3.1	512
128	The transcription factor FLC confers a flowering response to vernalization by repressing meristem competence and systemic signaling in Arabidopsis. <i>Genes and Development</i> , 2006, 20, 898-912.	2.7	744
129	THE MOLECULAR GENETICS OF PHOTO PERIODIC RESPONSES:COMPARISONS BETWEEN LONG-DAY AND SHORT-DAY SPECIES. , 2006, , 605-625.		0
130	The Family of CONSTANS-Like Genes in <i>Physcomitrella patens</i> . <i>Plant Biology</i> , 2005, 7, 266-275.	1.8	55
131	Photoperiodic flowering of Arabidopsis: integrating genetic and physiological approaches to characterization of the floral stimulus. <i>Plant, Cell and Environment</i> , 2005, 28, 54-66.	2.8	126
132	Cell signalling and gene regulation. <i>Current Opinion in Plant Biology</i> , 2005, 8, 457-461.	3.5	4
133	Circadian clock components in Arabidopsis I. The terminal flower 1 enhances the early flowering phenotype of a mutant, <i>lhy cca1</i> . <i>Plant Biotechnology</i> , 2005, 22, 311-317.	0.5	7
134	Distinct Roles of GIGANTEA in Promoting Flowering and Regulating Circadian Rhythms in Arabidopsis. <i>Plant Cell</i> , 2005, 17, 2255-2270.	3.1	408
135	A rapid and versatile combined DNA/RNA extraction protocol and its application to the analysis of a novel DNA marker set polymorphic between <i>Arabidopsis thaliana</i> ecotypes Col-0 and <i>Landsberg erecta</i> . <i>Plant Methods</i> , 2005, 1, 4.	1.9	67
136	Circadian clock components in Arabidopsis II. LHY/CCA1 regulate the floral integrator gene SOC1 in both GI-dependent and -independent pathways. <i>Plant Biotechnology</i> , 2005, 22, 319-325.	0.5	10
137	CONSTANS acts in the phloem to regulate a systemic signal that induces photoperiodic flowering of Arabidopsis. <i>Development (Cambridge)</i> , 2004, 131, 3615-3626.	1.2	573
138	The Molecular Basis of Diversity in the Photoperiodic Flowering Responses of Arabidopsis and Rice. <i>Plant Physiology</i> , 2004, 135, 677-684.	2.3	271
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