

George Coupland

List of Publications by Year in descending order

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218
papers

34,769
citations

3334

91
h-index

3650

180
g-index

246
all docs

246
docs citations

246
times ranked

17024
citing authors

#	ARTICLE	IF	CITATIONS
1	FT Protein Movement Contributes to Long-Distance Signaling in Floral Induction of Arabidopsis. <i>Science</i> , 2007, 316, 1030-1033.	12.6	1,855
2	The CONSTANS gene of arabidopsis promotes flowering and encodes a protein showing similarities to zinc finger transcription factors. <i>Cell</i> , 1995, 80, 847-857.	28.9	1,287
3	Distinct Roles of CONSTANS Target Genes in Reproductive Development of Arabidopsis. <i>Science</i> , 2000, 288, 1613-1616.	12.6	1,272
4	CONSTANS mediates between the circadian clock and the control of flowering in Arabidopsis. <i>Nature</i> , 2001, 410, 1116-1120.	27.8	1,258
5	The genetic basis of flowering responses to seasonal cues. <i>Nature Reviews Genetics</i> , 2012, 13, 627-639.	16.3	1,200
6	Photoreceptor Regulation of CONSTANS Protein in Photoperiodic Flowering. <i>Science</i> , 2004, 303, 1003-1006.	12.6	1,089
7	Regulation and Identity of Florigen: FLOWERING LOCUS T Moves Center Stage. <i>Annual Review of Plant Biology</i> , 2008, 59, 573-594.	18.7	889
8	The late elongated hypocotyl Mutation of Arabidopsis Disrupts Circadian Rhythms and the Photoperiodic Control of Flowering. <i>Cell</i> , 1998, 93, 1219-1229.	28.9	805
9	Control of Flowering Time. <i>Plant Cell</i> , 2002, 14, S111-S130.	6.6	785
10	A Polycomb-group gene regulates homeotic gene expression in Arabidopsis. <i>Nature</i> , 1997, 386, 44-51.	27.8	760
11	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.	5.9	744
12	GIGANTEA: a circadian clock-controlled gene that regulates photoperiodic flowering in Arabidopsis and encodes a protein with several possible membrane-spanning domains. <i>EMBO Journal</i> , 1999, 18, 4679-4688.	7.8	691
13	CONSTANS acts in the phloem to regulate a systemic signal that induces photoperiodic flowering of Arabidopsis. <i>Development (Cambridge)</i> , 2004, 131, 3615-3626.	2.5	573
14	LHY and CCA1 Are Partially Redundant Genes Required to Maintain Circadian Rhythms in Arabidopsis. <i>Developmental Cell</i> , 2002, 2, 629-641.	7.0	572
15	Arabidopsis TFL2/LHP1 Specifically Associates with Genes Marked by Trimethylation of Histone H3 Lysine 27. <i>PLoS Genetics</i> , 2007, 3, e86.	3.5	537
16	SnapShot: Control of Flowering in Arabidopsis. <i>Cell</i> , 2010, 141, 550-550.e2.	28.9	529
17	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.	6.6	512
18	Arabidopsis 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.	7.0	493

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19	The Evolution of CONSTANS-Like Gene Families in Barley, Rice, and Arabidopsis. <i>Plant Physiology</i> , 2003, 131, 1855-1867.	4.8	463
20	Antagonistic regulation of flowering-time gene SOC1 by CONSTANS and FLC via separate promoter motifs. <i>EMBO Journal</i> , 2002, 21, 4327-4337.	7.8	432
21	Arabidopsis COP1 shapes the temporal pattern of CO accumulation conferring a photoperiodic flowering response. <i>EMBO Journal</i> , 2008, 27, 1277-1288.	7.8	424
22	Distinct Roles of GIGANTEA in Promoting Flowering and Regulating Circadian Rhythms in Arabidopsis. <i>Plant Cell</i> , 2005, 17, 2255-2270.	6.6	408
23	Functional importance of conserved domains in the flowering-time gene CONSTANS demonstrated by analysis of mutant alleles and transgenic plants. <i>Plant Journal</i> , 2002, 28, 619-631.	5.7	397
24	Mutagenesis of Plants Overexpressing CONSTANS Demonstrates Novel Interactions among Arabidopsis Flowering-Time Genes. <i>Plant Cell</i> , 2000, 12, 885-900.	6.6	360
25	Activation of floral meristem identity genes in Arabidopsis. <i>Nature</i> , 1996, 384, 59-62.	27.8	351
26	Effects of Genetic Perturbation on Seasonal Life History Plasticity. <i>Science</i> , 2009, 323, 930-934.	12.6	340
27	Mutations in the Arabidopsis Gene IMMUTANS Cause a Variegated Phenotype by Inactivating a Chloroplast Terminal Oxidase Associated with Phytoene Desaturation. <i>Plant Cell</i> , 1999, 11, 57-68.	6.6	326
28	PEP1 regulates perennial flowering in <i>Arabis alpina</i> . <i>Nature</i> , 2009, 459, 423-427.	27.8	325
29	The <i>Arabidopsis</i> B-Box Zinc Finger Family. <i>Plant Cell</i> , 2009, 21, 3416-3420.	6.6	306
30	EARLY FLOWERING4 Recruitment of EARLY FLOWERING3 in the Nucleus Sustains the <i>Arabidopsis</i> Circadian Clock. <i>Plant Cell</i> , 2012, 24, 428-443.	6.6	275
31	<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.	6.6	274
32	Arabidopsis SPA proteins regulate photoperiodic flowering and interact with the floral inducer CONSTANS to regulate its stability. <i>Development (Cambridge)</i> , 2006, 133, 3213-3222.	2.5	272
33	The Molecular Basis of Diversity in the Photoperiodic Flowering Responses of Arabidopsis and Rice. <i>Plant Physiology</i> , 2004, 135, 677-684.	4.8	271
34	A Molecular Framework for Auxin-Mediated Initiation of Flower Primordia. <i>Developmental Cell</i> , 2013, 24, 271-282.	7.0	262
35	ACAULIS5, an Arabidopsis gene required for stem elongation, encodes a spermine synthase. <i>EMBO Journal</i> , 2000, 19, 4248-4256.	7.8	252
36	Induction of flowering by seasonal changes in photoperiod. <i>EMBO Journal</i> , 2004, 23, 1217-1222.	7.8	252

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37	A Dissociation insertion causes a semidominant mutation that increases expression of TINY, an Arabidopsis gene related to APETALA2.. Plant Cell, 1996, 8, 659-671.	6.6	248
38	The CCAAT binding factor can mediate interactions between CONSTANS-like proteins and DNA. Plant Journal, 2006, 46, 462-476.	5.7	247
39	Root-associated fungal microbiota of nonmycorrhizal <i>Arabidopsis thaliana</i> and its contribution to plant phosphorus nutrition. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E9403-E9412.	7.1	239
40	Shedding light on the circadian clock and the photoperiodic control of flowering. Current Opinion in Plant Biology, 2003, 6, 13-19.	7.1	228
41	Analysis of Natural Allelic Variation at Flowering Time Loci in the Landsberg erecta and Cape Verde Islands Ecotypes of Arabidopsis thaliana. Genetics, 1998, 149, 749-764.	2.9	225
42	Comparative mapping in Arabidopsis and Brassica, fine scale genome collinearity and congruence of genes controlling flowering time. Plant Journal, 1996, 9, 13-20.	5.7	222
43	ALBINO3, an Arabidopsis nuclear gene essential for chloroplast differentiation, encodes a chloroplast protein that shows homology to proteins present in bacterial membranes and yeast mitochondria.. Plant Cell, 1997, 9, 717-730.	6.6	209
44	Mechanisms of Age-Dependent Response to Winter Temperature in Perennial Flowering of <i>Arabidopsis thaliana</i> . Science, 2013, 340, 1094-1097.	12.6	207
45	A Nuclear Protease Required for Flowering-Time Regulation in Arabidopsis Reduces the Abundance of SMALL UBIQUITIN-RELATED MODIFIER Conjugates. Plant Cell, 2003, 15, 2308-2319.	6.6	204
46	Characterization of the ethanol-inducible alc gene-expression system in Arabidopsis thaliana. Plant Journal, 2001, 28, 225-235.	5.7	198
47	Genetic and spatial interactions between <i>FT</i> , <i>TSF</i> and <i>SVP</i> during the early stages of floral induction in Arabidopsis. Plant Journal, 2009, 60, 614-625.	5.7	194
48	Spatially distinct regulatory roles for gibberellins in the promotion of flowering of <i>Arabidopsis</i> under long photoperiods. Development (Cambridge), 2012, 139, 2198-2209.	2.5	193
49	The quest for florigen: a review of recent progress. Journal of Experimental Botany, 2006, 57, 3395-3403.	4.8	185
50	A Circadian Rhythm Set by Dusk Determines the Expression of <i>FT</i> Homologs and the Short-Day Photoperiodic Flowering Response in <i>Pharbitis</i> . Plant Cell, 2007, 19, 2988-3000.	6.6	181
51	Circadian Clock Proteins LHY and CCA1 Regulate SVP Protein Accumulation to Control Flowering in <i>Arabidopsis</i> . Plant Cell, 2008, 20, 2960-2971.	6.6	180
52	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. Plant Cell, 2012, 24, 444-462.	6.6	178
53	Multi-layered Regulation of SPL15 and Cooperation with SOC1 Integrate Endogenous Flowering Pathways at the Arabidopsis Shoot Meristem. Developmental Cell, 2016, 37, 254-266.	7.0	174
54	Cytokinin promotes flowering of Arabidopsis via transcriptional activation of the <i>FT</i> paralogue <i>TSF</i> . Plant Journal, 2011, 65, 972-979.	5.7	172

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55	Plant development goes like clockwork. <i>Trends in Genetics</i> , 2010, 26, 296-306.	6.7	166
56	Phenotypic assay for excision of the maize controlling element <i>Ac</i> in tobacco. <i>EMBO Journal</i> , 1987, 6, 1547-1554.	7.8	159
57	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.	7.1	159
58	Site-directed mutagenesis in <i>Arabidopsis thaliana</i> using dividing tissue-targeted RGEN of the CRISPR/Cas system to generate heritable null alleles. <i>Planta</i> , 2015, 241, 271-284.	3.2	159
59	Nitrate regulates floral induction in <i>Arabidopsis</i> , acting independently of light, gibberellin and autonomous pathways. <i>Planta</i> , 2011, 233, 539-552.	3.2	158
60	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.	4.8	158
61	Genome expansion of <i>Arabis alpina</i> linked with retrotransposition and reduced symmetric DNA methylation. <i>Nature Plants</i> , 2015, 1, 14023.	9.3	156
62	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.	5.5	155
63	Combinatorial activities of <i>SHORT VEGETATIVE PHASE</i> and <i>FLOWERING LOCUS C</i> define distinct modes of flowering regulation in <i>Arabidopsis</i> . <i>Genome Biology</i> , 2015, 16, 31.	8.8	150
64	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.	17.5	149
65	<i>Arabidopsis</i> florigen FT binds to diurnally oscillating phospholipids that accelerate flowering. <i>Nature Communications</i> , 2014, 5, 3553.	12.8	143
66	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.	5.7	136
67	Identification of pathways directly regulated by <i>SHORT VEGETATIVE PHASE</i> during vegetative and reproductive development in <i>Arabidopsis</i> . <i>Genome Biology</i> , 2013, 14, R56.	8.8	134
68	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.	9.8	133
69	<i>SHORT VEGETATIVE PHASE</i> 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.	7.1	132
70	<i>ALBINO3</i> , an <i>Arabidopsis</i> Nuclear Gene Essential for Chloroplast Differentiation, Encodes a Chloroplast Protein That Shows Homology to Proteins Present in Bacterial Membranes and Yeast Mitochondria. <i>Plant Cell</i> , 1997, 9, 717.	6.6	131
71	Comparative Analysis of Flowering in Annual and Perennial Plants. <i>Current Topics in Developmental Biology</i> , 2010, 91, 323-348.	2.2	130
72	The maize transposable element system <i>Ac/Ds</i> as a mutagen in <i>Arabidopsis</i> : identification of an albino mutation induced by <i>Ds</i> insertion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 10370-10374.	7.1	129

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73	Time measurement and the control of flowering in plants. <i>BioEssays</i> , 2000, 22, 38-47.	2.5	127
74	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.	5.7	126
75	Elevated levels of Activator transposase mRNA are associated with high frequencies of Dissociation excision in Arabidopsis.. <i>Plant Cell</i> , 1992, 4, 583-595.	6.6	123
76	Response of plant development to environment: control of flowering by daylength and temperature. <i>Current Opinion in Plant Biology</i> , 2000, 3, 37-42.	7.1	122
77	Genetic and environmental control of flowering time in Arabidopsis. <i>Trends in Genetics</i> , 1995, 11, 393-397.	6.7	121
78	EARLY BOLTING IN SHORT DAYS Is Related to Chromatin Remodeling Factors and Regulates Flowering in Arabidopsis by Repressing FT. <i>Plant Cell</i> , 2003, 15, 1552-1562.	6.6	121
79	Genome-scale Arabidopsis promoter array identifies targets of the histone acetyltransferase GCN5. <i>Plant Journal</i> , 2008, 56, 493-504.	5.7	120
80	Signalling for developmental plasticity. <i>Trends in Plant Science</i> , 2004, 9, 309-314.	8.8	117
81	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.	5.7	117
82	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.	6.6	117
83	Sequences near the termini are required for transposition of the maize transposon Ac in transgenic tobacco plants.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1989, 86, 9385-9388.	7.1	113
84	<i>miR824</i> Regulated AGAMOUS-LIKE16 Contributes to Flowering Time Repression in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 2024-2037.	6.6	112
85	Analysis of Flowering Time Control in Arabidopsis by Comparison of Double and Triple Mutants. <i>Plant Physiology</i> , 2001, 126, 1085-1091.	4.8	111
86	The Control of Flowering Time and Floral Identity in Arabidopsis1. <i>Plant Physiology</i> , 1998, 117, 1-8.	4.8	110
87	Transposition of the maize transposable element Ac in <i>Solanum tuberosum</i> . <i>Molecular Genetics and Genomics</i> , 1988, 213, 285-290.	2.4	106
88	Chlamydomonas CONSTANS and the Evolution of Plant Photoperiodic Signaling. <i>Current Biology</i> , 2009, 19, 359-368.	3.9	106
89	The (r)evolution of gene regulatory networks controlling Arabidopsis plant reproduction: a two-decade history. <i>Journal of Experimental Botany</i> , 2014, 65, 4731-4745.	4.8	106
90	The <i>GI</i> - <i>CDF</i> module of Arabidopsis affects freezing tolerance and growth as well as flowering. <i>Plant Journal</i> , 2015, 81, 695-706.	5.7	104

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91	Conserved structure and function of the Arabidopsis flowering time gene CONSTANS in Brassica napus. <i>Plant Molecular Biology</i> , 1998, 37, 763-772.	3.9	103
92	PSEUDO RESPONSE REGULATORS stabilize CONSTANS protein to promote flowering in response to day length. <i>EMBO Journal</i> , 2017, 36, 904-918.	7.8	103
93	Characterization of the maize transposable element <i>Ac</i> by internal deletions. <i>EMBO Journal</i> , 1988, 7, 3653-3659.	7.8	101
94	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.	5.7	100
95	Competence to Flower: Age-Controlled Sensitivity to Environmental Cues. <i>Plant Physiology</i> , 2017, 173, 36-46.	4.8	100
96	DELLA-Interacting SWI3C Core Subunit of Switch/Sucrose Nonfermenting Chromatin Remodeling Complex Modulates Gibberellin Responses and Hormonal Cross Talk in Arabidopsis. <i>Plant Physiology</i> , 2013, 163, 305-317.	4.8	98
97	early in short days 4, a mutation in Arabidopsis that causes early flowering and reduces the mRNA abundance of the floral repressor FLC. <i>Development (Cambridge)</i> , 2002, 129, 5349-5361.	2.5	95
98	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.	4.8	91
99	Flowering responses to seasonal cues: what's new?. <i>Current Opinion in Plant Biology</i> , 2014, 21, 120-127.	7.1	91
100	SUMO conjugation in plants. <i>Planta</i> , 2004, 220, 1-8.	3.2	86
101	Phenotypic assay for excision of the maize controlling element <i>Ac</i> in tobacco. <i>EMBO Journal</i> , 1987, 6, 1547-54.	7.8	84
102	Plant Phase Transitions Make a SPLash. <i>Cell</i> , 2009, 138, 625-627.	28.9	80
103	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.	2.5	78
104	Evolutionary conservation of cold-induced antisense RNAs of FLOWERING LOCUS C in Arabidopsis thaliana perennial relatives. <i>Nature Communications</i> , 2014, 5, 4457.	12.8	72
105	early bolting in short days: An Arabidopsis Mutation That Causes Early Flowering and Partially Suppresses the Floral Phenotype of leafy. <i>Plant Cell</i> , 2001, 13, 1011-1024.	6.6	71
106	Phloem transport of flowering signals. <i>Current Opinion in Plant Biology</i> , 2008, 11, 687-694.	7.1	71
107	Arabidopsis A BOUT DE SOUFFLE, Which Is Homologous with Mammalian Carnitine Acyl Carrier, Is Required for Postembryonic Growth in the Light. <i>Plant Cell</i> , 2002, 14, 2161-2173.	6.6	69
108	REGIA, An EU Project on Functional Genomics of Transcription Factors from Arabidopsis thaliana. <i>Comparative and Functional Genomics</i> , 2002, 3, 102-108.	2.0	69

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109	The impact of chromatin regulation on the floral transition. <i>Seminars in Cell and Developmental Biology</i> , 2008, 19, 560-573.	5.0	69
110	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.	3.5	69
111	A regulatory circuit conferring varied flowering response to cold in annual and perennial plants. <i>Science</i> , 2019, 363, 409-412.	12.6	69
112	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.	7.1	68
113	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 Landsberg erecta. <i>Plant Methods</i> , 2005, 1, 4.	4.3	67
114	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.	7.1	66
115	Characterization of the maize transposable element <i>Ac</i> by internal deletions. <i>EMBO Journal</i> , 1988, 7, 3653-9.	7.8	66
116	Transposons as tools for the isolation of plant genes. <i>Trends in Biotechnology</i> , 1991, 9, 31-37.	9.3	65
117	DOF-binding sites additively contribute to guard cell-specificity of <i>AtMYB60</i> promoter. <i>BMC Plant Biology</i> , 2011, 11, 162.	3.6	65
118	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.	3.9	63
119	Mutation of a family 8 glycosyltransferase gene alters cell wall carbohydrate composition and causes a humidity-sensitive semi-sterile dwarf phenotype in <i>Arabidopsis</i> . <i>Plant Molecular Biology</i> , 2003, 53, 687-701.	3.9	61
120	Phosphorylation of <i>CONSTANS</i> and its <i>COP1</i> -dependent degradation during photoperiodic flowering of <i>Arabidopsis</i> . <i>Plant Journal</i> , 2015, 84, 451-463.	5.7	59
121	The sugar transporter <i>SWEET10</i> acts downstream of <i>FLOWERING LOCUS T</i> during floral transition of <i>Arabidopsis thaliana</i> . <i>BMC Plant Biology</i> , 2020, 20, 53.	3.6	59
122	The Family of <i>CONSTANS</i> -Like Genes in <i>Physcomitrella patens</i> . <i>Plant Biology</i> , 2005, 7, 266-275.	3.8	55
123	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.	7.1	55
124	Antisense suppression of the <i>Arabidopsis PIF3</i> gene does not affect circadian rhythms but causes early flowering and increases FT expression. <i>FEBS Letters</i> , 2004, 557, 259-264.	2.8	54
125	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.	5.7	53
126	Genetics of homology-dependent gene silencing in <i>Arabidopsis</i> ; a role for methylation. <i>Plant Journal</i> , 1997, 12, 791-804.	5.7	52

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127	Distinct roles for Arabidopsis SUMO protease ESD4 and its closest homolog ELS1. <i>Planta</i> , 2011, 233, 63-73.	3.2	52
128	Floral induction in Arabidopsis thaliana 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.	4.8	51
129	Chromosome walking with YAC clones in Arabidopsis: isolation of 1700 kb of contiguous DNA on chromosome 5, including a 300 kb region containing the flowering-time gene CO. <i>Molecular Genetics and Genomics</i> , 1993, 239, 145-157.	2.4	50
130	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.	7.1	50
131	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.	8.9	49
132	Functional Divergence of the Arabidopsis Florigen-Interacting bZIP Transcription Factors FD and FDP. <i>Cell Reports</i> , 2020, 31, 107717.	6.4	49
133	Possible role of EARLY FLOWERING 3 (ELF3) in clock-dependent floral regulation by SHORT VEGETATIVE PHASE (SVP) in Arabidopsis thaliana. <i>New Phytologist</i> , 2009, 182, 838-850.	7.3	48
134	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.	3.5	48
135	The genetics of stamenoid petal production in oilseed rape (<i>Brassica napus</i>) and equivalent variation in Arabidopsis thaliana. <i>Theoretical and Applied Genetics</i> , 1997, 94, 731-736.	3.6	44
136	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.	2.3	44
137	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.	5.6	44
138	Ds elements on all five Arabidopsis chromosomes and assessment of their utility for transposon tagging. <i>Plant Journal</i> , 1997, 11, 145-148.	5.7	42
139	Elevated salicylic acid levels conferred by increased expression of <i>ISOCHORISMATE SYNTHASE</i> 1 contribute to hyperaccumulation of <i>SUMO</i> 1 conjugates in the Arabidopsis mutant <i>early in short days 4</i> . <i>Plant Journal</i> , 2014, 79, 206-219.	5.7	42
140	SWP73 Subunits of Arabidopsis SWI/SNF Chromatin Remodeling Complexes Play Distinct Roles in Leaf and Flower Development. <i>Plant Cell</i> , 2015, 27, 1889-1906.	6.6	42
141	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.	4.8	41
142	A rice single cell transcriptomic atlas defines the developmental trajectories of rice floret and inflorescence meristems. <i>New Phytologist</i> , 2022, 234, 494-512.	7.3	41
143	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.	2.3	38
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