

Charles P Scutt

List of Publications by Year in descending order

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Version: 2024-02-01

48
papers

1,818
citations

257357

24
h-index

276775

41
g-index

50
all docs

50
docs citations

50
times ranked

1926
citing authors

#	ARTICLE	IF	CITATIONS
1	A derived ZW chromosome system in <i>Amborella trichopoda</i> , representing the sister lineage to all other extant flowering plants. <i>New Phytologist</i> , 2022, 233, 1636-1642.	3.5	10
2	Custom methods to identify conserved genetic modules applied to novel transcriptomic data from <i>Amborella trichopoda</i> . <i>Journal of Experimental Botany</i> , 2022, 73, 2487-2498.	2.4	2
3	The Origin of Angiosperms. , 2021, , 663-682.		1
4	Immediate targets of ETTIN suggest a key role for pectin methylesterase inhibitors in the control of <i>Arabidopsis</i> gynecium development. <i>Plant Signaling and Behavior</i> , 2020, 15, 1771937.	1.2	8
5	Flowering plants return to the sea. <i>Journal of Experimental Botany</i> , 2019, 70, 4591-4593.	2.4	0
6	Transcriptomics of manually isolated <i>Amborella trichopoda</i> egg apparatus cells. <i>Plant Reproduction</i> , 2019, 32, 15-27.	1.3	16
7	Allocation of the epidermis to stomata relates to stomatal physiological control: Stomatal factors involved in the evolutionary diversification of the angiosperms and development of amphistomaty. <i>Environmental and Experimental Botany</i> , 2018, 151, 55-63.	2.0	67
8	Evidence for the Regulation of Gynoecium Morphogenesis by <i>ETTIN</i> via Cell Wall Dynamics. <i>Plant Physiology</i> , 2018, 178, 1222-1232.	2.3	25
9	Evidence for the Extensive Conservation of Mechanisms of Ovule Integument Development Since the Most Recent Common Ancestor of Living Angiosperms. <i>Frontiers in Plant Science</i> , 2018, 9, 1352.	1.7	17
10	The Origin of Angiosperms. , 2018, , 1-20.		9
11	A link between LEAFY and <i>W</i> gene homologues in <i>Welwitschia mirabilis</i> sheds light on ancestral mechanisms prefiguring floral development. <i>New Phytologist</i> , 2017, 216, 469-481.	3.5	33
12	The morphophysiological dormancy in <i>Amborella trichopoda</i> seeds is a pleisiomorphic trait in angiosperms. <i>Annals of Botany</i> , 2017, 119, mcw244.	1.4	12
13	Dioecy in <i>Amborella trichopoda</i> : evidence for genetically based sex determination and its consequences for inferences of the breeding system in early angiosperms. <i>Annals of Botany</i> , 2017, 119, mcw278.	1.4	14
14	Evolution of the YABBY gene family in seed plants. <i>Evolution & Development</i> , 2016, 18, 116-126.	1.1	87
15	The analysis of Gene Regulatory Networks in plant evo-devo. <i>Journal of Experimental Botany</i> , 2016, 67, 2549-2563.	2.4	11
16	The <i>Amborella</i> vacuolar processing enzyme family. <i>Frontiers in Plant Science</i> , 2015, 6, 618.	1.7	14
17	A Conserved Role for the NAM/miR164 Developmental Module Reveals a Common Mechanism Underlying Carpel Margin Fusion in Monocarpous and Syncarpous Eurosids. <i>Frontiers in Plant Science</i> , 2015, 6, 1239.	1.7	19
18	Current trends and future directions in flower development research. <i>Annals of Botany</i> , 2014, 114, 1399-1406.	1.4	45

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19	Evolution of the ARF Gene Family in Land Plants: Old Domains, New Tricks. <i>Molecular Biology and Evolution</i> , 2013, 30, 45-56.	3.5	196
20	A Light-Regulated Genetic Module Was Recruited to Carpel Development in <i>Arabidopsis</i> following a Structural Change to SPATULA. <i>Plant Cell</i> , 2012, 24, 2812-2825.	3.1	66
21	FILAMENTOUS FLOWER controls lateral organ development by acting as both an activator and a repressor. <i>BMC Plant Biology</i> , 2012, 12, 176.	1.6	80
22	Cabomba as a model for studies of early angiosperm evolution. <i>Annals of Botany</i> , 2011, 108, 589-598.	1.4	30
23	Insights from ANA-grade angiosperms into the early evolution of CUP-SHAPED COTYLEDON genes. <i>Annals of Botany</i> , 2011, 107, 1511-1519.	1.4	30
24	Parallel structural evolution of auxin response factors in the angiosperms. <i>Plant Journal</i> , 2010, 63, 952-959.	2.8	76
25	Carpel Development. <i>Advances in Botanical Research</i> , 2010, 55, 1-73.	0.5	65
26	The evolutionary-developmental analysis of plant microRNAs. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2010, 365, 469-476.	1.8	30
27	The analysis of entire gene promoters by surface plasmon resonance. <i>Plant Journal</i> , 2009, 59, 851-858.	2.8	15
28	Functional Conservation between CRABS CLAW Orthologues from Widely Diverged Angiosperms. <i>Annals of Botany</i> , 2007, 100, 651-657.	1.4	49
29	The Evolution of Plant Development: Past, Present and Future: Preface. <i>Annals of Botany</i> , 2007, 100, 599-601.	1.4	4
30	An evolutionary perspective on the regulation of carpel development. <i>Journal of Experimental Botany</i> , 2006, 57, 2143-2152.	2.4	75
31	Analysis of members of the <i>Silene latifolia</i> Cys2/His2 zinc-finger transcription factor family during dioecious flower development and in a novel stamen-defective mutant <i>ssf1</i> . <i>Planta</i> , 2005, 220, 559-571.	1.6	19
32	Evidence that CRABS CLAW and TOUSLED have conserved their roles in carpel development since the ancestor of the extant angiosperms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 4649-4654.	3.3	118
33	Molecular approaches to the study of sex determination in dioecious <i>Silene latifolia</i> . , 2004, , 51-71.		0
34	The Identification of Candidate Genes for a Reverse Genetic Analysis of Development and Function in the <i>Arabidopsis</i> Gynoecium. <i>Plant Physiology</i> , 2003, 132, 653-665.	2.3	31
35	Male Specific Genes from Dioecious White Campion Identified by Fluorescent Differential Display. <i>Plant and Cell Physiology</i> , 2002, 43, 563-572.	1.5	21
36	Techniques for the removal of marker genes from transgenic plants. <i>Biochimie</i> , 2002, 84, 1119-1126.	1.3	38

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37	Intrachromosomal recombination between attP regions as a tool to remove selectable marker genes from tobacco transgenes. <i>Nature Biotechnology</i> , 2000, 18, 442-445.	9.4	151
38	Morphological and molecular analysis of a double-flowered mutant of the dioecious plant white campion showing both meristic and homeotic effects. , 1999, 25, 267-279.		8
39	The Men-10 cDNA encodes a novel form of proline-rich protein expressed in the tapetum of dioecious <i>Silene latifolia</i> . <i>Sexual Plant Reproduction</i> , 1998, 11, 236-240.	2.2	9
40	Laser isolation of plant sex chromosomes: studies on the DNA composition of the X and Y sex chromosomes of <i>Silene latifolia</i> . <i>Genome</i> , 1997, 40, 705-715.	0.9	53
41	Sex Determination in Dioecious <i>Silene latifolia</i> (Effects of the Y Chromosome and the Parasitic Smut) <i>Tj ETQq1 1 0.784314 rgBT /Over</i> 114, 969-979.	2.3	57
42	High-Stringency Subtraction for the Identification of Differentially Regulated cDNA Clones. <i>BioTechniques</i> , 1997, 23, 468-474.	0.8	8
43	Spatial expression dynamics of Men-9 delineate the third floral whorl in male and female flowers of dioecious <i>Silene latifolia</i> . <i>Plant Journal</i> , 1997, 12, 155-168.	2.8	30
44	Differential Screening. , 1997, , 1-22.		6
45	Cloning of PCP1, a member of a family of pollen coat protein (PCP) genes from <i>Brassica oleracea</i> encoding novel cysteine-rich proteins involved in pollen-stigma interactions. <i>Plant Journal</i> , 1996, 10, 303-313.	2.8	54
46	Okadaic acid causes breakdown of self-incompatibility in <i>Brassica oleracea</i> : evidence for the involvement of protein phosphatases in the incompatible response. <i>Sexual Plant Reproduction</i> , 1993, 6, 282.	2.2	25
47	An S5 self-incompatibility allele-specific cDNA sequence from <i>Brassica oleracea</i> shows high homology to the SLR2 gene. <i>Molecular Genetics and Genomics</i> , 1992, 232, 240-246.	2.4	38
48	A cDNA encoding an S-locus specific glycoprotein from <i>Brassica oleracea</i> plants containing the S5 self-incompatibility allele. <i>Molecular Genetics and Genomics</i> , 1990, 220, 409-413.	2.4	44