John L Bowman

List of Publications by Year
in descending order

140 papers	19,271 citations	10779 66 h-index	8358 138 g-index
164 all docs	164 docs citations	164 times ranked	14049 citing authors

#	Article	IF	Citations
1	Hormonal and genetic control of pluripotency in bryophyte model systems. Current Opinion in Plant Biology, 2024, 77, 102486.	7.3	5
2	Reflections on the ABC model of flower development. Plant Cell, 2024, 36, 1334-1357.	7.6	16
3	The landscape of transcription factor promoter activity during vegetative development in Marchantia. Plant Cell, 2024, 36, 2140-2159.	7.6	6
4	Dual Regulation of Cytochrome P450 Gene Expression by Two Distinct Small RNAs, a Novel tasiRNA and miRNA, in <i>Marchantia polymorpha</i>. Plant and Cell Physiology, 2024, 65, 1115-1134.	3.5	1
5	The auronidin flavonoid pigments of the liverwort <i>Marchantia polymorpha</i> form polymers that modify cell wall properties. Plant Journal, 2024, 120, 1159-1175.	6.1	0
6	Control of sporophyte secondary cell wall development in Marchantia by a Class II KNOX gene. Current Biology, 2024, 34, 5213-5222.e5.	3.9	1
7	<scp>PINâ€FORMED</scp> is required for shoot phototropism/gravitropism and facilitates meristem formation in <i>Marchantia polymorpha</i>. New Phytologist, 2023, 238, 1498-1515.	8.2	7
8	Green land: Multiple perspectives on green algal evolution and the earliest land plants. American Journal of Botany, 2023, 110, .	2.2	23
9	The fate of sex chromosomes during the evolution of monoicy from dioicy in liverworts. Current Biology, 2023, 33, 3597-3609.e3.	3.9	12
10	The liverwort Marchantia polymorpha, a model for all ages. Current Topics in Developmental Biology, 2022, , 1-32.	0.0	16
11	KANADI promotes thallus differentiation and FRâ€induced gametangiophore formation in the liverwort <i>Marchantia</i>. New Phytologist, 2022, 234, 1377-1393.	8.2	9
12	Stress, senescence, and specialized metabolites in bryophytes. Journal of Experimental Botany, 2022, 73, 4396-4411.	5.1	16
13	The nature of nurture: the conserved role of tapetalâ€like cells in sporogenesis between mosses and angiosperms. New Phytologist, 2022, , .	8.2	1
14	The Evolution of Complex Multicellularity in Streptophytes. , 2022, , 325-342.		0
15	<i>CLASS-II KNOX</i>genes coordinate spatial and temporal ripening in tomato. Plant Physiology, 2022, 190, 657-668.	5.4	13
16	The renaissance and enlightenment of<i>Marchantia</i>as a model system. Plant Cell, 2022, 34, 3512-3542.	7.6	52
17	MarpolBase Expression: A Web-Based, Comprehensive Platform for Visualization and Analysis of Transcriptomes in the Liverwort <i>Marchantia polymorpha</i>. Plant and Cell Physiology, 2022, 63, 1745-1755.	3.5	22
18	A transporter of 1â€aminocyclopropaneâ€1â€carboxylic acid affects thallus growth and fertility in <i>Marchantia polymorpha</i>. New Phytologist, 2022, 236, 2103-2114.	8.2	5

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19	The single <i>Marchantia polymorpha FERONIA</i> homolog reveals an ancestral role in regulating cellular expansion and integrity. Development (Cambridge), 2022, 149, .	3.0	10
20	The origin of a land flora. Nature Plants, 2022, 8, 1352-1369.	7.0	50
21	On the Evolutionary Origins of Land Plant Auxin Biology. Cold Spring Harbor Perspectives in Biology, 2021, 13, a040048.	7.4	10
22	DEFECTIVE EMBRYO AND MERISTEMS genes are required for cell division and gamete viability in Arabidopsis. PLoS Genetics, 2021, 17, e1009561.	3.3	4
23	Author response: Gamete expression of TALE class HD genes activates the diploid sporophyte program in Marchantia polymorpha. , 2021, , .		0
24	Rates and patterns of molecular evolution in bryophyte genomes, with focus on complex thalloid liverworts, Marchantiopsida. Molecular Phylogenetics and Evolution, 2021, 165, 107295.	2.9	14
25	Identification of the sex-determining factor in the liverwort Marchantia polymorpha reveals unique evolution of sex chromosomes in a haploid system. Current Biology, 2021, 31, 5522-5532.e7.	3.9	40
26	Transcriptional and Morpho-Physiological Responses of Marchantia polymorpha upon Phosphate Starvation. International Journal of Molecular Sciences, 2020, 21, 8354.	4.5	22
27	Ethylene-independent functions of the ethylene precursor ACC in Marchantia polymorpha. Nature Plants, 2020, 6, 1335-1344.	7.0	49
28	Oil Body Formation in Marchantia polymorpha Is Controlled by MpC1HDZ and Serves as a Defense against Arthropod Herbivores. Current Biology, 2020, 30, 2815-2828.e8.	3.9	58
29	Chromatin Organization in Early Land Plants Reveals an Ancestral Association between H3K27me3, Transposons, and Constitutive Heterochromatin. Current Biology, 2020, 30, 573-588.e7.	3.9	133
30	The Evolution of Flavonoid Biosynthesis: A Bryophyte Perspective. Frontiers in Plant Science, 2020, 11, .	4.2	150
31	Cellulose Synthesis â€“ Central Components and Their Evolutionary Relationships. Trends in Plant Science, 2019, 24, 402-412.	15.4	63
32	Control of proliferation in the haploid meristem by CLE peptide signaling in Marchantia polymorpha. PLoS Genetics, 2019, 15, e1007997.	3.3	56
33	Evolution and co-option of developmental regulatory networks in early land plants. Current Topics in Developmental Biology, 2019, , 35-53.	0.0	28
34	Something ancient and something neofunctionalizedâ€"evolution of land plant hormone signaling pathways. Current Opinion in Plant Biology, 2019, 47, 64-72.	7.3	50
35	Class C <scp>ARF</scp>s evolved before the origin of land plants and antagonize differentiation and developmental transitions in <i>Marchantia polymorpha</i>. New Phytologist, 2018, 218, 1612-1630.	8.2	67
36	Evolutionary history of <scp>HOMEODOMAIN LEUCINE ZIPPER</scp> transcription factors during plant transition to land. New Phytologist, 2018, 219, 408-421.	8.2	29

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37	Genetic analysis of the liverwort <i>Marchantia polymorpha</i> reveals that R2R3<scp>MYB</scp> activation of flavonoid production in response to abiotic stress is an ancient character in land plants. New Phytologist, 2018, 218, 554-566.	8.2	101
38	Terpenoid Secondary Metabolites in Bryophytes: Chemical Diversity, Biosynthesis and Biological Functions. Critical Reviews in Plant Sciences, 2018, 37, 210-231.	5.6	64
39	Co-expression and Transcriptome Analysis of Marchantia polymorpha Transcription Factors Supports Class C ARFs as Independent Actors of an Ancient Auxin Regulatory Module. Frontiers in Plant Science, 2018, 9, .	4.2	39
40	Micro<scp>RNA</scp>s in <i>Marchantia polymorpha</i>. New Phytologist, 2018, 220, 409-416.	8.2	20
41	3D Body Evolution: Adding a New Dimension to Colonize the Land. Current Biology, 2018, 28, R838-R840.	3.9	1
42	UVR8â€mediated induction of flavonoid biosynthesis for UVB tolerance is conserved between the liverwort <i>Marchantia polymorpha</i> and flowering plants. Plant Journal, 2018, 96, 503-517.	6.1	99
43	Extensive epigenetic reprogramming during the life cycle of Marchantia polymorpha. Genome Biology, 2018, 19, .	8.4	45
44	Marchantia liverworts as a proxy to plantsâ€™ basal microbiomes. Scientific Reports, 2018, 8, .	3.7	47
45	The KNOXI Transcription Factor SHOOT MERISTEMLESS Regulates Floral Fate in Arabidopsis. Plant Cell, 2018, 30, 1309-1321.	7.6	26
46	Insights into Land Plant Evolution Garnered from the Marchantia polymorpha Genome. Cell, 2017, 171, 287-304.e15.	35.1	837
47	A Genetic Screen for Impaired Systemic RNAi Highlights the Crucial Role of DICER-LIKE 2. Plant Physiology, 2017, 175, 1424-1437.	5.4	66
48	Evolution of the Metabolic Network Leading to Ascorbate Synthesis and Degradation Using Marchantia polymorpha as a Model System. , 2017, , 417-430.		1
49	Evolution of the YABBY gene family in seed plants. Evolution & Development, 2016, 18, 116-126.	2.0	68
50	Evolution in the Cycles of Life. Annual Review of Genetics, 2016, 50, 133-154.	7.7	86
51	Field Guide to Plant Model Systems. Cell, 2016, 167, 325-339.	35.1	89
52	Microbial-type terpene synthase genes occur widely in nonseed land plants, but not in seed plants. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12328-12333.	7.7	75
53	Molecular Diversity of Terpene Synthases in the Liverwort Marchantia polymorpha. Plant Cell, 2016, , tpc.00062.2016.	7.6	44
54	A Brief History of<i>Marchantia</i>from Greece to Genomics. Plant and Cell Physiology, 2016, 57, 210-229.	3.5	67

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55	Efficient and Inducible Use of Artificial MicroRNAs in<i>Marchantia polymorpha</i>. Plant and Cell Physiology, 2016, 57, 281-290.	3.5	64
56	Class III HD-Zip activity coordinates leaf development in Physcomitrella patens. Developmental Biology, 2016, 419, 184-197.	1.9	47
57	The Naming of Names: Guidelines for Gene Nomenclature in<i>Marchantia</i>. Plant and Cell Physiology, 2016, 57, 257-261.	3.5	51
58	Identification of miRNAs and Their Targets in the Liverwort<i>Marchantia polymorpha</i>by Integrating RNA-Seq and Degradome Analyses. Plant and Cell Physiology, 2016, 57, 339-358.	3.5	51
59	<i>Marchantia</i>: Past, Present and Future. Plant and Cell Physiology, 2016, 57, 205-209.	3.5	40
60	Marchantia. Current Biology, 2016, 26, R186-R187.	3.9	14
61	Profiling and Characterization of Small RNAs in the Liverwort,<i>Marchantia polymorpha</i>, Belonging to the First Diverged Land Plants. Plant and Cell Physiology, 2016, 57, 359-372.	3.5	47
62	Author response: Active suppression of a leaf meristem orchestrates determinate leaf growth. , 2016, , .		1
63	A Role of TDIF Peptide Signaling in Vascular Cell Differentiation is Conserved Among Euphyllophytes. Frontiers in Plant Science, 2015, 6, .	4.2	37
64	Auxin Produced by the Indole-3-Pyruvic Acid Pathway Regulates Development and Gemmae Dormancy in the Liverwort <i>Marchantia polymorpha</i>. Plant Cell, 2015, 27, 1650-1669.	7.6	119
65	Antagonistic Roles for KNOX1 and KNOX2 Genes in Patterning the Land Plant Body Plan Following an Ancient Gene Duplication. PLoS Genetics, 2015, 11, e1004980.	3.3	121
66	Comparative Analysis of the Conserved Functions of Arabidopsis DRL1 and Yeast KTI12. Molecules and Cells, 2015, 38, 243-250.	5.0	10
67	Auxin-Mediated Transcriptional System with a Minimal Set of Components Is Critical for Morphogenesis through the Life Cycle in Marchantia polymorpha. PLoS Genetics, 2015, 11, e1005084.	3.3	119
68	A Simple Auxin Transcriptional Response System Regulates Multiple Morphogenetic Processes in the Liverwort Marchantia polymorpha. PLoS Genetics, 2015, 11, e1005207.	3.3	143
69	Origin of a novel regulatory module by duplication and degeneration of an ancient plant transcription factor. Molecular Phylogenetics and Evolution, 2014, 81, 159-173.	2.9	11
70	Flower Development: Open Questions and Future Directions. Methods in Molecular Biology, 2014, , 103-124.	0.0	25
71	From cell to organism across space and time. Current Opinion in Plant Biology, 2013, 16, 542-544.	7.3	0
72	Walkabout on the long branches of plant evolution. Current Opinion in Plant Biology, 2013, 16, 70-77.	7.3	77

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73	My favourite flowering image. Journal of Experimental Botany, 2013, 64, 5779-5782.	5.1	1
74	Evolution of the Class IV HD-Zip Gene Family in Streptophytes. Molecular Biology and Evolution, 2013, 30, 2347-2365.	4.7	31
75	Genome-Wide Identification of KANADI1 Target Genes. PLoS ONE, 2013, 8, e77341.	2.5	58
76	Stomata: Active Portals for Flourishing on Land. Current Biology, 2011, 21, R540-R541.	3.9	13
77	Arabidopsis Homologs of the<i>Petunia</i>Â<i>HAIRY MERISTEM</i>Gene Are Required for Maintenance of Shoot and Root Indeterminacy Â Â Â. Plant Physiology, 2011, 155, 735-750.	5.4	108
78	Cell signalling by microRNA165/6 directs gene dose-dependent root cell fate. Nature, 2010, 465, 316-321.	40.1	691
79	Interplay of auxin, KANADI and Class III HD-ZIP transcription factors in vascular tissue formation. Development (Cambridge), 2010, 137, 975-984.	3.0	177
80	Differentiating Arabidopsis Shoots from Leaves by Combined YABBY Activities Â. Plant Cell, 2010, 22, 2113-2130.	7.6	226
81	Criteria for Annotation of Plant MicroRNAs. Plant Cell, 2009, 20, 3186-3190.	7.6	1,046
82	The<i>NGATHA</i>Distal Organ Development Genes Are Essential for Style Specification in<i>Arabidopsis</i>Â. Plant Cell, 2009, 21, 1373-1393.	7.6	108
83	The flowering hormone florigen functions as a general systemic regulator of growth and termination. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 8392-8397.	7.7	277
84	Auxin-Dependent Patterning and Gamete Specification in the <i>Arabidopsis</i> Female Gametophyte. Science, 2009, 324, 1684-1689.	38.2	238
85	Gene expression patterns in seed plant shoot meristems and leaves: homoplasy or homology?. Journal of Plant Research, 2009, 123, 43-55.	2.0	64
86	Evolution of plant microRNAs and their targets. Trends in Plant Science, 2008, 13, 343-349.	15.4	356
87	Patterning and Polarity in Seed Plant Shoots. Annual Review of Plant Biology, 2008, 59, 67-88.	24.7	104
88	Activity Range of Arabidopsis Small RNAs Derived from Different Biogenesis Pathways Â. Plant Physiology, 2008, 147, 58-62.	5.4	44
89	Signals Derived from<i>YABBY</i>Gene Activities in Organ Primordia Regulate Growth and Partitioning of<i>Arabidopsis</i>Shoot Apical Meristems. Plant Cell, 2008, 20, 1217-1230.	7.6	135
90	<i>REBELOTE</i>,<i>SQUINT</i>, and<i>ULTRAPETALA1</i>Function Redundantly in the Temporal Regulation of Floral Meristem Termination in<i>Arabidopsis thaliana</i>Â. Plant Cell, 2008, 20, 901-919.	7.6	88

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91	The Ancestral Developmental Tool Kit of Land Plants. International Journal of Plant Sciences, 2007, 168, 1-35.	1.4	223
92	Green Genesâ€"Comparative Genomics of the Green Branch of Life. Cell, 2007, 129, 229-234.	35.1	171
93	KANADI and Class III HD-Zip Gene Families Regulate Embryo Patterning and Modulate Auxin Flow during Embryogenesis in Arabidopsis. Plant Cell, 2007, 19, 495-508.	7.6	195
94	Freezing and desiccation tolerance in the moss Physcomitrella patens: An in situ Fourier transform infrared spectroscopic study. Biochimica Et Biophysica Acta - General Subjects, 2006, 1760, 1226-1234.	2.0	60
95	ABERRANT TESTA SHAPE encodes a KANADI family member, linking polarity determination to separation and growth of Arabidopsis ovule integuments. Plant Journal, 2006, 46, 522-531.	6.1	121
96	Evolution of Class III Homeodomainâ€"Leucine Zipper Genes in Streptophytes. Genetics, 2006, 173, 373-388.	4.2	122
97	Recruitment of CRABS CLAW to promote nectary development within the eudicot clade. Development (Cambridge), 2005, 132, 5021-5032.	3.0	154
98	Multiple Protein Regions Contribute to Differential Activities of YABBY Proteins inReproductive Development. Plant Physiology, 2005, 137, 651-662.	5.4	28
99	Activation of CRABS CLAW in the Nectaries and Carpels of Arabidopsis. Plant Cell, 2005, 17, 25-36.	7.6	125
100	Roles for Class III HD-Zip and KANADI Genes in Arabidopsis Root Development. Plant Physiology, 2004, 135, 2261-2270.	5.4	133
101	Molecular evidence for bicontinental hybridogenous genomic constitution in <i>Lepidium</i> sensu stricto (Brassicaceae) species from Australia and New Zealand. American Journal of Botany, 2004, 91, 254-261.	2.2	117
102	Promoter Bashing, microRNAs, and Knox Genes. New Insights, Regulators, and Targets-of-Regulation in the Establishment of Lateral Organ Polarity in Arabidopsis. Plant Physiology, 2004, 135, 685-694.	5.4	44
103	The Arabidopsis thaliana SNF2 homolog AtBRM controls shoot development and flowering. Development (Cambridge), 2004, 131, 4965-4975.	3.0	147
104	Ancient microRNA target sequences in plants. Nature, 2004, 428, 485-486.	40.1	347
105	Class III HD-Zip gene regulation, the golden fleece of ARGONAUTE activity?. BioEssays, 2004, 26, 938-942.	2.3	55
106	Asymmetric leaf development and blade expansion in Arabidopsisare mediated by KANADI and YABBY activities. Development (Cambridge), 2004, 131, 2997-3006.	3.0	333
107	Radial Patterning of Arabidopsis Shoots by Class III HD-ZIP and KANADI Genes. Current Biology, 2003, 13, 1768-1774.	3.9	926
108	Plant genetics: a decade of integration. Nature Genetics, 2003, 33, 294-304.	16.3	30

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109	Allopolyploidization and evolution of species with reduced floral structures in Lepidium L. (Brassicaceae). Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 16835-16840.	7.7	63
110	A Surveillance System Regulates Selective Entry of RNA into the Shoot Apex. Plant Cell, 2002, 14, 1497-1508.	7.6	139
111	YABBY Polarity Genes Mediate the Repression of KNOX Homeobox Genes in Arabidopsis. Plant Cell, 2002, 14, 2761-2770.	7.6	204
112	Establishment of polarity in angiosperm lateral organs. Trends in Genetics, 2002, 18, 134-141.	13.0	249
113	Turning floral organs into leaves, leaves into floral organs. Current Opinion in Genetics and Development, 2001, 11, 449-456.	3.4	127
114	Establishment of polarity in lateral organs of plants. Current Biology, 2001, 11, 1251-1260.	3.9	542
115	Chloroplast DNA phylogeny and biogeography of Lepidium (Brassicaceae). American Journal of Botany, 2001, 88, 2051-2063.	2.2	119
116	The<i>Arabidopsis</i>nectary is an ABC-independent floral structure. Development (Cambridge), 2001, 128, 4657-4667.	3.0	80
117	SHATTERPROOF MADS-box genes control seed dispersal in Arabidopsis. Nature, 2000, 404, 766-770.	40.1	747
118	The YABBY gene family and abaxial cell fate. Current Opinion in Plant Biology, 2000, 3, 17-22.	7.3	218
119	Formation and maintenance of the shoot apical meristem. Trends in Plant Science, 2000, 5, 110-115.	15.4	178
120	Axial patterning in leaves and other lateral organs. Current Opinion in Genetics and Development, 2000, 10, 399-404.	3.4	30
121	Evolutionary Changes in Floral Structure withinLepidiumL. (Brassicaceae). International Journal of Plant Sciences, 1999, 160, 917-929.	1.4	57
122	4 Molecular Genetics of Gynoecium Development in Arabidopsis. Current Topics in Developmental Biology, 1999, , 155-205.	0.0	132
123	Distinct Mechanisms Promote Polarity Establishment in Carpels of Arabidopsis. Cell, 1999, 99, 199-209.	35.1	329
124	Patterns of Petal and Stamen Reduction in Australian Species of Lepidium L. (Brassicaceae). International Journal of Plant Sciences, 1998, 159, 65-74.	1.4	31
125	Evolutionary conservation of angiosperm flower development at the molecular and genetic levels. Journal of Biosciences, 1997, 22, 515-527.	1.5	112
126	Manipulating floral organ identity. Current Biology, 1993, 3, 90-93.	3.9	9

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127	Control of flower development in <i>Arabidopsis thaliana</i> by <i>APETALA1</i> and interacting genes. Development (Cambridge), 1993, 119, 721-743.	3.0	577
128	Vectors for plant transformation and cosmid libraries. Gene, 1992, 117, 161-167.	2.4	21
129	Manipulation of flower structure in transgenic tobacco. Cell, 1992, 71, 133-143.	35.1	224
130	Negative regulation of the Arabidopsis homeotic gene AGAMOUS by the APETALA2 product. Cell, 1991, 65, 991-1002.	35.1	600
131	Expression of the Arabidopsis Floral Homeotic Gene AGAMOUS Is Restricted to Specific Cell Types Late in Flower Development. Plant Cell, 1991, 3, 749.	7.6	48
132	A genetic and molecular model for flower development in <i>Arabidopsis thaliana</i>. Development (Cambridge), 1991, 113, 157-167.	3.0	132
133	The protein encoded by the Arabidopsis homeotic gene agamous resembles transcription factors. Nature, 1990, 346, 35-39.	40.1	1,416
134	Genes Directing Flower Development in Arabidopsis. Plant Cell, 1989, 1, 37.	7.6	217
135	MicroRNAs: Micro-managing the Plant Genome. , 0, , 244-278.		3
136	Active suppression of a leaf meristem orchestrates determinate leaf growth. ELife, 0, 5, .	1.6	120
137	Establishment of Polarity in Lateral Organs of Seed Plants. , 0, , 288-316.		0
138	<scp>MicroRNAs</scp>: Microâ€Managing the Plant Genome. , 0, , 244-278.		0
139	Gamete expression of TALE class HD genes activates the diploid sporophyte program in Marchantia polymorpha. ELife, 0, 10, .	1.6	39
140	Phosphate Starvation Triggers Transcriptional Changes in the Biosynthesis and Signaling Pathways of Phytohormones in Marchantia polymorphaÂ. Biology and Life Sciences Forum, 0, 90, 89.	0.0	1