

Philip N Benfey

List of Publications by Year in descending order

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191
papers

27,111
citations

7096

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157
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294
all docs

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docs citations

294
times ranked

18333
citing authors

#	ARTICLE	IF	CITATIONS
1	An Auxin-Dependent Distal Organizer of Pattern and Polarity in the Arabidopsis Root. <i>Cell</i> , 1999, 99, 463-472.	28.9	1,233
2	A Gene Expression Map of the Arabidopsis Root. <i>Science</i> , 2003, 302, 1956-1960.	12.6	1,161
3	A High-Resolution Root Spatiotemporal Map Reveals Dominant Expression Patterns. <i>Science</i> , 2007, 318, 801-806.	12.6	1,048
4	The SHORT-ROOT Gene Controls Radial Patterning of the Arabidopsis Root through Radial Signaling. <i>Cell</i> , 2000, 101, 555-567.	28.9	1,007
5	The SCARECROW Gene Regulates an Asymmetric Cell Division That Is Essential for Generating the Radial Organization of the Arabidopsis Root. <i>Cell</i> , 1996, 86, 423-433.	28.9	998
6	Transcriptional Regulation of ROS Controls Transition from Proliferation to Differentiation in the Root. <i>Cell</i> , 2010, 143, 606-616.	28.9	926
7	Intercellular movement of the putative transcription factor SHR in root patterning. <i>Nature</i> , 2001, 413, 307-311.	27.8	766
8	Cell signalling by microRNA165/6 directs gene dose-dependent root cell fate. <i>Nature</i> , 2010, 465, 316-321.	27.8	739
9	The auxin influx carrier LAX3 promotes lateral root emergence. <i>Nature Cell Biology</i> , 2008, 10, 946-954.	10.3	715
10	Cell Identity Mediates the Response of <i>Arabidopsis</i> Roots to Abiotic Stress. <i>Science</i> , 2008, 320, 942-945.	12.6	700
11	The GRAS gene family in Arabidopsis: sequence characterization and basic expression analysis of the SCARECROW-LIKE genes. <i>Plant Journal</i> , 1999, 18, 111-119.	5.7	572
12	The bHLH Transcription Factor POPEYE Regulates Response to Iron Deficiency in <i>Arabidopsis</i> Roots. <i>Plant Cell</i> , 2010, 22, 2219-2236.	6.6	561
13	Control of <i>Arabidopsis</i> Root Development. <i>Annual Review of Plant Biology</i> , 2012, 63, 563-590.	18.7	558
14	Oscillating Gene Expression Determines Competence for Periodic <i>Arabidopsis</i> Root Branching. <i>Science</i> , 2010, 329, 1306-1311.	12.6	532
15	An Evolutionarily Conserved Mechanism Delimiting SHR Movement Defines a Single Layer of Endodermis in Plants. <i>Science</i> , 2007, 316, 421-425.	12.6	522
16	A novel two-component hybrid molecule regulates vascular morphogenesis of the Arabidopsis root. <i>Genes and Development</i> , 2000, 14, 2938-2943.	5.9	499
17	An Auxin Gradient and Maximum in the <i>Arabidopsis</i> Root Apex Shown by High-Resolution Cell-Specific Analysis of IAA Distribution and Synthesis. <i>Plant Cell</i> , 2009, 21, 1659-1668.	6.6	439
18	Regulation of plant root system architecture: implications for crop advancement. <i>Current Opinion in Biotechnology</i> , 2015, 32, 93-98.	6.6	351

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19	Receptor-Like Kinase ACR4 Restricts Formative Cell Divisions in the <i>Arabidopsis</i> Root. <i>Science</i> , 2008, 322, 594-597.	12.6	342
20	COBRA encodes a putative GPI-anchored protein, which is polarly localized and necessary for oriented cell expansion in <i>Arabidopsis</i> . <i>Genes and Development</i> , 2001, 15, 1115-1127.	5.9	335
21	Genetic evidence that the endodermis is essential for shoot gravitropism in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 1998, 14, 425-430.	5.7	334
22	COBRA, an <i>Arabidopsis</i> Extracellular Glycosyl-Phosphatidyl Inositol-Anchored Protein, Specifically Controls Highly Anisotropic Expansion through Its Involvement in Cellulose Microfibril Orientation. <i>Plant Cell</i> , 2005, 17, 1749-1763.	6.6	321
23	Imaging and Analysis Platform for Automatic Phenotyping and Trait Ranking of Plant Root Systems. <i>Plant Physiology</i> , 2010, 152, 1148-1157.	4.8	306
24	A Gene Regulatory Network for Root Epidermis Cell Differentiation in <i>Arabidopsis</i> . <i>PLoS Genetics</i> , 2012, 8, e1002446.	3.5	306
25	Transcriptional Profile of the <i>Arabidopsis</i> Root Quiescent Center. <i>Plant Cell</i> , 2005, 17, 1908-1925.	6.6	288
26	Whole-Genome Analysis of the SHORT-ROOT Developmental Pathway in <i>Arabidopsis</i> . <i>PLoS Biology</i> , 2006, 4, e143.	5.6	283
27	GiA Roots: software for the high throughput analysis of plant root system architecture. <i>BMC Plant Biology</i> , 2012, 12, 116.	3.6	279
28	Cell type-specific expression profiling in plants via cell sorting of protoplasts from fluorescent reporter lines. <i>Nature Methods</i> , 2005, 2, 615-619.	19.0	276
29	A Bistable Circuit Involving SCARECROW-RETINOBLASTOMA Integrates Cues to Inform Asymmetric Stem Cell Division. <i>Cell</i> , 2012, 150, 1002-1015.	28.9	273
30	Bimodular auxin response controls organogenesis in <i>Arabidopsis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 2705-2710.	7.1	271
31	3D phenotyping and quantitative trait locus mapping identify core regions of the rice genome controlling root architecture. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E1695-704.	7.1	261
32	Transcriptional and posttranscriptional regulation of transcription factor expression in <i>Arabidopsis</i> roots. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 6055-6060.	7.1	257
33	High-Resolution Expression Map of the <i>Arabidopsis</i> Root Reveals Alternative Splicing and lincRNA Regulation. <i>Developmental Cell</i> , 2016, 39, 508-522.	7.0	245
34	From lab to field, new approaches to phenotyping root system architecture. <i>Current Opinion in Plant Biology</i> , 2011, 14, 310-317.	7.1	235
35	Super-resolution ribosome profiling reveals unannotated translation events in <i>Arabidopsis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E7126-E7135.	7.1	222
36	Mechanisms Regulating SHORT-ROOT Intercellular Movement. <i>Current Biology</i> , 2004, 14, 1847-1851.	3.9	203

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37	Auxin minimum triggers the developmental switch from cell division to cell differentiation in the <i>Arabidopsis</i> root. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7641-E7649.	7.1	193
38	Cell Identity Regulators Link Development and Stress Responses in the <i>Arabidopsis</i> Root. Developmental Cell, 2011, 21, 770-782.	7.0	178
39	From Genotype to Phenotype: Systems Biology Meets Natural Variation. Science, 2008, 320, 495-497.	12.6	170
40	Regulation of Plant Stem Cell Quiescence by a Brassinosteroid Signaling Module. Developmental Cell, 2014, 30, 36-47.	7.0	164
41	Unique cell-type-specific patterns of DNA methylation in the root meristem. Nature Plants, 2016, 2, 16058.	9.3	159
42	Down and out in <i>Arabidopsis</i> : the formation of lateral roots. Trends in Plant Science, 1997, 2, 390-396.	8.8	149
43	A stele-enriched gene regulatory network in the <i>Arabidopsis</i> root. Molecular Systems Biology, 2011, 7, 459.	7.2	145
44	MYB36 regulates the transition from proliferation to differentiation in the <i>Arabidopsis</i> root. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 12099-12104.	7.1	145
45	The COBRA Family of Putative GPI-Anchored Proteins in <i>Arabidopsis</i> . A New Fellowship in Expansion. Plant Physiology, 2002, 130, 538-548.	4.8	143
46	The protein expression landscape of the <i>Arabidopsis</i> root. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 6811-6818.	7.1	140
47	High-resolution experimental and computational profiling of tissue-specific known and novel miRNAs in <i>Arabidopsis</i> . Genome Research, 2012, 22, 163-176.	5.5	140
48	Periodic root branching in <i>Arabidopsis</i> requires synthesis of an uncharacterized carotenoid derivative. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E1300-9.	7.1	139
49	Root development. Current Biology, 2000, 10, R813-R815.	3.9	138
50	To branch or not to branch: the role of pre-patterning in lateral root formation. Development (Cambridge), 2013, 140, 4301-4310.	2.5	137
51	Plant Stem Cell Niches: Standing the Test of Time. Cell, 2008, 132, 553-557.	28.9	132
52	Omics meet networks—using systems approaches to infer regulatory networks in plants. Current Opinion in Plant Biology, 2010, 13, 126-131.	7.1	132
53	High-resolution metabolic mapping of cell types in plant roots. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E1232-41.	7.1	131
54	Transcriptional control of tissue formation throughout root development. Science, 2015, 350, 426-430.	12.6	128

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55	RGF1 controls root meristem size through ROS signalling. <i>Nature</i> , 2020, 577, 85-88.	27.8	128
56	Combining Expression and Comparative Evolutionary Analysis. The COBRA Gene Family. <i>Plant Physiology</i> , 2007, 143, 172-187.	4.8	125
57	Analyzing Lateral Root Development: How to Move Forward. <i>Plant Cell</i> , 2012, 24, 15-20.	6.6	125
58	MicroRNA miR396 Regulates the Switch between Stem Cells and Transit-Amplifying Cells in Arabidopsis Roots. <i>Plant Cell</i> , 2015, 27, 3354-3366.	6.6	125
59	A broad competence to respond to SHORT ROOT revealed by tissue-specific ectopic expression. <i>Development (Cambridge)</i> , 2004, 131, 2817-2826.	2.5	124
60	Genotypic recognition and spatial responses by rice roots. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 2670-2675.	7.1	124
61	The AUXIN BINDING PROTEIN 1 Is Required for Differential Auxin Responses Mediating Root Growth. <i>PLoS ONE</i> , 2009, 4, e6648.	2.5	124
62	Both the conserved GRAS domain and nuclear localization are required for SHORT-ROOT movement. <i>Plant Journal</i> , 2009, 57, 785-797.	5.7	123
63	Intercellular Communication during Plant Development. <i>Plant Cell</i> , 2011, 23, 855-864.	6.6	119
64	ASYMMETRIC CELL DIVISION IN PLANTS. <i>Annual Review of Plant Biology</i> , 1999, 50, 505-537.	14.8	117
65	Protonophore- and pH-insensitive glucose and sucrose accumulation detected by FRET nanosensors in Arabidopsis root tips. <i>Plant Journal</i> , 2008, 56, 948-962.	5.7	116
66	Spatial Coordination between Stem Cell Activity and Cell Differentiation in the Root Meristem. <i>Developmental Cell</i> , 2013, 26, 405-415.	7.0	113
67	Paired-End Analysis of Transcription Start Sites in <i>Arabidopsis</i> Reveals Plant-Specific Promoter Signatures. <i>Plant Cell</i> , 2014, 26, 2746-2760.	6.6	112
68	Î²-Cyclocitral is a conserved root growth regulator. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 10563-10567.	7.1	111
69	Genes and networks regulating root anatomy and architecture. <i>New Phytologist</i> , 2015, 208, 26-38.	7.3	108
70	A single-cell Arabidopsis root atlas reveals developmental trajectories in wild-type and cell identity mutants. <i>Developmental Cell</i> , 2022, 57, 543-560.e9.	7.0	106
71	Maturation of the Ground Tissue of the Root Is Regulated by Gibberellin and SCARECROW and Requires SHORT-ROOT. <i>Plant Physiology</i> , 2005, 138, 636-640.	4.8	103
72	Interplay between SCARECROW, GA and LIKE HETEROCHROMATIN PROTEIN 1 in ground tissue patterning in the Arabidopsis root. <i>Plant Journal</i> , 2009, 58, 1016-1027.	5.7	103

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73	The Genetic and Molecular Basis of Root Development. Annual Review of Plant Biology, 1994, 45, 25-45.	14.3	101
74	A microfluidic device and computational platform for high-throughput live imaging of gene expression. Nature Methods, 2012, 9, 1101-1106.	19.0	100
75	Systems Approaches to Identifying Gene Regulatory Networks in Plants. Annual Review of Cell and Developmental Biology, 2008, 24, 81-103.	9.4	96
76	Molecular Analysis of the SCARECROW Gene in Maize Reveals a Common Basis for Radial Patterning in Diverse Meristems. Plant Cell, 2000, 12, 1307-1318.	6.6	95
77	X-Ray Computed Tomography Reveals the Response of Root System Architecture to Soil Texture. Plant Physiology, 2016, 171, 2028-2040.	4.8	87
78	Spatiotemporal signalling in plant development. Nature Reviews Genetics, 2013, 14, 631-644.	16.3	84
79	Not just another hole in the wall: understanding intercellular protein trafficking. Genes and Development, 2005, 19, 189-195.	5.9	82
80	Tracking transcription factor mobility and interaction in Arabidopsis roots with fluorescence correlation spectroscopy. ELife, 2016, 5, .	6.0	79
81	Conservation and Diversification of SCARECROW in Maize. Plant Molecular Biology, 2005, 59, 619-630.	3.9	73
82	Cell wall remodeling and vesicle trafficking mediate the root clock in <i>Arabidopsis</i> . Science, 2020, 370, 819-823.	12.6	73
83	Advanced imaging techniques for the study of plant growth and development. Trends in Plant Science, 2014, 19, 304-310.	8.8	72
84	Regulation of Division and Differentiation of Plant Stem Cells. Annual Review of Cell and Developmental Biology, 2018, 34, 289-310.	9.4	72
85	The Lateral Root Cap Acts as an Auxin Sink that Controls Meristem Size. Current Biology, 2019, 29, 1199-1205.e4.	3.9	72
86	Plant immune system activation is necessary for efficient root colonization by auxin-secreting beneficial bacteria. Cell Host and Microbe, 2021, 29, 1507-1520.e4.	11.0	70
87	Getting to the root of plant biology: impact of the Arabidopsis genome sequence on root research. Plant Journal, 2010, 61, 992-1000.	5.7	67
88	G-quadruplex structures trigger RNA phase separation. Nucleic Acids Research, 2019, 47, 11746-11754.	14.5	67
89	Anchorene is a carotenoid-derived regulatory metabolite required for anchor root formation in <i>Arabidopsis</i> . Science Advances, 2019, 5, eaaw6787.	10.3	67
90	Control of Arabidopsis lateral root primordium boundaries by MYB36. New Phytologist, 2017, 213, 105-112.	7.3	65

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91	Novel technologies for emission reduction complement conservation agriculture to achieve negative emissions from row-crop production. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	64
92	GTL1 and DF1 regulate root hair growth through transcriptional repression of <i>ROOT HAIR DEFECTIVE 6-LIKE 4</i> in <i>Arabidopsis</i> . Development (Cambridge), 2018, 145, .	2.5	63
93	Small but Mighty: Functional Peptides Encoded by Small ORFs in Plants. Proteomics, 2018, 18, e1700038.	2.2	63
94	Getting to the root of plant development: the genetics of Arabidopsis root formation. Trends in Genetics, 1994, 10, 84-88.	6.7	61
95	Integrated detection of natural antisense transcripts using strand-specific RNA sequencing data. Genome Research, 2013, 23, 1730-1739.	5.5	58
96	Quantitative Trait Locus Mapping Reveals Regions of the Maize Genome Controlling Root System Architecture. Plant Physiology, 2015, 167, 1487-1496.	4.8	58
97	Signals that regulate stem cell activity during plant development. Current Opinion in Genetics and Development, 2005, 15, 388-394.	3.3	57
98	Cell type-specific transcriptional profiling: implications for metabolite profiling. Plant Journal, 2012, 70, 5-17.	5.7	57
99	High-throughput imaging and analysis of root system architecture in <i>Brachypodium distachyon</i> under differential nutrient availability. Philosophical Transactions of the Royal Society B: Biological Sciences, 2012, 367, 1559-1569.	4.0	54
100	Establishment of Expression in the SHORTROOT-SCARECROW Transcriptional Cascade through Opposing Activities of Both Activators and Repressors. Developmental Cell, 2016, 39, 585-596.	7.0	54
101	Two New Loci, PLEIADE and HYADE, Implicate Organ-Specific Regulation of Cytokinesis in Arabidopsis. Plant Physiology, 2002, 130, 312-324.	4.8	50
102	A SIMPLE Pipeline for Mapping Point Mutations. Plant Physiology, 2017, 174, 1307-1313.	4.8	50
103	Root layers: complex regulation of developmental patterning. Current Opinion in Genetics and Development, 2008, 18, 354-361.	3.3	49
104	Stem cells. Current Biology, 1999, 9, R171-R172.	3.9	48
105	Functional genomics of root growth and development in Arabidopsis. Current Opinion in Plant Biology, 2009, 12, 165-171.	7.1	48
106	Uncovering Gene Regulatory Networks Controlling Plant Cell Differentiation. Trends in Genetics, 2017, 33, 529-539.	6.7	47
107	Framework for gradual progression of cell ontogeny in the <i>Arabidopsis</i> root meristem. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8922-E8929.	7.1	46
108	An auxin-regulable oscillatory circuit drives the root clock in <i>Arabidopsis</i> . Science Advances, 2021, 7, .	10.3	46

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109	Reconstructing spatiotemporal gene expression data from partial observations. <i>Bioinformatics</i> , 2009, 25, 2581-2587.	4.1	45
110	Mechanism and function of root circumnutation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	45
111	Symmetry Breaking in Plants: Molecular Mechanisms Regulating Asymmetric Cell Divisions in <i>Arabidopsis</i> . <i>Cold Spring Harbor Perspectives in Biology</i> , 2009, 1, a000497-a000497.	5.5	40
112	Shootward and rootward: peak terminology for plant polarity. <i>Trends in Plant Science</i> , 2010, 15, 593-594.	8.8	39
113	Information processing without brains – the power of intercellular regulators in plants. <i>Development (Cambridge)</i> , 2010, 137, 1215-1226.	2.5	38
114	Integrative systems biology: an attempt to describe a simple weed. <i>Current Opinion in Plant Biology</i> , 2012, 15, 162-167.	7.1	38
115	Fluorescence-Activated Cell Sorting in Plant Developmental Biology. <i>Methods in Molecular Biology</i> , 2010, 655, 313-319.	0.9	38
116	Analysis of SCARECROW expression using a rapid system for assessing transgene expression in <i>Arabidopsis</i> roots. <i>Plant Journal</i> , 1997, 12, 957-963.	5.7	37
117	Apical–basal polarity: why plant cells don't stand on their heads. <i>Trends in Plant Science</i> , 2006, 11, 12-14.	8.8	37
118	A Lin28 homologue reprograms differentiated cells to stem cells in the moss <i>Physcomitrella patens</i> . <i>Nature Communications</i> , 2017, 8, 14242.	12.8	37
119	Minimum requirements for changing and maintaining endodermis cell identity in the <i>Arabidopsis</i> root. <i>Nature Plants</i> , 2018, 4, 586-595.	9.3	37
120	Network building: transcriptional circuits in the root. <i>Current Opinion in Plant Biology</i> , 2004, 7, 582-588.	7.1	36
121	Mechanism of Dual Targeting of the Phytochrome Signaling Component HEMERA/pTAC12 to Plastids and the Nucleus. <i>Plant Physiology</i> , 2017, 173, 1953-1966.	4.8	36
122	Transcription factors and hormones: new insights into plant cell differentiation. <i>Current Opinion in Cell Biology</i> , 2006, 18, 710-714.	5.4	35
123	Unraveling the Dynamic Transcriptome. <i>Plant Cell</i> , 2006, 18, 2101-2111.	6.6	35
124	<i>Arabidopsis thaliana</i> as a model organism in systems biology. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2009, 1, 372-379.	6.6	35
125	Transcriptional networks in root cell fate specification. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2009, 1789, 315-325.	1.9	34
126	Evolution, Interactions, and Biological Networks. <i>PLoS Biology</i> , 2007, 5, e11.	5.6	33

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127	Trait-to-Gene. <i>Current Biology</i> , 2003, 13, 129-133.	3.9	32
128	A plant lipocalin promotes retinal-mediated oscillatory lateral root initiation. <i>Science</i> , 2021, 373, 1532-1536.	12.6	32
129	Distinct sensitivities to phosphate deprivation suggest that <scp>RGF</scp> peptides play disparate roles in <i>Arabidopsis thaliana</i> root development. <i>New Phytologist</i> , 2015, 207, 683-691.	7.3	31
130	Patterning the primary root in <i>Arabidopsis</i>. <i>Wiley Interdisciplinary Reviews: Developmental Biology</i> , 2012, 1, 675-691.	5.9	30
131	The <i>Arabidopsis</i> GRAS-type SCL28 transcription factor controls the mitotic cell cycle and division plane orientation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	30
132	Using Cauliflower to Find Conserved Non-Coding Regions in Arabidopsis. <i>Plant Physiology</i> , 2002, 129, 451-454.	4.8	29
133	Single-cell analysis of cell identity in the Arabidopsis root apical meristem: insights and opportunities. <i>Journal of Experimental Botany</i> , 2021, 72, 6679-6686.	4.8	28
134	Is the shoot a root with a view?. <i>Current Opinion in Plant Biology</i> , 1999, 2, 39-43.	7.1	26
135	Beyond Arabidopsis. <i>Translational Biology Meets Evolutionary Developmental Biology. Plant Physiology</i> , 2004, 135, 611-614.	4.8	26
136	Reconstructing regulatory network transitions. <i>Trends in Cell Biology</i> , 2011, 21, 442-451.	7.9	26
137	Plant science decadal vision 2020â€“2030: Reimagining the potential of plants for a healthy and sustainable future. <i>Plant Direct</i> , 2020, 4, e00252.	1.9	26
138	Time-based patterning in development. <i>Transcription</i> , 2011, 2, 124-129.	3.1	25
139	Detailed reconstruction of 3D plant root shape. , 2011, , .		24
140	High-Throughput RNA Isolation Technologies. <i>New Tools for High-Resolution Gene Expression Profiling in Plant Systems. Plant Physiology</i> , 2005, 138, 585-590.	4.8	23
141	Intergenic and Genic Sequence Lengths Have Opposite Relationships with Respect to Gene Expression. <i>PLoS ONE</i> , 2008, 3, e3670.	2.5	23
142	Sequence Requirements of the 5-Enolpyruvylshikimate-3-Phosphate Synthase 5'-Upstream Region for Tissue-Specific Expression in Flowers and Seedlings. <i>Plant Cell</i> , 1990, 2, 849.	6.6	22
143	Quantification of transcription factor expression from Arabidopsis images. <i>Bioinformatics</i> , 2006, 22, e323-e331.	4.1	21
144	Integrated functional networks of process, tissue, and developmental stage specific interactions in Arabidopsis thaliana. <i>BMC Systems Biology</i> , 2010, 4, 180.	3.0	21

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145	Transcriptional Switches Direct Plant Organ Formation and Patterning. <i>Current Topics in Developmental Biology</i> , 2012, 98, 229-257.	2.2	19
146	Systems biology. <i>Current Biology</i> , 2004, 14, R179-R180.	3.9	18
147	Cortex proliferation. <i>Plant Signaling and Behavior</i> , 2009, 4, 551-553.	2.4	18
148	Spatiotemporal analysis identifies ABF2 and ABF3 as key hubs of endodermal response to nitrate. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	17
149	Transcriptional Networks Controlling Plant Development: Fig. 1.. <i>Plant Physiology</i> , 2001, 125, 109-111.	4.8	16
150	Physiological mechanisms contributing to the QTL qDTY3.2 effects on improved performance of rice Moroberekan x Swarna BC2F3:4 lines under drought. <i>Rice</i> , 2018, 11, 43.	4.0	15
151	Axis formation and polarity in plants. <i>Current Opinion in Genetics and Development</i> , 2001, 11, 405-409.	3.3	13
152	Histone Deacetylase HDA19 Affects Root Cortical Cell Fate by Interacting with SCARECROW. <i>Plant Physiology</i> , 2019, 180, 276-288.	4.8	13
153	Auxin Action: Slogging out of the Swamp. <i>Current Biology</i> , 2002, 12, R389-R390.	3.9	12
154	Integrating gene flow, crop biology, and farm management in on-farm conservation of avocado (<i>Persea americana</i> , Lauraceae). <i>American Journal of Botany</i> , 2003, 90, 1619-1627.	1.7	11
155	Stem Cell Research Goes Underground: The RETINOBLASTOMA-RELATED Gene in Root Development. <i>Cell</i> , 2005, 123, 1180-1182.	28.9	10
156	Defining the Path from Stem Cells to Differentiated Tissue. <i>Current Topics in Developmental Biology</i> , 2016, 116, 35-43.	2.2	10
157	Genes that regulate plant development. <i>Plant Science</i> , 1992, 83, 115-126.	3.6	9
158	Toward a Systems Analysis of the Root. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2012, 77, 91-96.	1.1	9
159	Capturing in-field root system dynamics with RootTracker. <i>Plant Physiology</i> , 2021, 187, 1117-1130.	4.8	9
160	POWRS: Position-Sensitive Motif Discovery. <i>PLoS ONE</i> , 2012, 7, e40373.	2.5	9
161	Arabidopsis Functional Genomics. <i>Plant Physiology</i> , 2002, 129, 393-393.	4.8	8
162	Optimizing root system architecture in biofuel crops for sustainable energy production and soil carbon sequestration. <i>F1000 Biology Reports</i> , 2010, 2, 65.	4.0	8

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163	Large Cellular Inclusions Accumulate in Arabidopsis Roots Exposed to Low-Sulfur Conditions. <i>Plant Physiology</i> , 2015, 168, 1573-1589.	4.8	7
164	Growth control of root architecture. , 2012, , 373-386.		6
165	A Friend in Need (of Nutrients) Is aâ€¦. <i>Cell</i> , 2016, 165, 269-271.	28.9	5
166	Tissue-Specific Transcriptome Profiling in Arabidopsis Roots. <i>Methods in Molecular Biology</i> , 2017, 1610, 107-122.	0.9	5
167	Phage-Resistant Bacteria Reveal a Role for Potassium in Root Colonization. <i>MBio</i> , 2021, 12, e0140321.	4.1	5
168	A Common Switch used by Plants and Animals. <i>Cell</i> , 2004, 116, 4-5.	28.9	4
169	A Co-opted Regulator of Lateral Root Development Controls Nodule Organogenesis in Lotus. <i>Developmental Cell</i> , 2020, 52, 6-7.	7.0	4
170	Root development: Signaling down and around. <i>BioEssays</i> , 1997, 19, 959-965.	2.5	3
171	Development and Ecology in the Time of Systems Biology. <i>Developmental Cell</i> , 2004, 7, 329-330.	7.0	3
172	A systems approach to understanding root development. <i>Canadian Journal of Botany</i> , 2006, 84, 695-701.	1.1	3
173	Studying Root Development Using a Genomic Approach. , 0, , 325-351.		3
174	HEC of a Job Regulating Stem Cells. <i>Developmental Cell</i> , 2014, 28, 349-350.	7.0	3
175	Single-cell genomics revolutionizes plant development studies across scales. <i>Development (Cambridge)</i> , 2022, 149, .	2.5	3
176	managedpop: a computer simulation to project allelic diversity in managed populations with overlapping generations. <i>Molecular Ecology Notes</i> , 2002, 2, 615-617.	1.7	2
177	Developmental Networks. <i>Plant Physiology</i> , 2005, 138, 548-549.	4.8	2
178	Taking a Developmental Perspective on Systems Biology. <i>Developmental Cell</i> , 2011, 21, 27-28.	7.0	2
179	Arabidopsis as a Model for Systems Biology. , 2013, , 391-406.		2
180	VAP-RELATED SUPPRESSORS OF TOO MANY MOUTHS (VST) family proteins are regulators of root system architecture. <i>Plant Physiology</i> , 2021, 185, 457-468.	4.8	2

#	ARTICLE	IF	CITATIONS
181	Polarized Radicals. <i>Cell</i> , 2013, 153, 285-286.	28.9	1
182	Identifying Gene Regulatory Networks in Arabidopsis by In Silico Prediction, Yeast-1-Hybrid, and Inducible Gene Profiling Assays. <i>Methods in Molecular Biology</i> , 2016, 1370, 29-50.	0.9	1
183	Lateral Root Initiation: The Emergence of New Primordia Following Cell Death. <i>Current Biology</i> , 2020, 30, R121-R122.	3.9	1
184	Appointments and awards. <i>Plant Molecular Biology Reporter</i> , 1992, 10, 4-4.	1.8	0
185	International Symposium on the Molecular Genetics of Root Development. <i>Plant Molecular Biology Reporter</i> , 1993, 11, 60-64.	1.8	0
186	Signaling the tips: Regulation of stem cell function in plants. <i>Differentiation</i> , 2001, 68, 155-158.	1.9	0
187	Additions and corrections: A systems approach to understanding root development. <i>Canadian Journal of Botany</i> , 2006, 84, 1508.	1.1	0
188	Promoting Collaborative Interdisciplinary Research at the Duke Center for Systems Biology. <i>ACS Synthetic Biology</i> , 2012, 1, 153-155.	3.8	0
189	Arabidopsis Root. , 2013, , 2-1-2-17.		0
190	Reply to Amundson: Time to go to work. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2122842119.	7.1	0
191	Development and Systems Biology: Riding the Genomics Wave towards a Systems Understanding of Root Development. , 0, , 304-330.		0