Ralf Reski

List of Publications by Year
in descending order

Version: 2024-02-01

210 papers	13,329 citations	16451 64 h-index	30922 102 g-index
231 all docs	231 docs citations	231 times ranked	10960 citing authors

#	Article	IF	Citations
1	Viral suppressor of <scp>RNA</scp> silencing in vascular plants also interferes with the development of the bryophyte <scp>Physcomitrella patens</scp>. Plant, Cell and Environment, 2022, 45, 220-235.	5.7	3
2	Autopolyploidization affects transcript patterns and gene targeting frequencies in Physcomitrella. Plant Cell Reports, 2022, 41, 153-173.	5.6	5
3	O-methylated N-glycans Distinguish Mosses from Vascular Plants. Biomolecules, 2022, 12, 136.	4.0	8
4	A synthetic protein as efficient multitarget regulator against complement over-activation. Communications Biology, 2022, 5, 152.	4.4	9
5	Unexpected Arabinosylation after Humanization of Plant Protein N-Glycosylation. Frontiers in Bioengineering and Biotechnology, 2022, 10, 838365.	4.1	6
6	Process Engineering of Biopharmaceutical Production in Moss Bioreactors via Model-Based Description and Evaluation of Phytohormone Impact. Frontiers in Bioengineering and Biotechnology, 2022, 10, 837965.	4.1	5
7	Recombinant Spider Silk: Promises and Bottlenecks. Frontiers in Bioengineering and Biotechnology, 2022, 10, 835637.	4.1	35
8	Spindle motility skews division site determination during asymmetric cell division in Physcomitrella. Nature Communications, 2022, 13, 2488.	12.8	7
9	The mosaic oat genome gives insights into a uniquely healthy cereal crop. Nature, 2022, 606, 113-119.	27.8	70
10	Axenic in vitro cultivation of 19 peat moss (Sphagnum L.) species as a resource for basic biology, biotechnology, and paludiculture. New Phytologist, 2021, 229, 861-876.	7.3	28
11	Automated and semi-automated enhancement, segmentation and tracing of cytoskeletal networks in microscopic images: A review. Computational and Structural Biotechnology Journal, 2021, 19, 2106-2120.	4.1	16
12	Biopolymer segmentation from CLSM microscopy images using a convolutional neural network. Proceedings in Applied Mathematics and Mechanics, 2021, 20, e202000188.	0.2	2
13	Function of the HYDROXYCINNAMOYL-CoA:SHIKIMATE HYDROXYCINNAMOYL TRANSFERASE is evolutionarily conserved in embryophytes. Plant Cell, 2021, 33, 1472-1491.	6.6	45
14	Natural Products from Bryophytes: From Basic Biology to Biotechnological Applications. Critical Reviews in Plant Sciences, 2021, 40, 191-217.	5.7	33
15	Polyploidization within the Funariaceaeâ€"a key principle behind speciation, sporophyte reduction and the high variance of spore diameters?. Bryophyte Diversity and Evolution, 2021, 43, .	1.1	9
16	Expression of a human cDNA in moss results in spliced mRNAs and fragmentary protein isoforms. Communications Biology, 2021, 4, 964.	4.4	12
17	Medium optimization for biomass production of three peat moss (Sphagnum L.) species using fractional factorial design and response surface methodology. Bioresource Technology Reports, 2021, 15, 100729.	2.7	3
18	Convergence of sphingolipid desaturation across over 500 million years of plant evolution. Nature Plants, 2021, 7, 219-232.	9.3	31

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19	Testing a novel biotechnological passive sampler for monitoring atmospheric PAH pollution. Journal of Hazardous Materials, 2020, 381, 120949.	12.4	17
20	Mosses in biotechnology. Current Opinion in Biotechnology, 2020, 61, 21-27.	6.6	39
21	Stable Protein Sialylation in Physcomitrella. Frontiers in Plant Science, 2020, 11, 610032.	3.6	21
22	A NanoFE simulation-based surrogate machine learning model to predict mechanical functionality of protein networks from live confocal imaging. Computational and Structural Biotechnology Journal, 2020, 18, 2774-2788.	4.1	6
23	Chloroplasts require glutathione reductase to balance reactive oxygen species and maintain efficient photosynthesis. Plant Journal, 2020, 103, 1140-1154.	5.7	47
24	Editorial overview: Plant biotechnology. Current Opinion in Biotechnology, 2020, 61, iii-v.	6.6	0
25	Development of a method for protonema proliferation of peat moss (Sphagnum squarrosum) through regeneration analysis. New Phytologist, 2019, 221, 1160-1171.	7.3	10
26	Biosurface properties and lead adsorption in a clone of Sphagnum palustre (Mosses): Towards a unified protocol of biomonitoring of airborne heavy metal pollution. Chemosphere, 2019, 236, 124375.	8.2	15
27	Recombinant Production of MFHR1, A Novel Synthetic Multitarget Complement Inhibitor, in Moss Bioreactors. Frontiers in Plant Science, 2019, 10, 260.	3.6	24
28	Critical Evaluation of Strategies for the Production of Blood Coagulation Factors in Plant-Based Systems. Frontiers in Plant Science, 2019, 10, 261.	3.6	10
29	Single-cell transcriptome analysis of Physcomitrella leaf cells during reprogramming using microcapillary manipulation. Nucleic Acids Research, 2019, 47, 4539-4553.	14.5	39
30	ABA-Induced Vegetative Diaspore Formation in Physcomitrella patens. Frontiers in Plant Science, 2019, 10, 315.	3.6	30
31	The plastid skeleton: a source of ideas in the nano range. , 2019, , 163-166.		3
32	Das Plastidenskelett: ein Ideengeber im Nanobereich. , 2019, , 163-166.		0
33	Evolutive approaches to explorative design methods in architecture. , 2019, , 134-141.		0
34	Evolutive AnsÃtze fÃ¼r explorative Entwurfsmethoden in der Architektur. , 2019, , 134-141.		0
35	The Physcomitrella patens gene atlas project: largeâ€scale <scp>RNA</scp>â€seq based expression data. Plant Journal, 2018, 95, 168-182.	5.7	115
36	RecQ Helicases Function in Development, DNA Repair, and Gene Targeting in Physcomitrella patens. Plant Cell, 2018, 30, 717-736.	6.6	44

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37	The single berberine bridge enzyme homolog of PhyscomitrellaÂpatens is a cellobiose oxidase. FEBS Journal, 2018, 285, 1923-1943.	4.7	17
38	The loss of SMG1 causes defects in quality control pathways in Physcomitrella patens. Nucleic Acids Research, 2018, 46, 5822-5836.	14.5	24
39	Enabling the water-to-land transition. Nature Plants, 2018, 4, 67-68.	9.3	31
40	The MFHR1 Fusion Protein Is a Novel Synthetic Multitarget Complement Inhibitor with Therapeutic Potential. Journal of the American Society of Nephrology: JASN, 2018, 29, 1141-1153.	6.1	28
41	The Physcomitrella patens chromosomeâ€scale assembly reveals moss genome structure and evolution. Plant Journal, 2018, 93, 515-533.	5.7	406
42	Featureâ€based Classification of Protein Networks using Confocal Microscopy Imaging and Machine Learning. Proceedings in Applied Mathematics and Mechanics, 2018, 18, e201800246.	0.2	1
43	Moss-made complement therapeutics. Molecular Immunology, 2018, 102, 209.	2.2	0
44	Host Cell Proteome ofPhyscomitrella patensHarbors Proteases and Protease Inhibitors under Bioproduction Conditions. Journal of Proteome Research, 2018, 17, 3749-3760.	3.7	19
45	Physcomitrella patens, a versatile synthetic biology chassis. Plant Cell Reports, 2018, 37, 1409-1417.	5.6	50
46	Computational 3D imaging to quantify structural components and assembly of protein networks. Acta Biomaterialia, 2018, 69, 206-217.	8.3	14
47	Quantitative moss cell biology. Current Opinion in Plant Biology, 2018, 46, 39-47.	7.1	17
48	Cytological analysis and structural quantification of FtsZ1-2 and FtsZ2-1 network characteristics in Physcomitrella patens. Scientific Reports, 2018, 8, 11165.	3.3	14
49	Sphagnum palustre clone vs native Pseudoscleropodium purum : A first trial in the field to validate the future of the moss bag technique. Environmental Pollution, 2017, 225, 323-328.	7.5	29
50	Strigolactone biosynthesis is evolutionarily conserved, regulated by phosphate starvation and contributes to resistance against phytopathogenic fungi in a moss, Physcomitrella patens. New Phytologist, 2017, 216, 455-468.	7.3	121
51	A phenol-enriched cuticle is ancestral to lignin evolution in land plants. Nature Communications, 2017, 8, 14713.	12.8	157
52	Moss-Produced, Glycosylation-Optimized Human Factor H for Therapeutic Application in Complement Disorders. Journal of the American Society of Nephrology: JASN, 2017, 28, 1462-1474.	6.1	43
53	Dithiol disulphide exchange in redox regulation of chloroplast enzymes in response to evolutionary and structural constraints. Plant Science, 2017, 255, 1-11.	3.6	38
54	Treatment of experimental C3 Glomerulopathy by human complement factor H produced in glycosylation-optimized Physcomitrella patens. Molecular Immunology, 2017, 89, 120.	2.2	8

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55	Selfing in Haploid Plants and Efficacy of Selection: Codon Usage Bias in the Model Moss Physcomitrella patens. Genome Biology and Evolution, 2017, 9, 1528-1546.	2.5	21
56	Microscopy of Physcomitrella patens sperm cells. Plant Methods, 2017, 13, 33.	4.3	13
57	Approaches to Characterize Organelle, Compartment, or Structure Purity. Methods in Molecular Biology, 2017, 1511, 13-28.	0.9	4
58	The mitochondrial proteome of the moss Physcomitrella patens. Mitochondrion, 2017, 33, 38-44.	3.4	5
59	Alternation of generations â€“ unravelling the underlying molecular mechanism of a 165â€yearâ€old botanical observation. Plant Biology, 2016, 18, 549-551.	3.8	19
60	Spatioâ€temporal patterning of arginylâ€<scp>tRNA</scp> protein transferase (<scp>ATE</scp>) contributes to gametophytic development in a moss. New Phytologist, 2016, 209, 1014-1027.	7.3	35
61	Chloroplast FBPase and SBPase are thioredoxin-linked enzymes with similar architecture but different evolutionary histories. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6779-6784.	7.1	60
62	Identification of Targets and Interaction Partners of Arginyl-tRNA Protein Transferase in the Moss Physcomitrella patens. Molecular and Cellular Proteomics, 2016, 15, 1808-1822.	3.8	25
63	Best options for the exposure of traditional and innovative moss bags: A systematic evaluation in three European countries. Environmental Pollution, 2016, 214, 362-373.	7.5	61
64	The Polycomb group protein CLF emerges as a specific tri-methylase of H3K27 regulating gene expression and development in Physcomitrella patens. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2016, 1859, 860-870.	1.9	17
65	Molecular and chemical characterization of a Sphagnum palustre clone: Key steps towards a standardized and sustainable moss bag technique. Ecological Indicators, 2016, 71, 388-397.	6.3	29
66	Origin and function of stomata in the moss Physcomitrella patens. Nature Plants, 2016, 2, 16179.	9.3	138
67	A single homeobox gene triggers phase transition, embryogenesis and asexual reproduction. Nature Plants, 2016, 2, 15209.	9.3	116
68	Analysis of confocal microscopy image data of Physcomitrella chloroplasts to reveal adaptation principles leading to structural stability at the nanoscale. Proceedings in Applied Mathematics and Mechanics, 2016, 16, 69-70.	0.2	4
69	Genetic analysis of Physcomitrella patens identifies ABSCISIC ACID NON-RESPONSIVE (ANR), a regulator of ABA responses unique to basal land plants and required for desiccation tolerance. Plant Cell, 2016, 28, tpc.00091.2016.	6.6	98
70	Chemical and structural characterization of copper adsorbed on mosses (Bryophyta). Journal of Hazardous Materials, 2016, 308, 343-354.	12.4	20
71	Implications of plant glycans in the development of innovative vaccines. Expert Review of Vaccines, 2016, 15, 915-925.	4.4	26
72	The Genome of the Model Moss Physcomitrella patens. Advances in Botanical Research, 2016, 78, 97-140.	1.1	9

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73	Metal and proton adsorption capacities of natural and cloned Sphagnum mosses. Journal of Colloid and Interface Science, 2016, 461, 326-334.	9.4	34
74	Can mosses serve as model organisms for forest research?. Annals of Forest Science, 2016, 73, 135-146.	2.0	13
75	Analysis of Physcomitrella Chloroplasts to Reveal Adaptation Principles Leading to Structural Stability at the Nano-Scale. Biologically-inspired Systems, 2016, , 261-275.	0.2	6
76	Evolutionary Processes as Models for Exploratory Design. Biologically-inspired Systems, 2016, , 295-318.	0.2	3
77	Mossâ€made pharmaceuticals: from bench to bedside. Plant Biotechnology Journal, 2015, 13, 1191-1198.	8.3	95
78	Mitochondrial Dynamics and the ER: The Plant Perspective. Frontiers in Cell and Developmental Biology, 2015, 3, 78.	3.7	49
79	Clonal in vitro propagation of peat mosses (Sphagnum L.) as novel green resources for basic and applied research. Plant Cell, Tissue and Organ Culture, 2015, 120, 1037-1049.	2.3	42
80	An Env-derived multi-epitope HIV chimeric protein produced in the moss Physcomitrella patens is immunogenic in mice. Plant Cell Reports, 2015, 34, 425-433.	5.6	31
81	Matrix solid phase dispersion method for determination of polycyclic aromatic hydrocarbons in moss. Journal of Chromatography A, 2015, 1406, 19-26.	3.7	20
82	Stomatal Guard Cells Co-opted an Ancient ABA-Dependent Desiccation Survival System to Regulate Stomatal Closure. Current Biology, 2015, 25, 928-935.	3.9	154
83	DNA METHYLTRANSFERASE 1 is involved in mCG and mCCG DNA methylation and is essential for sporophyte development in Physcomitrella patens. Plant Molecular Biology, 2015, 88, 387-400.	3.9	45
84	Insights from the cold transcriptome of <scp>P</scp>hyscomitrella patens: global specialization pattern of conserved transcriptional regulators and identification of orphan genes involved in cold acclimation. New Phytologist, 2015, 205, 869-881.	7.3	84
85	Gene Targeting for Precision Glyco-Engineering: Production of Biopharmaceuticals Devoid of Plant-Typical Glycosylation in Moss Bioreactors. Methods in Molecular Biology, 2015, 1321, 213-224.	0.9	19
86	Applied Bryology - Bryotechnology. Bryophyte Diversity and Evolution, 2015, 31, 22.	1.1	12
87	Plasma Membrane-Targeted PIN Proteins Drive Shoot Development in a Moss. Current Biology, 2014, 24, 2776-2785.	3.9	133
88	Evolution and communication of subcellular compartments. Plant Signaling and Behavior, 2014, 9, e28993.	2.4	10
89	The potential ofPhyscomitrella patensas a platform for the production of plant-based vaccines. Expert Review of Vaccines, 2014, 13, 203-212.	4.4	15
90	ppdb: plant promoter database version 3.0. Nucleic Acids Research, 2014, 42, D1188-D1192.	14.5	61

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91	Glyco-engineering for biopharmaceutical production in moss bioreactors. Frontiers in Plant Science, 2014, 5, 346.	3.6	39
92	Quantitative Analysis of the Mitochondrial and Plastid Proteomes of the Moss Physcomitrella patens Reveals Protein Macrocompartmentation and Microcompartmentation. Plant Physiology, 2014, 164, 2081-2095.	4.8	61
93	Molecular evidence for convergent evolution and allopolyploid speciation within the Physcomitrium-Physcomitrellaspecies complex. BMC Evolutionary Biology, 2014, 14, 158.	3.2	48
94	Design and Characterization of a Modular Membrane Protein Anchor to Functionalize the Moss Physcomitrella patens with Extracellular Catalytic and/or Binding Activities. ACS Synthetic Biology, 2014, 3, 990-994.	3.8	5
95	Balanced activity of microRNA166/165 and its target transcripts from the class III homeodomain-leucine zipper family regulates root growth in Arabidopsis thaliana. Plant Cell Reports, 2014, 33, 945-953.	5.6	83
96	High contents of very long-chain polyunsaturated fatty acids in different moss species. Plant Cell Reports, 2014, 33, 245-254.	5.6	48
97	Isopentenyltransferase-1 (IPT1) knockout in Physcomitrella together with phylogenetic analyses of IPTs provide insights into evolution of plant cytokinin biosynthesis. Journal of Experimental Botany, 2014, 65, 2533-2543.	4.8	57
98	A red light-controlled synthetic gene expression switch for plant systems. Molecular BioSystems, 2014, 10, 1679-1688.	2.9	89
99	Largeâ€scale gene expression profiling data for the model moss <scp>P</scp>hyscomitrella patens aid understanding of developmental progression, culture and stress conditions. Plant Journal, 2014, 79, 530-539.	5.7	82
100	Reannotation and extended community resources for the genome of the non-seed plant Physcomitrella patens provide insights into the evolution of plant gene structures and functions. BMC Genomics, 2013, 14, 498.	2.8	170
101	The Plant Ontology as a Tool for Comparative Plant Anatomy and Genomic Analyses. Plant and Cell Physiology, 2013, 54, e1-e1.	3.1	131
102	Physcomitrella PpORS, Basal to Plant Type III Polyketide Synthases in Phylogenetic Trees, Is a Very Long Chain 2â€²-Oxoalkylresorcinol Synthase. Journal of Biological Chemistry, 2013, 288, 2767-2777.	3.4	19
103	A gene responsible for prolyl-hydroxylation of moss-produced recombinant human erythropoietin. Scientific Reports, 2013, 3, 3019.	3.3	50
104	System for Stable Î²-Estradiol-Inducible Gene Expression in the Moss Physcomitrella patens. PLoS ONE, 2013, 8, e77356.	2.5	71
105	Involvement of a Class III Peroxidase and the Mitochondrial Protein TSPO in Oxidative Burst Upon Treatment of Moss Plants with a Fungal Elicitor. Molecular Plant-Microbe Interactions, 2012, 25, 363-371.	2.6	66
106	DICER-LIKE3 Activity in Physcomitrella patens DICER-LIKE4 Mutants Causes Severe Developmental Dysfunction and Sterility. Molecular Plant, 2012, 5, 1281-1294.	8.3	45
107	Biosynthesis of allene oxides in Physcomitrella patens. BMC Plant Biology, 2012, 12, 228.	3.6	39
108	The relevance of compartmentation for cysteine synthesis in phototrophic organisms. Protoplasma, 2012, 249, 147-155.	2.1	22

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109	Mossâ€based production of asialoâ€erythropoietin devoid of Lewis A and other plantâ€typical carbohydrate determinants. Plant Biotechnology Journal, 2012, 10, 851-861.	8.3	74
110	Glycoprotein production in moss bioreactors. Plant Cell Reports, 2012, 31, 453-460.	5.6	57
111	Metabolite profiling of the moss Physcomitrella patens reveals evolutionary conservation of osmoprotective substances. Plant Cell Reports, 2012, 31, 427-436.	5.6	78
112	Production of biologically active recombinant human factor H in Physcomitrella. Plant Biotechnology Journal, 2011, 9, 373-383.	8.3	86
113	PpASCL, a moss ortholog of antherâ€specific chalcone synthaseâ€like enzymes, is a hydroxyalkylpyrone synthase involved in an evolutionarily conserved sporopollenin biosynthesis pathway. New Phytologist, 2011, 192, 855-868.	7.3	48
114	MicroRNA534a control of BLADEâ€ONâ€PETIOLE 1 and 2 mediates juvenileâ€toâ€adult gametophyte transition in Physcomitrella patens. Plant Journal, 2011, 65, 661-674.	5.7	29
115	Simultaneous isolation of pure and intact chloroplasts and mitochondria from moss as the basis for sub-cellular proteomics. Plant Cell Reports, 2011, 30, 205-215.	5.6	53
116	Plant biotechnology in support of the Millennium Goals. Plant Cell Reports, 2011, 30, 245-247.	5.6	4
117	Plant biotechnology in support of the Millennium Goals II. Plant Cell Reports, 2011, 30, 677-679.	5.6	4
118	A Novel Calcium Binding Site in the Slow Vacuolar Cation Channel TPC1 Senses Luminal Calcium Levels. Plant Cell, 2011, 23, 2696-2707.	6.6	93
119	Genome-Wide Phylogenetic Comparative Analysis of Plant Transcriptional Regulation: A Timeline of Loss, Gain, Expansion, and Correlation with Complexity. Genome Biology and Evolution, 2010, 2, 488-503.	2.5	174
120	Microarray analysis of the moss Physcomitrella patens reveals evolutionarily conserved transcriptional regulation of salt stress and abscisic acid signalling. Plant Molecular Biology, 2010, 72, 27-45.	3.9	110
121	Identification and characterization of NAGNAG alternative splicing in the moss Physcomitrella patens. BMC Plant Biology, 2010, 10, 76.	3.6	13
122	Targeted knockâ€out of a gene encoding sulfite reductase in the moss Physcomitrella patens affects gametophytic and sporophytic development. FEBS Letters, 2010, 584, 2271-2278.	2.8	18
123	The moss Physcomitrella patens contains cyclopentenones but no jasmonates: mutations in allene oxide cyclase lead to reduced fertility and altered sporophyte morphology. New Phytologist, 2010, 188, 740-749.	7.3	125
124	THE SPECIATION HISTORY OF THEPHYSCOMITRIUM-PHYSCOMITRELLASPECIES COMPLEX. Evolution; International Journal of Organic Evolution, 2010, 64, 217-231.	2.3	59
125	Overexpression of the Arabidopsis Gene UPRIGHT ROSETTE Reveals a Homeostatic Control for Indole-3-Acetic Acid Â Â. Plant Physiology, 2010, 153, 1311-1320.	4.8	22
126	Transcriptional Control of Gene Expression by MicroRNAs. Cell, 2010, 140, 111-122.	28.9	431

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127	Regulation of stem cell maintenance by the Polycomb protein FIE has been conserved during land plant evolution. Development (Cambridge), 2009, 136, 2433-2444.	2.5	133
128	Dead end for auxin conjugates in Physcomitrella?. Plant Signaling and Behavior, 2009, 4, 116-118.	2.4	23
129	Targeted Gene Knockouts Reveal Overlapping Functions of the Five Physcomitrella patens FtsZ Isoforms in Chloroplast Division, Chloroplast Shaping, Cell Patterning, Plant Development, and Gravity Sensing. Molecular Plant, 2009, 2, 1359-1372.	8.3	44
130	Challenges to our current view on chloroplasts. Biological Chemistry, 2009, 390, 731-738.	2.5	22
131	The evolution of nuclear auxin signalling. BMC Evolutionary Biology, 2009, 9, 126.	3.2	115
132	Auxin-binding proteins without KDEL sequence in the moss Funaria hygrometrica. Plant Cell Reports, 2009, 28, 1747-1758.	5.6	14
133	Functional crossâ€kingdom conservation of mammalian and moss (Physcomitrella patens) transcription, translation and secretion machineries. Plant Biotechnology Journal, 2009, 7, 73-86.	8.3	41
134	Moss (Physcomitrella patens) GH3 proteins act in auxin homeostasis. New Phytologist, 2009, 181, 323-338.	7.3	129
135	Prediction of dual protein targeting to plant organelles. New Phytologist, 2009, 183, 224-236.	7.3	73
136	Current achievements in the production of complex biopharmaceuticals with moss bioreactors. Bioprocess and Biosystems Engineering, 2008, 31, 3-9.	3.4	89
137	Exploring plant biodiversity: the Physcomitrella genome and beyond. Trends in Plant Science, 2008, 13, 542-549.	8.8	132
138	The Physcomitrella Genome Reveals Evolutionary Insights into the Conquest of Land by Plants. Science, 2008, 319, 64-69.	12.6	1,712
139	A sequenceâ€anchored genetic linkage map for the moss, Physcomitrella patens. Plant Journal, 2008, 56, 855-866.	5.7	42
140	A P _{IIB} -type Ca ²⁺ -ATPase is essential for stress adaptation in Physcomitrella patens. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 19555-19560.	7.1	116
141	Specific Gene Silencing by Artificial MicroRNAs in Physcomitrella patens: An Alternative to Targeted Gene Knockouts Â Â. Plant Physiology, 2008, 148, 684-693.	4.8	109
142	PlanTAPDB, a Phylogeny-Based Resource of Plant Transcription-Associated Proteins. Plant Physiology, 2007, 143, 1452-1466.	4.8	79
143	The Putative Moss 3â€²-Phosphoadenosine-5â€²-phosphosulfate Reductase Is a Novel Form of Adenosine-5â€²-phosphosulfate Reductase without an Iron-Sulfur Cluster. Journal of Biological Chemistry, 2007, 282, 22930-22938.	3.4	37
144	A mitochondrial protein homologous to the mammalian peripheralâ€type benzodiazepine receptor is essential for stress adaptation in plants. Plant Journal, 2007, 51, 1004-1018.	5.7	83

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145	Filamentous temperatureâ€sensitive Z (FtsZ) isoforms specifically interact in the chloroplasts and in the cytosol of Physcomitrella patens. New Phytologist, 2007, 176, 299-310.	7.3	30
146	Evidence for the rapid expansion of microRNA-mediated regulation in early land plant evolution. BMC Plant Biology, 2007, 7, 13.	3.6	108
147	Evolutionary conservation of plant gibberellin signalling pathway components. BMC Plant Biology, 2007, 7, 65.	3.6	93
148	An ancient genome duplication contributed to the abundance of metabolic genes in the moss Physcomitrella patens. BMC Evolutionary Biology, 2007, 7, 130.	3.2	171
149	The role of the novel adenosine 5â€²-phosphosulfate reductase in regulation of sulfate assimilation of Physcomitrella patens. Plant Molecular Biology, 2007, 65, 667-676.	3.9	15
150	Dating the early evolution of plants: detection and molecular clock analyses of orthologs. Molecular Genetics and Genomics, 2007, 278, 393-402.	2.1	103
151	Moss bioreactors producing improved biopharmaceuticals. Current Opinion in Biotechnology, 2007, 18, 393-398.	6.6	73
152	Rapid Alteration of the Phosphoproteome in the MossPhyscomitrellapatensafter Cytokinin Treatment. Journal of Proteome Research, 2006, 5, 2283-2293.	3.7	32
153	Biosynthesis of C9-aldehydes in the moss Physcomitrella patensâ˜†. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2006, 1761, 301-312.	2.4	54
154	Moss Systems Biology en Route: Phytohormones in Physcomitrella Development. Plant Biology, 2006, 8, 397-406.	3.8	135
155	Identification of genic moss SSR markers and a comparative analysis of twenty-four algal and plant gene indices reveal species-specific rather than group-specific characteristics of microsatellites. BMC Plant Biology, 2006, 6, 9.	3.6	78
156	High throughput metabolic screen ofPhyscomitrellatransformants. Bryologist, 2006, 109, 247-256.	0.6	12
157	The mechanism of gene targeting in Physcomitrella patens: homologous recombination, concatenation and multiple integration. Nucleic Acids Research, 2006, 34, 6205-6214.	14.5	126
158	Enhanced recovery of a secreted recombinant human growth factor using stabilizing additives and by co-expression of human serum albumin in the moss Physcomitrella patens. Plant Biotechnology Journal, 2005, 3, 331-340.	8.3	68
159	Gene clusters involved in anaerobic benzoate degradation of Geobacter metallireducens. Molecular Microbiology, 2005, 58, 1238-1252.	2.5	147
160	Protein encoding genes in an ancient plant: analysis of codon usage, retained genes and splice sites in a moss, Physcomitrella patens. BMC Genomics, 2005, 6, 43.	2.8	56
161	Isolation and characterisation of three moss-derived beta-tubulin promoters suitable for recombinant expression. Current Genetics, 2005, 47, 111-120.	1.7	33
162	From axenic spore germination to molecular farming. Plant Cell Reports, 2005, 23, 513-521.	5.6	36

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163	EST Sequencing from Embryogenic Cyclamen persicum Cell Cultures Identifies a High Proportion of Transcripts Homologous to Plant Genes Involved in Somatic Embryogenesis. Journal of Plant Growth Regulation, 2005, 24, 102-115.	5.1	37
164	Physcomitrella patens is highly tolerant against drought, salt and osmotic stress. Planta, 2005, 220, 384-394.	3.2	205
165	Control of growth and differentiation of bioreactor cultures of Physcomitrella by environmental parameters. Plant Cell, Tissue and Organ Culture, 2005, 81, 307-311.	2.3	8
166	Cloning and expression of the tobacco CHLM sequence encoding Mg protoporphyrin IX methyltransferase and its interaction with Mg chelatase. Plant Molecular Biology, 2005, 57, 679-691.	3.9	47
167	Use of endogenous signal sequences for transient production and efficient secretion by moss (Physcomitrella patens) cells. BMC Biotechnology, 2005, 5, 30.	3.3	39
168	Effect of Ploidy Level on Growth, Differentiation, and Morphology in Physcomitrella patens. Bryologist, 2005, 108, 27-35.	0.6	29
169	A fast and flexible PEG-mediated transient expression system in plants for high level expression of secreted recombinant proteins. Journal of Biotechnology, 2005, 119, 332-342.	3.8	31
170	Moss (Physcomitrella patens) functional genomics -- Gene discovery and tool development, with implications for crop plants and human health. Briefings in Functional Genomics & Proteomics, 2005, 4, 48-57.	3.8	22
171	Isolation and Characterization ofALDH11A5, a Novel Non-Phosphorylating GAPDH cDNA from Physcomitrella patens*. Bryologist, 2004, 107, 385-387.	0.6	9
172	Targeted knockouts of Physcomitrella lacking plant-specific immunogenic N-glycans. Plant Biotechnology Journal, 2004, 2, 517-523.	8.3	221
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