Sabeeha S Merchant

List of Publications by Year
in descending order

134 papers	14,897 citations	17440 63 h-index	19749 117 g-index
151 all docs	151 docs citations	151 times ranked	13566 citing authors

#	Article	IF	Citations
1	Chlamydomonas ATX1 is essential for Cu distribution to multiple cuproâ€enzymes and maintenance of biomass in conditions demanding cuproâ€enzymeâ€dependent metabolic pathways. Plant Direct, 2022, 6, e383.	1.9	5
2	An optimized ChIPâ€Seq framework for profiling histone modifications in Chromochloris zofingiensis. Plant Direct, 2022, 6, e392.	1.9	0
3	Thank you and best wishes to Annette Kessler, peer review manager for The Plant Cell. Plant Cell, 2022, , .	6.6	1
4	Systematic characterization of gene function in the photosynthetic alga Chlamydomonas reinhardtii. Nature Genetics, 2022, 54, 705-714.	21.4	42
5	Simple steps to enable reproducibility: culture conditions affecting Chlamydomonas growth and elemental composition. Plant Journal, 2022, 111, 995-1014.	5.7	7
6	Co-expression networks in Chlamydomonas reveal significant rhythmicity in batch cultures and empower gene function discovery. Plant Cell, 2021, 33, 1058-1082.	6.6	31
7	Widespread polycistronic gene expression in green algae. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	30
8	Single-cell RNA sequencing of batch Chlamydomonas cultures reveals heterogeneity in their diurnal cycle phase. Plant Cell, 2021, 33, 1042-1057.	6.6	29
9	Single-cell visualization and quantification of trace metals in Chlamydomonas lysosome-related organelles. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	20
10	A pathogenic role for histone H3 copper reductase activity in a yeast model of Friedreichâ€™s ataxia. Science Advances, 2021, 7, eabj9889.	10.3	6
11	From economy to luxury: Copper homeostasis in Chlamydomonas and other algae. Biochimica Et Biophysica Acta - Molecular Cell Research, 2020, 1867, 118822.	4.1	35
12	Coexpressed subunits of dual genetic origin define a conserved supercomplex mediating essential protein import into chloroplasts. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 32739-32749.	7.1	30
13	An epigenetic gene silencing pathway selectively acting on transgenic DNA in the green alga Chlamydomonas. Nature Communications, 2020, 11, 6269.	12.8	58
14	The histone H3-H4 tetramer is a copper reductase enzyme. Science, 2020, 369, 59-64.	12.6	60
15	An atypical short-chain dehydrogenaseâ€“reductase functions in the relaxation of photoprotective qH in Arabidopsis. Nature Plants, 2020, 6, 154-166.	9.3	27
16	The lichen symbiosis re-viewed through the genomes of Cladonia grayi and its algal partner Asterochloris glomerata. BMC Genomics, 2019, 20, 605.	2.8	98
17	Manganese co-localizes with calcium and phosphorus in Chlamydomonas acidocalcisomes and is mobilized in manganese-deficient conditions. Journal of Biological Chemistry, 2019, 294, 17626-17641.	3.4	53
18	Multiomics resolution of molecular events during a day in the life of Chlamydomonas. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 2374-2383.	7.1	133

#	Article	IF	Citations
19	The Plant Cell Celebrates 30 Years of Publishing the Best Work in Plant Biology. Plant Cell, 2019, 31, 1-1.	6.6	1
20	A Series of Fortunate Events: Introducing Chlamydomonas as a Reference Organism. Plant Cell, 2019, 31, 1682-1707.	6.6	169
21	Reflections on The Plant Cell Classics. Plant Cell, 2019, 31, 1185-1185.	6.6	0
22	Comparative and Functional Algal Genomics. Annual Review of Plant Biology, 2019, 70, 605-638.	18.7	76
23	Regulation of Oxygenic Photosynthesis during Trophic Transitions in the Green Alga Chromochloris zofingiensis. Plant Cell, 2019, 31, 579-601.	6.6	61
24	Thank You, Editors and Reviewers of The Plant Cell. Plant Cell, 2019, 31, 2807-2812.	6.6	0
25	The Plant Cell Is Accepting Applications for Assistant Features Editors. Plant Cell, 2019, 31, tpc.00787.2019.	6.6	1
26	A Look Back from the Helm of The Plant Cell. Plant Cell, 2019, 31, 2813-2813.	6.6	1
27	Using YFP as a Reporter of Gene Expression in the Green Alga Chlamydomonas reinhardtii. Methods in Molecular Biology, 2018, 1755, 135-148.	0.9	0
28	<scp>RAF</scp>2 is a RuBis<scp>CO</scp> assembly factor in Arabidopsis thaliana. Plant Journal, 2018, 94, 146-156.	5.7	22
29	The Plant Cell Welcomes Assistant Features Editors. Plant Cell, 2018, 30, 1-2.	6.6	5
30	Highâ€throughput sequencing of the chloroplast and mitochondrion of Chlamydomonas reinhardtii to generate improved de novo assemblies, analyze expression patterns and transcript speciation, and evaluate diversity among laboratory strains and wild isolates. Plant Journal, 2018, 93, 545-565.	5.7	90
31	Thank You, Editors and Reviewers of The Plant Cell. Plant Cell, 2018, 30, 2873-2879.	6.6	0
32	A Gelatin Microdroplet Platform for Highâ€Throughput Sorting of Hyperproducing Singleâ€Cellâ€Derived Microalgal Clones. Small, 2018, 14, e1803315.	10.0	52
33	In situ architecture of the algal nuclear pore complex. Nature Communications, 2018, 9, 2361.	12.8	107
34	Chromosome-level genome assembly and transcriptome of the green alga Chromochloris zofingiensis illuminates astaxanthin production. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E4296-E4305.	7.1	131
35	Chlamydomonas reinhardtii LFO1 Is an IsdG Family Heme Oxygenase. MSphere, 2017, 2, .	2.9	15
36	Bilin-Dependent Photoacclimation in Chlamydomonas reinhardtii. Plant Cell, 2017, 29, 2711-2726.	6.6	36

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37	Journal Impact: Brave New World. Plant Cell, 2017, 29, 2071-2074.	6.6	0
38	Regulating cellular trace metal economy in algae. Current Opinion in Plant Biology, 2017, 39, 88-96.	7.1	52
39	Endoplasmic reticulumâ€“mitochondria junction is required for iron homeostasis. Journal of Biological Chemistry, 2017, 292, 13197-13204.	3.4	40
40	Algae as nutritional and functional food sources: revisiting our understanding. Journal of Applied Phycology, 2017, 29, 949-982.	2.8	984
41	Thank You, Editors and Reviewers of The Plant Cell. Plant Cell, 2017, 29, 2941-2947.	6.6	0
42	A bioactive peptide amidating enzyme is required for ciliogenesis. ELife, 2017, 6, .	6.0	28
43	Genome and methylome of the oleaginous diatom Cyclotella cryptica reveal genetic flexibility toward a high lipid phenotype. Biotechnology for Biofuels, 2016, 9, 258.	6.2	87
44	Ni induces the CRR1-dependent regulon revealing overlap and distinction between hypoxia and Cu deficiency responses in Chlamydomonas reinhardtii. Metallomics, 2016, 8, 679-691.	2.4	27
45	Genetically Programmed Changes in Photosynthetic Cofactor Metabolism in Copper-deficient Chlamydomonas. Journal of Biological Chemistry, 2016, 291, 19118-19131.	3.4	13
46	The Plant Cell Begins Opt-in Publishing of Peer Review Reports. Plant Cell, 2016, 28, 2343-2343.	6.6	2
47	Exploiting algal NADPH oxidase for biophotovoltaic energy. Plant Biotechnology Journal, 2016, 14, 22-28.	8.3	37
48	Early eukaryotic origins for cilia-associated bioactive peptide amidating activity. Journal of Cell Science, 2016, 129, 943-56.	2.0	24
49	Copper status of exposed microorganisms influences susceptibility to metallic nanoparticles. Environmental Toxicology and Chemistry, 2016, 35, 1148-1158.	4.3	7
50	Genomeâ€wide analysis on Chlamydomonas reinhardtii reveals the impact of hydrogen peroxide on protein stress responses and overlap with other stress transcriptomes. Plant Journal, 2015, 84, 974-988.	5.7	55
51	The Plant Cell in the New Age of Scientific Publishing. Plant Cell, 2015, 27, 303-305.	6.6	2
52	Dissecting the contributions of <scp>GC</scp> content and codon usage to gene expression in the model alga Chlamydomonas reinhardtii. Plant Journal, 2015, 84, 704-717.	5.7	113
53	The Thylakoid Membrane Protein CGL160 Supports CF1CF0 ATP Synthase Accumulation in Arabidopsis thaliana. PLoS ONE, 2015, 10, e0121658.	2.5	29
54	A Dedicated Type II NADPH Dehydrogenase Performs the Penultimate Step in the Biosynthesis of Vitamin K1 in Synechocystis and Arabidopsis. Plant Cell, 2015, 27, 1730-1741.	6.6	50

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55	Identification of Coq11, a New Coenzyme Q Biosynthetic Protein in the CoQ-Synthome in Saccharomyces cerevisiae. Journal of Biological Chemistry, 2015, 290, 7517-7534.	3.4	65
56	The Plant CellIntroduces Breakthrough Reports: A New Forum for Cutting-Edge Plant Research. Plant Cell, 2015, , tpc.15.00862.	6.6	1
57	Copper economy in Chlamydomonas : Prioritized allocation and reallocation of copper to respiration vs. photosynthesis. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 2644-2651.	7.1	79
58	PHOTOSYSTEM II PROTEIN33, a Protein Conserved in the Plastid Lineage, Is Associated with the Chloroplast Thylakoid Membrane and Provides Stability to Photosystem II Supercomplexes in Arabidopsis. Plant Physiology, 2015, 167, 481-492.	4.8	46
59	Activation of Autophagy by Metals in Chlamydomonas reinhardtii. Eukaryotic Cell, 2015, 14, 964-973.	3.4	29
60	Redesigning photosynthesis to sustainably meet global food and bioenergy demand. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8529-8536.	7.1	751
61	Dynamic changes in the transcriptome and methylome of Chlamydomonas reinhardtii throughout its life cycle. Plant Physiology, 2015, 169, pp.00861.2015.	4.8	51
62	Chlamydomonas Genome Resource for Laboratory Strains Reveals a Mosaic of Sequence Variation, Identifies True Strain Histories, and Enables Strain-Specific Studies. Plant Cell, 2015, 27, 2335-2352.	6.6	102
63	Critical role ofChlamydomonas reinhardtiiferredoxin-5 in maintaining membrane structure and dark metabolism. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 14978-14983.	7.1	58
64	High-Resolution Profiling of a Synchronized Diurnal Transcriptome from Chlamydomonas reinhardtii Reveals Continuous Cell and Metabolic Differentiation. Plant Cell, 2015, 27, 2743-69.	6.6	195
65	Interpretation of the Genome in Synchronized Chlamydomonas Cells. FASEB Journal, 2015, 29, 485.1.	0.5	0
66	Evolution of a plant-specific copper chaperone family for chloroplast copper homeostasis. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E5480-7.	7.1	57
67	A Thylakoid Membrane Protein Harboring a DnaJ-type Zinc Finger Domain Is Required for Photosystem I Accumulation in Plants. Journal of Biological Chemistry, 2014, 289, 30657-30667.	3.4	64
68	Subcellular metal imaging identifies dynamic sites of Cu accumulation in Chlamydomonas. Nature Chemical Biology, 2014, 10, 1034-1042.	8.0	143
69	The Path to Triacylglyceride Obesity in the sta6 Strain of Chlamydomonas reinhardtii. Eukaryotic Cell, 2014, 13, 591-613.	3.4	143
70	Lysosome-related Organelles as Mediators of Metal Homeostasis. Journal of Biological Chemistry, 2014, 289, 28129-28136.	3.4	114
71	RBF1, a Plant Homolog of the Bacterial Ribosome-Binding Factor RbfA, Acts in Processing of the Chloroplast 16S Ribosomal RNA. Plant Physiology, 2014, 164, 201-215.	4.8	48
72	The Chlamydomonas genome project: a decade on. Trends in Plant Science, 2014, 19, 672-680.	8.8	145

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73	Nitrogen-Sparing Mechanisms in Chlamydomonas Affect the Transcriptome, the Proteome, and Photosynthetic Metabolism. Plant Cell, 2014, 26, 1410-1435.	6.6	314
74	Phosphoprotein SAK1 is a regulator of acclimation to singlet oxygen in Chlamydomonas reinhardtii. ELife, 2014, 3, e02286.	6.0	45
75	Systems-Level Analysis of Nitrogen Starvation-Induced Modifications of Carbon Metabolism in a Chlamydomonas reinhardtii Starchless Mutant. Plant Cell, 2013, 25, 4305-4323.	6.6	176
76	Functional Modeling Identifies Paralogous Solanesyl-diphosphate Synthases That Assemble the Side Chain of Plastoquinone-9 in Plastids. Journal of Biological Chemistry, 2013, 288, 27594-27606.	3.4	44
77	Remodeling of Membrane Lipids in Iron-starved Chlamydomonas. Journal of Biological Chemistry, 2013, 288, 30246-30258.	3.4	77
78	COPPER RESPONSE REGULATOR1â€“Dependent and â€“Independent Responses of the Chlamydomonas reinhardtii Transcriptome to Dark Anoxia. Plant Cell, 2013, 25, 3186-3211.	6.6	77
79	Iron economy in Chlamydomonas reinhardtii. Frontiers in Plant Science, 2013, 4, 337.	3.6	65
80	The Proteome of Copper, Iron, Zinc, and Manganese Micronutrient Deficiency in Chlamydomonas reinhardtii. Molecular and Cellular Proteomics, 2013, 12, 65-86.	3.8	85
81	Retrograde bilin signaling enables Chlamydomonas greening and phototrophic survival. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 3621-3626.	7.1	107
82	Zinc Deficiency Impacts CO2 Assimilation and Disrupts Copper Homeostasis in Chlamydomonas reinhardtii. Journal of Biological Chemistry, 2013, 288, 10672-10683.	3.4	72
83	Fe Sparing and Fe Recycling Contribute to Increased Superoxide Dismutase Capacity in Iron-Starved Chlamydomonas reinhardtii. Plant Cell, 2012, 24, 2649-2665.	6.6	82
84	Impact of Oxidative Stress on Ascorbate Biosynthesis in Chlamydomonas via Regulation of the VTC2 Gene Encoding a GDP-l-galactose Phosphorylase. Journal of Biological Chemistry, 2012, 287, 14234-14245.	3.4	93
85	Systems and Trans-System Level Analysis Identifies Conserved Iron Deficiency Responses in the Plant Lineage. Plant Cell, 2012, 24, 3921-3948.	6.6	142
86	Transcriptome Sequencing Identifies SPL7-Regulated Copper Acquisition Genes FRO4/FRO5 and the Copper Dependence of Iron Homeostasis in Arabidopsis. Plant Cell, 2012, 24, 738-761.	6.6	286
87	TAG, Youâ€™re it! Chlamydomonas as a reference organism for understanding algal triacylglycerol accumulation. Current Opinion in Biotechnology, 2012, 23, 352-363.	6.6	291
88	Elemental Economy. Advances in Microbial Physiology, 2012, 60, 91-210.	2.4	180
89	Transcriptome-Wide Changes in Chlamydomonas reinhardtii Gene Expression Regulated by Carbon Dioxide and the CO₂-Concentrating Mechanism Regulator CIA5/CCM1. Plant Cell, 2012, 24, 1876-1893.	6.6	180
90	Three Acyltransferases and Nitrogen-responsive Regulator Are Implicated in Nitrogen Starvation-induced Triacylglycerol Accumulation in Chlamydomonas. Journal of Biological Chemistry, 2012, 287, 15811-15825.	3.4	379

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91	The ins and outs of algal metal transport. Biochimica Et Biophysica Acta - Molecular Cell Research, 2012, 1823, 1531-1552.	4.1	173
92	A revised mineral nutrient supplement increases biomass and growth rate in Chlamydomonas reinhardtii. Plant Journal, 2011, 66, 770-780.	5.7	282
93	Algal Functional Annotation Tool: a web-based analysis suite to functionally interpret large gene lists using integrated annotation and expression data. BMC Bioinformatics, 2011, 12, 282.	2.6	84
94	The CRR1 Nutritional Copper Sensor in Chlamydomonas Contains Two Distinct Metal-Responsive Domains Â Â. Plant Cell, 2011, 22, 4098-4113.	6.6	93
95	The GreenCut2 Resource, a Phylogenomically Derived Inventory of Proteins Specific to the Plant Lineage. Journal of Biological Chemistry, 2011, 286, 21427-21439.	3.4	113
96	Systems Biology Approach in Chlamydomonas Reveals Connections between Copper Nutrition and Multiple Metabolic Steps Â Â. Plant Cell, 2011, 23, 1273-1292.	6.6	204
97	Enzymatic properties of the ferredoxin-dependent nitrite reductase from Chlamydomonas reinhardtii. Evidence for hydroxylamine as a late intermediate in ammonia production. Photosynthesis Research, 2010, 103, 67-77.	2.9	35
98	Phylogenomic analysis of the Chlamydomonas genome unmasks proteins potentially involved in photosynthetic function and regulation. Photosynthesis Research, 2010, 106, 3-17.	2.9	51
99	Trophic status of Chlamydomonas reinhardtii influences the impact of iron deficiency on photosynthesis. Photosynthesis Research, 2010, 105, 39-49.	2.9	80
100	RNA-Seq Analysis of Sulfur-Deprived Chlamydomonas Cells Reveals Aspects of Acclimation Critical for Cell Survival. Plant Cell, 2010, 22, 2058-2084.	6.6	253
101	The Elements of Plant Micronutrients. Plant Physiology, 2010, 154, 512-515.	4.8	69
102	His protects heme as it crosses the membrane. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 10069-10070.	7.1	16
103	Pattern of Expression and Substrate Specificity of Chloroplast Ferredoxins from Chlamydomonas reinhardtii. Journal of Biological Chemistry, 2009, 284, 25867-25878.	3.4	122
104	Two Chlamydomonas CTR Copper Transporters with a Novel Cys-Met Motif Are Localized to the Plasma Membrane and Function in Copper Assimilation. Plant Cell, 2009, 21, 928-943.	6.6	94
105	A Ferroxidase Encoded by FOX1 Contributes to Iron Assimilation under Conditions of Poor Iron Nutrition in Chlamydomonas. Eukaryotic Cell, 2008, 7, 541-545.	3.4	33
106	FER1 and FER2 Encoding Two Ferritin Complexes in Chlamydomonas reinhardtii Chloroplasts Are Regulated by Iron. Genetics, 2008, 179, 137-147.	2.9	57
107	Manganese Deficiency in Chlamydomonas Results in Loss of Photosystem II and MnSOD Function, Sensitivity to Peroxides, and Secondary Phosphorus and Iron Deficiency. Plant Physiology, 2007, 143, 263-277.	4.8	149
108	FEA1 , FEA2 , and FRE1 , Encoding Two Homologous Secreted Proteins and a Candidate Ferrireductase, Are Expressed Coordinately with FOX1 and FTR1 in Iron-Deficient Chlamydomonas reinhardtii. Eukaryotic Cell, 2007, 6, 1841-1852.	3.4	121

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109	The Chlamydomonas Genome Reveals the Evolution of Key Animal and Plant Functions. Science, 2007, 318, 245-250.	12.6	2,354
110	Precious metal economy. Cell Metabolism, 2006, 4, 99-101.	16.2	4
111	Between a rock and a hard place: Trace element nutrition in Chlamydomonas. Biochimica Et Biophysica Acta - Molecular Cell Research, 2006, 1763, 578-594.	4.1	202
112	A regulator of nutritional copper signaling in Chlamydomonas is an SBP domain protein that recognizes the GTAC core of copper response element. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 18730-18735.	7.1	292
113	The Light Reactions: A Guide to Recent Acquisitions for the Picture Gallery. Plant Cell, 2005, 17, 648-663.	6.6	86
114	Genetic Dissection of Nutritional Copper Signaling in Chlamydomonas Distinguishes Regulatory and Target Genes. Genetics, 2004, 168, 795-807.	2.9	82
115	Arabidopsis CHL27, located in both envelope and thylakoid membranes, is required for the synthesis of protochlorophyllide. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 16119-16124.	7.1	195
116	Copper Response Element and Crr1-Dependent Ni 2+ -Responsive Promoter for Induced, Reversible Gene Expression in Chlamydomonas reinhardtii. Eukaryotic Cell, 2003, 2, 995-1002.	3.4	62
117	Essential Histidine and Tryptophan Residues in CcsA, a System II Polytopic Cytochrome c Biogenesis Protein. Journal of Biological Chemistry, 2003, 278, 2593-2603.	3.4	66
118	Reciprocal Expression of Two Candidate Di-Iron Enzymes Affecting Photosystem I and Light-Harvesting Complex Accumulation. Plant Cell, 2002, 14, 673-688.	6.6	136
119	Copper-Dependent Iron Assimilation Pathway in the Model Photosynthetic EukaryoteChlamydomonas reinhardtii. Eukaryotic Cell, 2002, 1, 736-757.	3.4	184
120	Adaptation to Fe-deficiency requires remodeling of the photosynthetic apparatus. EMBO Journal, 2002, 21, 6709-6720.	7.8	240
121	The Crd1 gene encodes a putative di-iron enzyme required for photosystem I accumulation in copper deficiency and hypoxia in Chlamydomonas reinhardtii. EMBO Journal, 2000, 19, 2139-2151.	7.8	170
122	Coordinate Copper- and Oxygen-responsive Cyc6 andCpx1 Expression in Chlamydomonas Is Mediated by the Same Element. Journal of Biological Chemistry, 2000, 275, 6080-6089.	3.4	114
123	ADAPTATION OF SCENEDESMUS OBLIQUUS (CHLOROPHYCEAE) TO COPPER-DEFICIENCY: TRANSCRIPTIONAL REGULATION OF PCY1 BUT NOT CPX1. Journal of Phycology, 1999, 35, 1253-1263.	2.3	3
124	Mmicular mechanisms of cytochromecbiogenesis: three distinct systems. Molecular Microbiology, 1998, 29, 383-396.	2.5	266
125	[18] Copper-responsive gene expression during adaptation to copper deficiency. Methods in Enzymology, 1998, 297, 263-279.	1.0	75
126	POSTTRANSLATIONAL ASSEMBLY OF PHOTOSYNTHETIC METALLOPROTEINS. Annual Review of Plant Biology, 1998, 49, 25-51.	14.3	80

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127	Genetic Analysis of Chloroplast c-Type Cytochrome Assembly in Chlamydomonas reinhardtii: One Chloroplast Locus and at Least Four Nuclear Loci Are Required for Heme Attachment. Genetics, 1998, 148, 681-692.	2.9	68
128	Biosynthesis of cytochrome f in Chiamydomonas reinhardtii: analysis of the pathway in gabaculine-treated cells and in the heme attachment mutant B6. Molecular Genetics and Genomics, 1995, 246, 156-165.	2.4	36
129	Two Copper-Responsive Elements Associated with the Chlamydomonas Cyc6 Gene Function as Targets for Transcriptional Activators. Plant Cell, 1995, 7, 623.	6.6	40
130	Degradation of Plastocyanin in Copper-deficient Chlamydomonas reinhardtii. Journal of Biological Chemistry, 1995, 270, 23504-23510.	3.4	77
131	In Vivo Competition between Plastocyanin and a Copper-Dependent Regulator of the Chlamydomonas reinhardtii Cytochrome c₆ Gene. Plant Physiology, 1992, 100, 319-326.	4.8	33
132	Heavy Metal-Activated Synthesis of Peptides in Chlamydomonas reinhardtii. Plant Physiology, 1992, 98, 127-136.	4.8	139
133	Synthesis and Turnover of the Chloroplast Coupling Factor 1 in Chlamydomonas reinhardi. Plant Physiology, 1984, 75, 781-787.	4.8	19
134	Isolation, purification, and characterization of coupling factor 1 from Chlamydomonas reinhardi. Biochemistry, 1981, 20, 5476-5482.	2.5	62