

Wataru Sakamoto

List of Publications by Year in descending order

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130
papers

6,722
citations

50276

46
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71685

76
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133
all docs

133
docs citations

133
times ranked

8040
citing authors

#	ARTICLE	IF	CITATIONS
1	Coordinated Regulation and Complex Formation of YELLOW VARIEGATED1 and YELLOW VARIEGATED2, Chloroplastic FtsH Metalloproteases Involved in the Repair Cycle of Photosystem II in Arabidopsis Thylakoid Membranes. <i>Plant Cell</i> , 2003, 15, 2843-2855.	6.6	276
2	PROTEIN DEGRADATION MACHINERIES IN PLASTIDS. <i>Annual Review of Plant Biology</i> , 2006, 57, 599-621.	18.7	202
3	Highly Oxidized Peroxisomes Are Selectively Degraded via Autophagy in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2013, 25, 4967-4983.	6.6	195
4	The Variegated Mutants Lacking Chloroplastic FtsHs Are Defective in D1 Degradation and Accumulate Reactive Oxygen Species. <i>Plant Physiology</i> , 2009, 151, 1790-1801.	4.8	189
5	The VAR1 locus of <i>Arabidopsis</i> encodes a chloroplastic FtsH and is responsible for leaf variegation in the mutant alleles. <i>Genes To Cells</i> , 2002, 7, 769-780.	1.2	185
6	The YELLOW VARIEGATED (VAR2) Locus Encodes a Homologue of FtsH, an ATP-Dependent Protease in <i>Arabidopsis</i> . <i>Plant and Cell Physiology</i> , 2000, 41, 1334-1346.	3.1	184
7	Widespread Endogenization of Genome Sequences of Non-Retroviral RNA Viruses into Plant Genomes. <i>PLoS Pathogens</i> , 2011, 7, e1002146.	4.7	173
8	Functional characterization of key structural genes in rice flavonoid biosynthesis. <i>Planta</i> , 2008, 228, 1043-1054.	3.2	160
9	The Balance between Protein Synthesis and Degradation in Chloroplasts Determines Leaf Variegation in <i>Arabidopsis</i> yellow variegated Mutants. <i>Plant Cell</i> , 2007, 19, 1313-1328.	6.6	149
10	Cooperative D1 Degradation in the Photosystem II Repair Mediated by Chloroplastic Proteases in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2012, 159, 1428-1439.	4.8	147
11	Altered mitochondrial gene expression in a maternal distorted leaf mutant of <i>Arabidopsis</i> induced by chloroplast mutator.. <i>Plant Cell</i> , 1996, 8, 1377-1390.	6.6	135
12	Chloroplast Biogenesis: Control of Plastid Development, Protein Import, Division and Inheritance. <i>The Arabidopsis Book</i> , 2008, 6, e0110.	0.5	129
13	Functional divergence of the TFL1-like gene family in <i>Arabidopsis</i> revealed by characterization of a novel homologue. <i>Genes To Cells</i> , 2001, 6, 327-336.	1.2	128
14	Protein Quality Control in Chloroplasts: A Current Model of D1 Protein Degradation in the Photosystem II Repair Cycle. <i>Journal of Biochemistry</i> , 2009, 146, 463-469.	1.7	127
15	Physical interaction between peroxisomes and chloroplasts elucidated by in situ laser analysis. <i>Nature Plants</i> , 2015, 1, 15035.	9.3	118
16	Chloroplast Proteases: Updates on Proteolysis within and across Suborganellar Compartments. <i>Plant Physiology</i> , 2016, 171, 2280-2293.	4.8	118
17	The Purple leaf (Pl) Locus of Rice: the Plw Allele has a Complex Organization and Includes Two Genes Encoding Basic Helix-Loop-Helix Proteins Involved in Anthocyanin Biosynthesis. <i>Plant and Cell Physiology</i> , 2001, 42, 982-991.	3.1	117
18	FtsH Protease in the Thylakoid Membrane: Physiological Functions and the Regulation of Protease Activity. <i>Frontiers in Plant Science</i> , 2018, 9, 855.	3.6	117

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19	In vivo analysis of <i>Chlamydomonas</i> chloroplast <i>petD</i> gene expression using stable transformation of beta-glucuronidase translational fusions.. Proceedings of the National Academy of Sciences of the United States of America, 1993, 90, 497-501.	7.1	114
20	Leaf-variegated mutations and their responsible genes in <i>Arabidopsis thaliana</i> .. Genes and Genetic Systems, 2003, 78, 1-9.	0.7	114
21	Essential Role of VIPP1 in Chloroplast Envelope Maintenance in <i>Arabidopsis</i> . Plant Cell, 2012, 24, 3695-3707.	6.6	107
22	White Leaf Sectors in yellow variegated2 Are Formed by Viable Cells with Undifferentiated Plastids. Plant Physiology, 2007, 144, 952-960.	4.8	104
23	<i>NYC4</i> , the rice ortholog of <i>Arabidopsis</i> <i>THF1</i> , is involved in the degradation of chlorophyll "a" protein complexes during leaf senescence. Plant Journal, 2013, 74, 652-662.	5.7	98
24	Amyloplast-Localized SUBSTANDARD STARCH GRAIN4 Protein Influences the Size of Starch Grains in Rice Endosperm. Plant Physiology, 2014, 164, 623-636.	4.8	98
25	Selective Elimination of Membrane-Damaged Chloroplasts via Microautophagy. Plant Physiology, 2018, 177, 1007-1026.	4.8	91
26	Putative phospholipid hydroperoxide glutathione peroxidase gene from <i>Arabidopsis thaliana</i> induced by oxidative stress.. Genes and Genetic Systems, 1997, 72, 311-316.	0.7	90
27	Essentials of Proteolytic Machineries in Chloroplasts. Molecular Plant, 2017, 10, 4-19.	8.3	90
28	Function of the <i>Chlamydomonas reinhardtii</i> <i>petD</i> 5' untranslated region in regulating the accumulation of subunit IV of the cytochrome b6/f complex. Plant Journal, 1994, 6, 503-512.	5.7	89
29	<i>Arabidopsis</i> ELONGATED MITOCHONDRIA1 Is Required for Localization of DYNAMIN-RELATED PROTEIN3A to Mitochondrial Fission Sites. Plant Cell, 2008, 20, 1555-1566.	6.6	89
30	The lack of mitochondrial <i>AtFtsH4</i> protease alters <i>Arabidopsis</i> leaf morphology at the late stage of rosette development under short-day photoperiod. Plant Journal, 2009, 59, 685-699.	5.7	80
31	A mutation of the <i>CRUMPLED LEAF</i> gene that encodes a protein localized in the outer envelope membrane of plastids affects the pattern of cell division, cell differentiation, and plastid division in <i>Arabidopsis</i> . Plant Journal, 2004, 38, 448-459.	5.7	79
32	A single gene of chloroplast origin codes for mitochondrial and chloroplastic methionyl-tRNA synthetase in <i>Arabidopsis thaliana</i> . Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 11014-11019.	7.1	79
33	Structural basis for VIPP1 oligomerization and maintenance of thylakoid membrane integrity. Cell, 2021, 184, 3643-3659.e23.	28.9	76
34	Activation of the heterotrimeric G protein β -subunit <i>GPA1</i> suppresses the <i>ftsH</i> -mediated inhibition of chloroplast development in <i>Arabidopsis</i> . Plant Journal, 2009, 58, 1041-1053.	5.7	73
35	A Rapid, Direct Observation Method to Isolate Mutants with Defects in Starch Grain Morphology in Rice. Plant and Cell Physiology, 2010, 51, 728-741.	3.1	69
36	<i>Arabidopsis thaliana</i> vegetative storage protein (VSP) genes: gene organization and tissue-specific expression. Plant Molecular Biology, 1998, 38, 565-576.	3.9	68

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37	An EAR-Dependent Regulatory Module Promotes Male Germ Cell Division and Sperm Fertility in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 2098-2113.	6.6	67
38	Influence of Chloroplastic Photo-Oxidative Stress on Mitochondrial Alternative Oxidase Capacity and Respiratory Properties: A Case Study with <i>Arabidopsis</i> yellow variegated 2. <i>Plant and Cell Physiology</i> , 2008, 49, 592-603.	3.1	66
39	New Insights into the Types and Function of Proteases in Plastids. <i>International Review of Cell and Molecular Biology</i> , 2010, 280, 185-218.	3.2	66
40	Arrested Differentiation of Proplastids into Chloroplasts in Variegated Leaves Characterized by Plastid Ultrastructure and Nucleoid Morphology. <i>Plant and Cell Physiology</i> , 2009, 50, 2069-2083.	3.1	62
41	Identification and Characterization of High Molecular Weight Complexes Formed by Matrix AAA Proteases and Prohibitins in Mitochondria of <i>Arabidopsis thaliana</i> . <i>Journal of Biological Chemistry</i> , 2010, 285, 12512-12521.	3.4	62
42	Amyloplast Membrane Protein SUBSTANDARD STARCH GRAIN6 Controls Starch Grain Size in Rice Endosperm. <i>Plant Physiology</i> , 2016, 170, 1445-1459.	4.8	61
43	Multiple Intracellular Locations of Lon Protease in <i>Arabidopsis</i> : Evidence for the Localization of AtLon4 to Chloroplasts. <i>Plant and Cell Physiology</i> , 2007, 48, 881-885.	3.1	60
44	Phosphorylation of photosystem II core proteins prevents undesirable cleavage of D1 and contributes to the fine-tuned repair of photosystem II. <i>Plant Journal</i> , 2014, 79, 312-321.	5.7	60
45	Nucleases in higher plants and their possible involvement in DNA degradation during leaf senescence. <i>Journal of Experimental Botany</i> , 2014, 65, 3835-3843.	4.8	58
46	The FtsH Protease Heterocomplex in <i>Arabidopsis</i> : Dispensability of Type-B Protease Activity for Proper Chloroplast Development. <i>Plant Cell</i> , 2010, 22, 3710-3725.	6.6	57
47	Chloroplast DNA Dynamics: Copy Number, Quality Control and Degradation. <i>Plant and Cell Physiology</i> , 2018, 59, 1120-1127.	3.1	56
48	A Conserved, Mg ²⁺ -Dependent Exonuclease Degrades Organelle DNA during <i>Arabidopsis</i> Pollen Development. <i>Plant Cell</i> , 2011, 23, 1608-1624.	6.6	53
49	Vegetative and Sperm Cell-Specific Aquaporins of <i>Arabidopsis</i> Highlight the Vacuolar Equipment of Pollen and Contribute to Plant Reproduction. <i>Plant Physiology</i> , 2014, 164, 1697-1706.	4.8	50
50	VIPP1 Has a Disordered C-Terminal Tail Necessary for Protecting Photosynthetic Membranes against Stress. <i>Plant Physiology</i> , 2016, 171, 1983-1995.	4.8	50
51	The Model Plant <i>Medicago truncatula</i> Exhibits Biparental Plastid Inheritance. <i>Plant and Cell Physiology</i> , 2008, 49, 81-91.	3.1	46
52	Comparative transcriptome analysis of green/white variegated sectors in <i>Arabidopsis</i> yellow variegated2: responses to oxidative and other stresses in white sectors. <i>Journal of Experimental Botany</i> , 2010, 61, 2433-2445.	4.8	46
53	Mitochondrial Dynamics in Plant Male Gametophyte Visualized by Fluorescent Live Imaging. <i>Plant and Cell Physiology</i> , 2008, 49, 1074-1083.	3.1	44
54	Plant mitochondrial rhomboid, AtRBL12, has different substrate specificity from its yeast counterpart. <i>Plant Molecular Biology</i> , 2008, 68, 159-171.	3.9	43

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55	FtsH proteases in chloroplasts and cyanobacteria. <i>Physiologia Plantarum</i> , 2005, 123, 386-390.	5.2	41
56	The Photosystem II Repair Cycle Requires FtsH Turnover through the EngA GTPase. <i>Plant Physiology</i> , 2018, 178, 596-611.	4.8	41
57	Ion gradients in xylem exudate and guttation fluid related to tissue ion levels along primary leaves of barley. <i>Plant, Cell and Environment</i> , 2013, 36, 1826-1837.	5.7	39
58	D1 fragmentation in photosystem II repair caused by photo-damage of a two-step model. <i>Photosynthesis Research</i> , 2015, 126, 409-416.	2.9	39
59	Mitochondrial Localization of AtOXA1, an Arabidopsis Homologue of Yeast Oxa1p Involved in the Insertion and Assembly of Protein Complexes in Mitochondrial Inner Membrane. <i>Plant and Cell Physiology</i> , 2000, 41, 1157-1163.	3.1	38
60	Organelle DNA degradation contributes to the efficient use of phosphate in seed plants. <i>Nature Plants</i> , 2018, 4, 1044-1055.	9.3	38
61	Impaired PSII Proteostasis Promotes Retrograde Signaling via Salicylic Acid. <i>Plant Physiology</i> , 2019, 180, 2182-2197.	4.8	38
62	Functional complementation of an oxa1- yeast mutation identifies an Arabidopsis thaliana cDNA involved in the assembly of respiratory complexes. <i>Plant Journal</i> , 1997, 12, 1319-1327.	5.7	37
63	Isolation of an Arabidopsis thaliana cDNA by complementation of a yeast abc1 deletion mutant deficient in complex III respiratory activity. <i>Gene</i> , 1998, 221, 117-125.	2.2	37
64	The Non-Mendelian Green Cotyledon Gene in Soybean Encodes a Small Subunit of Photosystem II. <i>Plant Physiology</i> , 2017, 173, 2138-2147.	4.8	37
65	Possible function of VIPP1 in thylakoids. <i>Plant Signaling and Behavior</i> , 2013, 8, e22860.	2.4	36
66	Visualization of Plastids in Pollen Grains: Involvement of FtsZ1 in Pollen Plastid Division. <i>Plant and Cell Physiology</i> , 2009, 50, 904-908.	3.1	35
67	Allelic characterization of the leaf-variegated mutation var2 identifies the conserved amino acid residues of FtsH that are important for ATP hydrolysis and proteolysis. <i>Plant Molecular Biology</i> , 2004, 56, 705-716.	3.9	34
68	A Phylogenetic Re-evaluation of Morphological Variations of Starch Grains among Poaceae Species. <i>Journal of Applied Glycoscience</i> (1999), 2013, 60, 37-44.	0.7	33
69	Different amounts of DNA in each mitochondrion in rice root. <i>Genes and Genetic Systems</i> , 2006, 81, 215-218.	0.7	32
70	Possible function of VIPP1 in maintaining chloroplast membranes. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2015, 1847, 831-837.	1.0	32
71	The strange evolutionary history of plant mitochondrial tRNAs and their aminoacyl-tRNA synthetases. <i>Journal of Molecular Evolution</i> , 1999, 90, 333-337.		29
72	Variegated Tobacco Leaves Generated by Chloroplast FtsH Suppression: Implication of FtsH Function in the Maintenance of Thylakoid Membranes. <i>Plant and Cell Physiology</i> , 2012, 53, 391-404.	3.1	28

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73	Isolation of mutants with aberrant mitochondrial morphology from <i>Arabidopsis thaliana</i> . <i>Genes and Genetic Systems</i> , 2004, 79, 301-305.	0.7	26
74	Reduction in amounts of mitochondrial DNA in the sperm cells as a mechanism for maternal inheritance in <i>Hordeum vulgare</i> . <i>Planta</i> , 2002, 216, 235-244.	3.2	25
75	RAD-seq-Based High-Density Linkage Map Construction and QTL Mapping of Biomass-Related Traits in Sorghum using the Japanese Landrace Takakibi NOG. <i>Plant and Cell Physiology</i> , 2020, 61, 1262-1272.	3.1	25
76	Rice CYO1, an ortholog of <i>Arabidopsis thaliana</i> cotyledon chloroplast biogenesis factor AtCYO1, is expressed in leaves and involved in photosynthetic performance. <i>Journal of Plant Physiology</i> , 2016, 207, 78-83.	3.5	23
77	Impairment of Lhca4, a subunit of LHCl, causes high accumulation of chlorophyll and the stay-green phenotype in rice. <i>Journal of Experimental Botany</i> , 2018, 69, 1027-1035.	4.8	22
78	Dielectric relaxation of vegetable-based polyurethane. <i>Journal of Materials Science</i> , 2003, 38, 1465-1470.	3.7	21
79	Chemically Induced Expression of Rice OSB2 under the Control of the OsPR1.1 Promoter Confers Increased Anthocyanin Accumulation in Transgenic Rice. <i>Journal of Agricultural and Food Chemistry</i> , 2007, 55, 1241-1247.	5.2	21
80	The Rubisco Chaperone BSD2 May Regulate Chloroplast Coverage in Maize Bundle Sheath Cells. <i>Plant Physiology</i> , 2017, 175, 1624-1633.	4.8	21
81	VIPP1 Involved in Chloroplast Membrane Integrity Has GTPase Activity in Vitro. <i>Plant Physiology</i> , 2018, 177, 328-338.	4.8	21
82	Tissue-specific organelle DNA degradation mediated by DPD1 exonuclease. <i>Plant Signaling and Behavior</i> , 2011, 6, 1391-1393.	2.4	18
83	Protection of Chloroplast Membranes by VIPP1 Rescues Aberrant Seedling Development in <i>Arabidopsis nyc1</i> Mutant. <i>Frontiers in Plant Science</i> , 2016, 7, 533.	3.6	18
84	Mutations defective in ribonucleotide reductase activity interfere with pollen plastid DNA degradation mediated by DPD1 exonuclease. <i>Plant Journal</i> , 2012, 70, 637-649.	5.7	17
85	Isolation and characterization of cDNA clones corresponding to the genes expressed preferentially in floral organs of <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1996, 32, 759-765.	3.9	16
86	MOLECULAR DIVERGENCE AND CHARACTERIZATION OF TWO CHLOROPLAST DIVISION GENES, <i>FTSZ1</i> AND <i>FTSZ2</i> , IN THE UNICELLULAR GREEN ALGA <i>NANNOCHLORIS BACILLARIS</i> (CHLOROPHYTA). <i>Journal of Phycology</i> , 2004, 40, 546-556.	2.3	16
87	Photosynthetic Responses to High Temperature and Strong Light Suggest Potential Post-flowering Drought Tolerance of Sorghum Japanese Landrace Takakibi. <i>Plant and Cell Physiology</i> , 2019, 60, 2086-2099.	3.1	15
88	Plant autophagy is responsible for peroxisomal transition and plays an important role in the maintenance of peroxisomal quality. <i>Autophagy</i> , 2014, 10, 936-937.	9.1	14
89	A Mutation in GIANT CHLOROPLAST Encoding a PARC6 Homolog Affects Spikelet Fertility in Rice. <i>Plant and Cell Physiology</i> , 2015, 56, 977-991.	3.1	14
90	Phosphorylation of the Chloroplastic Metalloprotease FtsH in <i>Arabidopsis</i> Characterized by Phos-Tag SDS-PAGE. <i>Frontiers in Plant Science</i> , 2019, 10, 1080.	3.6	14

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91	Genetic analysis of chlorophyll synthesis and degradation regulated by BALANCE of CHLOROPHYLL METABOLISM. <i>Plant Physiology</i> , 2022, 189, 419-432.	4.8	14
92	Distribution and quantitative variation of mitochondrial plasmid-like DNAs in cultivated rice (<i>Oryza</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.0	13
93	TERMINAL FLOWER 1-like genes in Brassica species. <i>Plant Science</i> , 1999, 142, 155-162.	3.6	13
94	Geometrical Formation of Compound Starch Grains in Rice Implements Voronoi Diagram. <i>Plant and Cell Physiology</i> , 2015, 56, pcv123.	3.1	13
95	Overexpression of BUNDLE SHEATH DEFECTIVE 2 improves the efficiency of photosynthesis and growth in <i>Arabidopsis</i> . <i>Plant Journal</i> , 2020, 102, 129-137.	5.7	13
96	Analysis of homology of small plasmid-like mitochondrial DNAs in the different cytoplasmic male sterile strains in rice. <i>Japanese Journal of Genetics</i> , 1989, 64, 49-56.	1.0	12
97	Analysis of mitochondrial DNAs from <i>Oryza glaberrima</i> and its cytoplasmic substituted line for <i>Oryza sativa</i> associated with cytoplasmic male sterility. <i>Japanese Journal of Genetics</i> , 1990, 65, 1-6.	1.0	12
98	Characterization of a Flower-Specific Gene Encoding a Putative Myrosinase Binding Protein in <i>Arabidopsis thaliana</i> . <i>Plant and Cell Physiology</i> , 1999, 40, 1287-1296.	3.1	12
99	High temperature causes breakdown of S haplotype-dependent stigmatic self-incompatibility in self-incompatible <i>Arabidopsis thaliana</i> . <i>Journal of Experimental Botany</i> , 2019, 70, 5745-5751.	4.8	10
100	Targeted proteome analysis of microalgae under high-light conditions by optimized protein extraction of photosynthetic organisms. <i>Journal of Bioscience and Bioengineering</i> , 2019, 127, 394-402.	2.2	10
101	Mutations in a Golden2-Like Gene Cause Reduced Seed Weight in Barley albino lemma 1 Mutants. <i>Plant and Cell Physiology</i> , 2021, 62, 447-457.	3.1	10
102	Isolation and molecular characterization of rbcS in the unicellular green alga <i>Nannochloris bacillaris</i> (Chlorophyta, Trebouxiophyceae). <i>Phycological Research</i> , 2005, 53, 67-76.	1.6	10
103	The BnALMT1 Protein that is an Aluminum-Activated Malate Transporter is Localized in the Plasma Membrane. <i>Plant Signaling and Behavior</i> , 2007, 2, 255-257.	2.4	9
104	In situ RNA hybridization using Technovit resin in <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology Reporter</i> , 1999, 17, 43-51.	1.8	8
105	Genetic dissection of QTLs associated with spikelet-related traits and grain size in sorghum. <i>Scientific Reports</i> , 2021, 11, 9398.	3.3	8
106	Isolation and molecular characterization of rbcS in the unicellular green alga <i>Nannochloris bacillaris</i> (Chlorophyta, Trebouxiophyceae). <i>Phycological Research</i> , 2005, 53, 67-76.	1.6	7
107	Phototropin and photosynthesis dependent mitochondrial positioning in <i>Arabidopsis thaliana</i> mesophyll cells. <i>Journal of Integrative Plant Biology</i> , 2020, 62, 1352-1371.	8.5	7
108	Phos-tag-based approach to study protein phosphorylation in the thylakoid membrane. <i>Photosynthesis Research</i> , 2021, 147, 107-124.	2.9	7

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109	Maintaining the Chloroplast Redox Balance through the PGR5-Dependent Pathway and the Trx System Is Required for Light-Dependent Activation of Photosynthetic Reactions. <i>Plant and Cell Physiology</i> , 2022, 63, 92-103.	3.1	7
110	Highly efficient visual selection of transgenic rice plants using green fluorescent protein or anthocyanin synthetic genes. <i>Plant Biotechnology</i> , 2011, 28, 107-110.	1.0	6
111	Sorghum Ionomics Reveals the Functional <i>SbHMA3a</i> Allele that Limits Excess Cadmium Accumulation in Grains. <i>Plant and Cell Physiology</i> , 2022, 63, 713-728.	3.1	6
112	<i>DOMINANT AWN INHIBITOR</i> Encodes the ALOG Protein Originating from Gene Duplication and Inhibits AWN Elongation by Suppressing Cell Proliferation and Elongation in Sorghum. <i>Plant and Cell Physiology</i> , 2022, 63, 901-918.	3.1	6
113	Linkage analysis of the nuclear homologues of mitochondrial plasmid-like DNAs in rice.. <i>Japanese Journal of Genetics</i> , 1991, 66, 597-607.	1.0	5
114	Isolation and characterization of Ty1/copia-like retrotransposons in mung bean (<i>Vigna radiata</i>). <i>Journal of Plant Research</i> , 2007, 120, 323-328.	2.4	5
115	Localization and expression of serine racemase in <i>Arabidopsis thaliana</i> . <i>Amino Acids</i> , 2009, 36, 587-590.	2.7	5
116	Plastid Protein Degradation During Leaf Development and Senescence: Role of Proteases and Chaperones. <i>Advances in Photosynthesis and Respiration</i> , 2013, , 453-477.	1.0	5
117	NB-LRR-encoding genes conferring susceptibility to organophosphate pesticides in sorghum. <i>Scientific Reports</i> , 2021, 11, 19828.	3.3	5
118	Reactive oxygen species derived from impaired quality control of Photosystem II are irrelevant to plasma-membrane NADPH oxidases. <i>Plant Signaling and Behavior</i> , 2010, 5, 264-266.	2.4	4
119	Overexpression of the protein disulfide isomerase AtCYO1 in chloroplasts slows dark-induced senescence in <i>Arabidopsis</i> . <i>BMC Plant Biology</i> , 2018, 18, 80.	3.6	4
120	Plastid Proteases. , 2014, , 359-389.		4
121	Possible compensatory role among chloroplast proteases under excess-light stress condition. <i>Plant Signaling and Behavior</i> , 2013, 8, e23198.	2.4	3
122	Taiwanâ€™Japan Plant Biology 2017 Spotlight Issue: From Light Signals/Signaling to Photosynthesis and Chloroplast Development. <i>Plant and Cell Physiology</i> , 2018, 59, 1099-1103.	3.1	2
123	Functional division of f-type and m-type thioredoxins to regulate the Calvin cycle and cyclic electron transport around photosystem I. <i>Journal of Plant Research</i> , 2022, , 1.	2.4	2
124	With Greetings and Hope for a Recoverable 2021: From the PCP Editor-In-Chief. <i>Plant and Cell Physiology</i> , 2021, 62, 219-221.	3.1	1
125	Cellular Dynamics: Cellular Systems in the Time Domain. <i>Plant Physiology</i> , 2018, 176, 12-15.	4.8	0
126	A 2020 Vision of the Next Four Yearsâ€™From the PCPâ€™s New Editor-in-Chief. <i>Plant and Cell Physiology</i> , 2020, 61, 671-672.	3.1	0

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127	Editorial Feature: Meet the PCP Editor-In-Chief“Wataru Sakamoto. Plant and Cell Physiology, 2021, 62, 222-223.	3.1	0
128	A Novel Link between Chloroplast Development and Stress Response Lessened by Leaf-Variegated Mutant. Advanced Topics in Science and Technology in China, 2013, , 669-673.	0.1	0
129	The Lattice-Like Structure Observed by Vipp1-GFP in Arabidopsis Chloroplasts. Advanced Topics in Science and Technology in China, 2013, , 394-397.	0.1	0
130	Distinctive in vitro ATP Hydrolysis Activity of AtVIPP1, a Chloroplastic ESCRT-III Superfamily Protein in Arabidopsis. Frontiers in Plant Science, 0, 13, .	3.6	0