

## List of Publications by Year in descending order

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81  
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36203

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docs citations

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times ranked

8659  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-Wide Analysis of Arabidopsis Pentatricopeptide Repeat Proteins Reveals Their Essential Role in Organelle Biogenesis[W]. <i>Plant Cell</i> , 2004, 16, 2089-2103.	3.1	1,132
2	Pentatricopeptide Repeat Proteins in Plants. <i>Annual Review of Plant Biology</i> , 2014, 65, 415-442.	8.6	842
3	Predotar: A tool for rapidly screening proteomes for N-terminal targeting sequences. <i>Proteomics</i> , 2004, 4, 1581-1590.	1.3	817
4	Pentatricopeptide repeat proteins: a socket set for organelle gene expression. <i>Trends in Plant Science</i> , 2008, 13, 663-670.	4.3	754
5	A Combinatorial Amino Acid Code for RNA Recognition by Pentatricopeptide Repeat Proteins. <i>PLoS Genetics</i> , 2012, 8, e1002910.	1.5	455
6	SUBA: the Arabidopsis Subcellular Database. <i>Nucleic Acids Research</i> , 2007, 35, D213-D218.	6.5	394
7	On the Expansion of the Pentatricopeptide Repeat Gene Family in Plants. <i>Molecular Biology and Evolution</i> , 2008, 25, 1120-1128.	3.5	329
8	Genome-Wide Analysis of mRNA Decay Rates and Their Determinants in <i>Arabidopsis thaliana</i> . <i>Plant Cell</i> , 2007, 19, 3418-3436.	3.1	296
9	Identification of the fertility restoration locus, <i>Rfo</i> , in radish, as a member of the pentatricopeptide repeat protein family. <i>EMBO Reports</i> , 2003, 4, 588-594.	2.0	291
10	Versatile Gene-Specific Sequence Tags for Arabidopsis Functional Genomics: Transcript Profiling and Reverse Genetics Applications. <i>Genome Research</i> , 2004, 14, 2176-2189.	2.4	282
11	SUBA3: a database for integrating experimentation and prediction to define the SUBcellular location of proteins in Arabidopsis. <i>Nucleic Acids Research</i> , 2012, 41, D1185-D1191.	6.5	272
12	Redefining the structural motifs that determine <i>scp</i> RNA binding and <i>scp</i> RNA editing by pentatricopeptide repeat proteins in land plants. <i>Plant Journal</i> , 2016, 85, 532-547.	2.8	267
13	The evolution of RNA editing and pentatricopeptide repeat genes. <i>New Phytologist</i> , 2011, 191, 37-47.	3.5	249
14	Rampant Gene Loss in the Underground Orchid <i>Rhizanthella gardneri</i> Highlights Evolutionary Constraints on Plastid Genomes. <i>Molecular Biology and Evolution</i> , 2011, 28, 2077-2086.	3.5	248
15	CLB19, a pentatricopeptide repeat protein required for editing of <i>rpoA</i> and <i>clpP</i> chloroplast transcripts. <i>Plant Journal</i> , 2008, 56, 590-602.	2.8	236
16	Pentatricopeptide Repeat Proteins with the DYW Motif Have Distinct Molecular Functions in RNA Editing and RNA Cleavage in <i>Arabidopsis</i> Chloroplasts. <i>Plant Cell</i> , 2009, 21, 146-156.	3.1	226
17	A hypothesis on the identification of the editing enzyme in plant organelles. <i>FEBS Letters</i> , 2007, 581, 4132-4138.	1.3	211
18	A Study of New <i>Arabidopsis</i> Chloroplast RNA Editing Mutants Reveals General Features of Editing Factors and Their Target Sites. <i>Plant Cell</i> , 2009, 21, 3686-3699.	3.1	179

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19	The Arabidopsis gene <i>YS1</i> encoding a DYW protein is required for editing of <i>rpoB</i> transcripts and the rapid development of chloroplasts during early growth. <i>Plant Journal</i> , 2009, 58, 82-96.	2.8	178
20	Sequence and transcript analysis of the Nco2.5 Ogura-specific fragment correlated with cytoplasmic male sterility in Brassica cybrids. <i>Molecular Genetics and Genomics</i> , 1992, 235, 340-348.	2.4	176
21	A rapid high-throughput method for the detection and quantification of RNA editing based on high-resolution melting of amplicons. <i>Nucleic Acids Research</i> , 2007, 35, e114.	6.5	167
22	The pentatricopeptide repeat gene <i>OTP51</i> with two LAGLIDADG motifs is required for the cis-splicing of plastid <i>yfc3</i> intron 2 in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 2008, 56, 157-168.	2.8	148
23	Plant RNA editing. <i>RNA Biology</i> , 2010, 7, 213-219.	1.5	146
24	Recent surprises in protein targeting to mitochondria and plastids. <i>Current Opinion in Plant Biology</i> , 2006, 9, 610-615.	3.5	145
25	A reevaluation of dual-targeting of proteins to mitochondria and chloroplasts. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2013, 1833, 253-259.	1.9	141
26	Two Interacting Proteins Are Necessary for the Editing of the NdhD-1 Site in <i>Arabidopsis</i> Plastids. <i>Plant Cell</i> , 2012, 24, 3684-3694.	3.1	130
27	RNAi for revealing and engineering plant gene functions. <i>Current Opinion in Biotechnology</i> , 2007, 18, 148-153.	3.3	128
28	<i>Small kernel1</i> encodes a pentatricopeptide repeat protein required for mitochondrial <i>nad7</i> transcript editing and seed development in maize ( <i>Zea mays</i> ) and rice ( <i>Oryza</i> )	2.8	127
29	Nucleotide and RNA Metabolism Prime Translational Initiation in the Earliest Events of Mitochondrial Biogenesis during Arabidopsis Germination. <i>Plant Physiology</i> , 2012, 158, 1610-1627.	2.3	124
30	Identification of a Pentatricopeptide Repeat Protein Implicated in Splicing of Intron 1 of Mitochondrial <i>nad7</i> Transcripts. <i>Journal of Biological Chemistry</i> , 2010, 285, 32192-32199.	1.6	123
31	<i>MAT4</i> , a maturase factor required for <i>nad1</i> pre-rRNA processing and maturation, is essential for holocomplex biogenesis in Arabidopsis mitochondria. <i>Plant Journal</i> , 2014, 78, 253-268.	2.8	110
32	OTP70 is a pentatricopeptide repeat protein of the E subgroup involved in splicing of the plastid transcript <i>rpoC1</i> . <i>Plant Journal</i> , 2011, 65, 532-542.	2.8	106
33	The cytidine deaminase signature HxxE(x)CxxC of DYW1 binds zinc and is necessary for RNA editing of <i>ndhD1</i> . <i>New Phytologist</i> , 2014, 203, 1090-1095.	3.5	100
34	A PORR domain protein required for <i>rpl2</i> and <i>ccmF</i> intron splicing and for the biogenesis of c-type cytochromes in Arabidopsis mitochondria. <i>Plant Journal</i> , 2012, 69, 996-1005.	2.8	99
35	<i>CSF1</i> , a nucleus-encoded CRM protein required for the processing of many mitochondrial introns, is involved in the biogenesis of respiratory complexes I and IV in Arabidopsis. <i>New Phytologist</i> , 2013, 199, 379-394.	3.5	98
36	An Arabidopsis Dual-Localized Pentatricopeptide Repeat Protein Interacts with Nuclear Proteins Involved in Gene Expression Regulation. <i>Plant Cell</i> , 2011, 23, 730-740.	3.1	96

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37	Mutations in an <i>Arabidopsis</i> Mitochondrial Transcription Termination Factor-Related Protein Enhance Thermotolerance in the Absence of the Major Molecular Chaperone HSP101. <i>Plant Cell</i> , 2012, 24, 3349-3365.	3.1	94
38	Pentatricopeptide Repeat Proteins in <i>Trypanosoma brucei</i> Function in Mitochondrial Ribosomes. <i>Molecular and Cellular Biology</i> , 2007, 27, 6876-6888.	1.1	92
39	The pentatricopeptide repeat protein OTP82 is required for RNA editing of plastid <i>ndhB</i> and <i>ndhG</i> transcripts. <i>Plant Journal</i> , 2010, 61, 339-349.	2.8	92
40	The plastid redox insensitive-2 mutant of <i>Arabidopsis</i> is impaired in PEP activity and high light-dependent plastid redox signalling to the nucleus. <i>Plant Journal</i> , 2012, 70, 279-291.	2.8	81
41	The Pentatricopeptide Repeat Proteins TANG2 and ORGANELLE TRANSCRIPT PROCESSING439 Are Involved in the Splicing of the Multipartite <i>nad5</i> Transcript Encoding a Subunit of Mitochondrial Complex I. <i>Plant Physiology</i> , 2014, 165, 1409-1416.	2.3	78
42	Duplication and Quadruplication of <i>Arabidopsis thaliana</i> Cysteinyl- and Asparaginyl-tRNA Synthetase Genes of Organellar Origin. <i>Journal of Molecular Evolution</i> , 2000, 50, 413-423.	0.8	76
43	Albinism in Plants: A Major Bottleneck in Wide Hybridization, Androgenesis and Doubled Haploid Culture. <i>Critical Reviews in Plant Sciences</i> , 2009, 28, 393-409.	2.7	76
44	The Pentatricopeptide Repeat Protein OTP87 Is Essential for RNA Editing of <i>nad7</i> and <i>atp1</i> Transcripts in <i>Arabidopsis</i> Mitochondria. <i>Journal of Biological Chemistry</i> , 2011, 286, 21361-21371.	1.6	76
45	<i>scp&gt;AEF&lt;/scp&gt;1&lt;/i&gt;/<i>scp&gt;MPR&lt;/scp&gt;25</i> is implicated in <i>scp&gt;RNA&lt;/scp&gt;</i> editing of plastid <i>atpF</i> and mitochondrial <i>nad5</i>, and also promotes <i>atpF</i> splicing in <i>Arabidopsis</i> and rice. <i>Plant Journal</i>, 2015, 81, 661-669.</i>	2.8	75
46	Predictable Alteration of Sequence Recognition by RNA Editing Factors from <i>Arabidopsis</i> . <i>Plant Cell</i> , 2015, 27, 403-416.	3.1	75
47	Potential dual targeting of an <i>Arabidopsis</i> archaeobacterial-like histidyl-tRNA synthetase to mitochondria and chloroplasts. <i>FEBS Letters</i> , 1998, 431, 39-44.	1.3	73
48	<i>Arabidopsis</i> tRNA Adenosine Deaminase Arginine Edits the Wobble Nucleotide of Chloroplast tRNA <sup>Arg</sup> (ACG) and Is Essential for Efficient Chloroplast Translation. <i>Plant Cell</i> , 2009, 21, 2058-2071.	3.1	69
49	The potential for manipulating <i>scp&gt;RNA&lt;/scp&gt;</i> with pentatricopeptide repeat proteins. <i>Plant Journal</i> , 2014, 78, 772-782.	2.8	64
50	Surrogate mutants for studying mitochondrially encoded functions. <i>Biochimie</i> , 2014, 100, 234-242.	1.3	61
51	The mitochondrial pentatricopeptide repeat protein <i>scp&gt;PPR&lt;/scp&gt;19</i> is involved in the stabilization of <i>NADH dehydrogenase 1</i> transcripts and is crucial for mitochondrial function and <i>Arabidopsis thaliana</i> development. <i>New Phytologist</i> , 2017, 215, 202-216.	3.5	60
52	The <i>scp&gt;E&lt;/scp&gt;</i> domains of pentatricopeptide repeat proteins from different organelles are not functionally equivalent for <i>scp&gt;RNA&lt;/scp&gt;</i> editing. <i>Plant Journal</i> , 2013, 74, 935-945.	2.8	58
53	PPR-SMRs. <i>RNA Biology</i> , 2013, 10, 1501-1510.	1.5	57
54	Mitochondrial Defects Confer Tolerance against Cellulose Deficiency. <i>Plant Cell</i> , 2016, 28, 2276-2290.	3.1	57

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55	Towards a plant model for enigmatic U $\alpha$ - $\alpha$ C RNA editing: the organelle genomes, transcriptomes, editomes and candidate RNA editing factors in the hornwort <i>Anthoceros agrestis</i> . <i>New Phytologist</i> , 2020, 225, 1974-1992.	3.5	57
56	<b>Organelle RNA editing</b>. <i>Wiley Interdisciplinary Reviews RNA</i> , 2011, 2, 493-506.	3.2	55
57	The Pentatricopeptide Repeat Protein EMB2654 Is Essential for Trans-Splicing of a Chloroplast Small Ribosomal Subunit Transcript. <i>Plant Physiology</i> , 2017, 173, 1164-1176.	2.3	52
58	Quantitative analysis of motifs contributing to the interaction between <scp>PLS</scp> subfamily members and their target <scp>RNA</scp> sequences in plastid <scp>RNA</scp> editing. <i>Plant Journal</i> , 2014, 80, 870-882.	2.8	51
59	Arabidopsis CSP41 proteins form multimeric complexes that bind and stabilize distinct plastid transcripts. <i>Journal of Experimental Botany</i> , 2012, 63, 1251-1270.	2.4	49
60	Evolutionary plasticity of restorer-of-fertility-like proteins in rice. <i>Scientific Reports</i> , 2016, 6, 35152.	1.6	46
61	A <scp>DYW</scp> protein knockout in <i><scp>P</scp>hyscomitrella</i> affects two closely spaced mitochondrial editing sites and causes a severe developmental phenotype. <i>Plant Journal</i> , 2013, 76, 420-432.	2.8	45
62	European consortia building integrated resources for Arabidopsis functional genomics. <i>Current Opinion in Plant Biology</i> , 2003, 6, 426-429.	3.5	44
63	The coordinated action of <scp>PPR</scp>4 and <scp>EMB</scp>2654 on each intron half mediates <i>trans</i> splicing of <i>rps12</i> transcripts in plant chloroplasts. <i>Plant Journal</i> , 2019, 100, 1193-1207.	2.8	42
64	The E domain of CRR2 participates in sequence-specific recognition of RNA in plastids. <i>New Phytologist</i> , 2019, 222, 218-229.	3.5	36
65	Plastid tRNA Genes trnC-GCA and trnN-GUU are essential for plant cell development. <i>Plant Journal</i> , 2007, 51, 751-762.	2.8	30
66	A synthetic RNA editing factor edits its target site in chloroplasts and bacteria. <i>Communications Biology</i> , 2021, 4, 545.	2.0	28
67	High intraspecific diversity of <i>Restorer</i> fertility-like genes in barley. <i>Plant Journal</i> , 2019, 97, 281-295.	2.8	24
68	In-silico identification of candidate genes for fertility restoration in cytoplasmic male sterile perennial ryegrass ( <i>Lolium perenne</i> L.). <i>Genome Biology and Evolution</i> , 2016, 9, evw047.	1.1	22
69	Editing of Chloroplast rps14 by PPR Editing Factor EMB2261 Is Essential for Arabidopsis Development. <i>Frontiers in Plant Science</i> , 2018, 9, 841.	1.7	18
70	Evolutionary Model of Plastidial RNA Editing in Angiosperms Presumed from Genome-Wide Analysis of <i>Amborella trichopoda</i> . <i>Plant and Cell Physiology</i> , 2019, 60, 2141-2151.	1.5	17
71	GeneFarm, structural and functional annotation of Arabidopsis gene and protein families by a network of experts. <i>Nucleic Acids Research</i> , 2004, 33, D641-D646.	6.5	16
72	Using the SUBcellular database for Arabidopsis proteins to localize the Deg protease family. <i>Frontiers in Plant Science</i> , 2014, 5, 396.	1.7	16

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73	Protein Complexes Implicated in RNA Editing in Plant Organelles. <i>Molecular Plant</i> , 2017, 10, 1255-1257.	3.9	14
74	In <i>Arabidopsis thaliana</i> distinct alleles encoding mitochondrial RNA PROCESSING FACTOR 4 support the generation of additional 5' termini of ccmB transcripts. <i>Plant Molecular Biology</i> , 2017, 93, 659-668.	2.0	13
75	Cofactor-independent RNA editing by a synthetic S-type PPR protein. <i>Synthetic Biology</i> , 2022, 7, ysab034.	1.2	12
76	A PPR protein involved in regulating nuclear genes encoding mitochondrial proteins?. <i>Plant Signaling and Behavior</i> , 2011, 6, 748-750.	1.2	11
77	Mitochondrial genomes as living "fossils". <i>BMC Biology</i> , 2013, 11, 30.	1.7	11
78	The GENOMES UNCOUPLED1 protein has an ancient, highly conserved role but not in retrograde signalling. <i>New Phytologist</i> , 2022, 236, 99-113.	3.5	11
79	Triticeae genome sequences reveal huge expansions of gene families implicated in fertility restoration. <i>Current Opinion in Plant Biology</i> , 2022, 66, 102166.	3.5	8
80	In Silico Methods for Identifying Organellar and Suborganellar Targeting Peptides in <i>Arabidopsis</i> Chloroplast Proteins and for Predicting the Topology of Membrane Proteins. <i>Methods in Molecular Biology</i> , 2011, 774, 243-280.	0.4	6
81	The Pentatricopeptide Repeat Protein MEF100 Is Required for the Editing of Four Mitochondrial Editing Sites in <i>Arabidopsis</i> . <i>Cells</i> , 2021, 10, 468.	1.8	4