

Supplemental Figure 1. RNA from pooled samples can be used to detected editing differences between control and silenced plants. Comparison of the sequencing electrophoretograms of *nad2* RT-PCR products obtained from control and silenced pools shows a noticeable difference in the height of the T peak at position 558. The difference in the T peaks reflects the differential editing extent of *nad2*-558 between control and silenced pools.



Supplemental Figure 2. An example of differential editing between pools of control and silenced plants assessed by electophoretogram traces of bulk RT-PCR products. Editing T is shown in red. A, *OrfX*-552 is more edited in control than in silenced pools. B, *MatR*-1771 is less edited in control than in silenced pools. Note that the height of the peaks does not reflect the actual editing extent as evaluated by PPE (43% vs. 26% and 60% vs. 76% for *orfX*-552 and *matR*-1771 respectively).



Supplemental Figure 3. Virus induced gene silencing (VIGS) of *REME1* results in increased editing of sites found in *rpl5*. A, Editing extent of C at position 92 is increased by an average of 15% in silenced plants ((85-74)/74 = 0.15, P < 5 10⁻⁵). B, The increase in editing is much less pronounced for the edited C at position 329 (P < 3 10⁻⁴). On the right of each group is the average editing extent with s.d.



Supplemental Figure 4. Editing extent of *nad2*-558 is decreased in *REME1* silenced plants. Editing extents were assayed by PPE. A and B show data from two independent experiments. A, Virus inoculation induces a significant decrease of *nad2*-558 editing extent (not inoculated vs. empty vector, P<2 10⁻²), but half the range of *REME1* specific silencing (empty vector vs. REME1-silenced, P<7 10⁻⁴). B, No effect of virus inoculation on *nad2*-558 editing extent in *REME1* silenced plants is more pronounced in B than in A (29% = (50-36)/36 x 100 vs. 18% (49-40)/49 x 100). On the right of each group is the average editing extent with s.d.



Supplemental Figure 5. REME1 is targeted to mitochondria. In the panels are shown the subcellular localization in onion epidermal cells of REME1 fused to GFP (left panel) and of the Cherry fluorescent marker addressed to mitochondria (right panel). In the middle panel is shown an overlay of the two pictures showing a clear co-localization of both fluorescent markers.



Supplemental Figure 6. Semi-quantitative RT-PCR of a reference gene (At4g26410) shows no difference in expression level between transgenic lines and parental accessions. RT-PCR products were amplified and electrophoresed on 1% agarose gels after a completed number of cycles indicated above each lane. On the left of each gel are indicated the origin of the RNA used for the RT-PCR: 1, 13, 5, 16, 2, and 14 are T_0 Ler plants expressing *REME1* Col allele and showing a high, low, and medium level of *nad2*-558 editing extent respectively. The same amount of template RNA (50 ng) was used for each reverse transcription reaction. The abundance of RT-PCR products is very similar for each template. This control experiment validates the difference observed in *REME1* expression level reported in Figure 10.



Supplemental Figure 7. Semi-quantitative RT-PCR is able to detect a three to four-fold reduction in *nad2* second intron spliced or intron adjoining product. Intron-spliced and intron-adjoining RT-PCR products for the second intron were amplified and electrophoresed on 2% agarose gels after a completed number of cycles indicated above each lane. –RT indicates a negative control in which the reverse transcriptase was omitted and was performed for each combination of primers at the maximum number of cycles (e.g. 31 for intron2-spliced). On the left of each gel are indicated the origin of the RNA used for the RT-PCR: 1 is a T₀ Ler plant expressing *REME1* Col allele, 1-d3X and 1-d4X indicate a three and four times dilution of the cDNA from 1. The difference between 1 and 1-d3X is readily observable as there is a shift of about 2 cycles to obtain the same amount of PCR product (e.g., intron-spliced amplification, 27 cycles for 1 vs. 29 cycles for 1-d3X). The difference in PCR products abundance is even more pronounced between 1 and 1-d4X.

Supplemental Table 1. Markers used in *REME1* fine mapping

Marker ^a	Category	Primers ^b	Amplicon length	Enzyme	Allele cut	Polymorphism
						coordinates (AGI map) ^c
1027K	CAPs	1027K-F1 = CCTTGATATAAACTTGACGATTGT	556 bp	RsaI	Ler	1027083
		1027K-R1 = ACGAACCGACAATATTATGTATGT	_			
1113K	CAPs	1113K-F1 = AAAGATCGGTAGACAAATCATTGA	381 bp	MfeI	Ler	1112836
		1113K-F1 = ATTCGCCGAGTTATTTGTAAAAA				
1135K	CAPs	1135K-F1 = ACTCTAGTGAGCCCAAAAGATGAC	406 bp	TscI	Col	1134935
		1135K-R1 = AGACCGGATGTGTTACCATCTG				
1150K	sequencing	1150K-F1 = GCGGCTTGCATACCAGTA	442 bp			11 SNPs
		1150K-R1 = CGTCACTAATCCTAAATCAGGTAC				1149913-1150153
1170K	sequencing	1170K-F1 = CTATTAAGCTATTTCAAGGGAAAAA	416 bp			1170362
		1170K-R1 = TCGATCGGTTATACTATGTTTAATC				
1181K	CAPs	1181K-F1 = AGACGCCTTGGTTTGATTGT	471 bp	PstI	Col	1181241
		1181K-R1 = ATTCGATTTTTCCGGTTTACTTC				
At2g03880	sequencing	At2g03880-F1 = ACCGGAAAAATCGAATGTCTGAGT	2316 bp			51 SNPs, 3 indels
		At2g03880-5' = TGGATTCATTGCAAAGTCATGG				1181511-1183701
		At2g03880-F2 = TGGTGGGTTTGCTCAGAATAGTA				
		At2g03880-F3 = CGGAGTATGCAGCTAAAAAGGTC				
		At2g03880-R1 = GTGCAGCAAGTGTCTTCCCATTAA				
1184K,1186K	sequencing	1184K-F1 = TCATGTCATCGCTACATCCTATTA	1917 bp			4 SNPs, 3 indels
		1184K-F2 = TTCCACTGCAAAGGTATTAAATA				1184613-1185817
		1186K-R1 = CATCCCCAAAACACAACCAGT				
1193K	CAPs	1193K-F1 = AAATCGATCTCTAGCTTCAACTAA	398 bp	EarI	Ler	1192628
		11931K-R1 = CTATCTTAGTGGCGAGCATGATC				
1220K	CAPs	1220K-F1 = AATGTTTCCCAATTAATAGTATAGATG	402 bp	EcoRV	Ler	1220166
		1220K-R1 = AAGCGACAATATAATAAAATTAGTTTT				
1276K	CAPs	1276K-F1 = GCATAAGTTAGGAGCAAGAAAGTT	381 bp	ClaI	Col	1275626
		1276K-R1 = GACGCGGATCAAAGAAAATA				

^aMarkers 1027K and 1276K were used to screen the 692 F2 plants. Recombinants were then genotyped with the other markers.

^bFor sequencing markers the internal markers used for sequencing are also given, the amplification were obtained with the pair marker-F1, marker-R1.

^cPhysical coordinates of the polymorphism are given in bp; when several SNPs correspond to a marker, the number is given and the coordinates of the first and the last SNP are also provided.

Coding sequences $atp1$ $atp1-Fl = CGCGGAATTAGACCTGCTATTAAC469 bpatp1-Rl = TGATTGAGGATTCCTAATGTGATGccb206ccb206ccb206-Fl = CAGCCTTGAAGTGAATGAATGAATGACTA661 bpccb382ccb382-R2 = GATTAGACCATGTTCCTGAAGGACCccb382-R2 = GATTAGACCATGTTCCTGAGAGTTTccb452ccb452-P2 = GAATCACTTCATGCCGACCTCccb452-P2 = GAATCACTTCATGCCGACCTCccb452-R2 = AAGTGTTTGGGCATTAATTGATTGG000000000000000000000000000000000000$	Gene	Primers ^a	Amplicon length
atp1 $atp1-F1 = CGCGGAATTAGACCTGCTATTAAC469 bpatp1-R1 = TGATTGAGGATTCCTAATGTGATGatp1-R1 = TGATTGAGGATTCCTAATGTGATGccb206ccb206-F1 = CAGCCTTGAAGTGAATGAATT661 bpccb206-F1 = CTAGCTTTGTGAAACTAATCGAGACCccb382ccb382ccb382-R2 = GATTAGACCATGTCCTGAGATTTccb452ccb452-F2 = GAATCACTTCATGCCGACCTCccb452-F2 = GATTGGGGGATTAATTGGCCGACCTCccb452-F2 = TCAGATACTCCATAGCCGACCTCCcox2-R1 = ATGATTGTCTCAAAGTGCCGATCCCmatRmatR-F1 = AGAGGCGATCAGAATGGTACTCGAmatR-F1 = AGAGGCCAGCAGAATGGTACTCGAmatR-F1 = AGAGGCCAGACATAGTGAGTAAATAAAA995 bpmatl-R1* = GTCTTGACCGGTCCGAGCTTCCmatRmatR-F1 = AGAGGCGATCAGAATGGTACTCAGTGmatR-F1 = AGGCCGTACCGTATCATGAGTAATAAAA995 bpmatl-R1* = AAGGTGCACTAGAGTAGCAGAAACnad2mad2-F1 = GACCGTACCCTTTCTTTGAAmad2-F2 = TTGATGCTTTTGTGAATTGCTATmad3mad3-F1 = AGCAAGGAGCGCAGAAAACAAAGTmad4-R1* = TTTGCCATGTTGCACTATGTATmad4-R1* = TTTGCCATGTTGCAATGCTATTmad5-F1 = TTTTCCGGACGTTTCCAAGGACAAACAmad5-F2 = GGTGATTTGGGACATAGCTAAGGCmad5-F2 = GGTGATTTGGGACATAGCTATGCmad6-F1 = CTTTCCCCCTCTCCAAACACAAATmad7-R1 = ATGCACGCTCTCCCAACCACAATAmad7-R2 = TCTCCCCGCTCCTCAAAAmad7-R2 = TCTCCCCGCTCCTCAAAAmad7-R2 = TCTCCCCGCCCCTCCAAAAmad7-R2 = TCTCCCCGCCCCTCCAAAAmad7-F2 = AAAGGATTGGAATGGATGmad7-F2 = AAGGCTACGGATTGGATGmad7-F1 = ATGACGACTAGGAAAAGGAAACCCGAGTAGmad7-F2 = TCCCCCGCTCCTCAAAA$	Coding sequ	lences	
atp1 atp1-F1 = CGCGGAATTAGACCTGCTATTAAC 469 bp atp1-R1 = TGATTGAGGATTCCTAATGTGATG atp1-R1 = TGATTGAGGATTCCTAATGTGATG ccb206 ccb206-F1 = CAGCCTTGAAGTGAATGAATT 661 bp ccb382 ccb382-R2 = GATTAGACCATGTTCCTGAGATT ccb382 ccb452-F2 = GAATCACTCATGCCGACCTC ccb452-F2 = GATTGGGCGTTCCTTCT ccb452-F2 = GATTGGGCGATTAATTGATCGTCCTTCT ccv2 cox2-F1 = ATGATTGTTCTAAAATGGTTATT 770 bp cox2-R2 = TCAGAATACTCATAAGTCCGATACC matR-R1 = GTCTTGACCGGGTCCGAGCTTCC 1927 bp matR-R1 = GTCTTGACCGGGTCCGAGCTTCC nad1-R1* = AAGGCCAGATCATGAGTAAATAAAA 995 bp nad1 nad1-F1 = AGGCCAGATCATGAGTAAATAAAAA 995 bp nad2-R1 = AAGGCTACCCTTTCTTTGAA nad2-F1 = GACCGTAACGTAAGTGACAGAAAC nad2-R1 = AGGCCAGACCCGTGTCAAAT nad2-F1 = TGATGCTTTTGAATTCATTGTAT nad2-R1 = ACGGCTTACCCTTTCTTGGAA nad2-F1 = AGGAAGCCGTACCCTTTCTTGAA nad3 nad3-F1 = AGGCAAGCGGAGAAACAAAGT 414 bp nad3-R1 = CCCCCATTTGGTGCACTAAGTACC nad4-R1* = TTTGCCAGTGTTCGAAAAGCAAAGT 147 bp nad4 nad4-F1 = ATGTAGACTATCGAAAAGAAAGAAAAGGCAAAT 147 bp nad5-F1 = TTAAAACTACTCACTATCGAAAGTAAAGAAAGAAAAGAA			
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ccb206ccb206-Fl = CAGCCTTGAAAGTGAATGAATT661 bpccb206-R1* = TTAATCTTGTAAACTAATCGAGACCccb382-R2 = GATTAGACCATGTTCCTGAGATTTccb452ccb452-F2 = GAATCACTTCATGCCGACCTCccb452-R2 = AAGTGTTGGGCATTACTGGCGACCTCccv2cox2-F1 = ATGATTGTTCTAAAATGGTTATT770 bpcox2-R2 = TCAGAATACTCATAAGGTCGATACCmatRmatR-F1 = AGAGGCGATCAGAATGGTACTCGAmatRmatR-F1 = AGAGGCGATCAGAATGGTACTCGAmatRmatR-F1 = AGGCCAGATCAGAATGGTACTCGAmatRmatR-F1 = AGGCCAGATCATGAGTAAAAAAAAAAAAAAAAAAAAAAA		atp1-R1 = TGATTGAGGATTCCTAATGTGATG	
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cox2cox2-F1 = ATGATTGTTCTAAAATGGTTATT770 bpcox2-R1* = GTTTGGGGGATTAATTGATTGGcox2-R2 = TCAGAATACTCATAAGTCGATACCmatRmatR-F1 = AGAGGCGATCAGAATGGTACTCGA1927 bpmatR-R1 = GTCTTGACCGGGTCCGAGCTTCCnad1-F1 = AGGCCAGATCATGAGTAAATAAAA995 bpnad1nad1-F1 = AGGCCAGATCATGAGTGACTCAGTG1578 bpnad2nad2-F1 = GACCGTAACGTAACGTAAGTGACTCAGTG1578 bpnad2-R1 = ACGGCCTACCCTTTCTTGAAnad2-R1 = ACGGCCTACCCTTTCTTGAAnad2-R1 = ACGGCCTACCCTTTCTTGAAnad2-F2 = TTGATGCTTTGAATTCATTGTATnad3nad3-F1 = AGCAAGGAGCGAGAAACAAAGT414 bpnad3-R1 = CCCCCATTTTGTGCCCTATC1479 bpnad4nad4-F1 = ATGTTAGACATTCTGTGAATGCTATT1479 bpnad5-R1 = TTGCCATGTTGCACTAAGTTACT1479 bpnad5-R1 = TTTTCCGAGCTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCCACTATCAAAATGAAAG1937 bpnad5-R1 = TAGATACTTCTGTTTGTCGAG536 bpnad6 = nad5-F3 = GAGATCTCATTCGAAGCTTTAGC1179 bpnad7 = ATGATACTTCCTCCTAAGAAGCCAAAT1179 bpnad7 = ATGATACTTCCTCCCAACACAATA1179 bpnad7-R1 = ATCCACCTCCCAAACACAATA1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA381 bporf114 orf114-F1 = CAACGGCGATTGGATG381 bporf114 = ATGGAGAAAGGAAAAGGAAACACGAATT381 bporf114 = ATGGAGAGGAGACACCGAGTAG381 bporf114 = ATGGAGAAAGGAACACCGAATGA381 bporf114-R1 = ATGGAGAGGAGGAATCCGTATA381 bp		ccb452-R2 = AAGTGTTTGGCCTTTCCTTCT	
$\begin{tabular}{lllllllllllllllllllllllllllllllllll$	cox2	cox2-F1 = ATGATTGTTCTAAAATGGTTATT	770 bp
cox2-R2 = TCAGAATACTCATAAGTCCGATACCmatRmatR-F1 = AGAGGCGATCAGAATAGTAACTCGA1927 bpmatRmatR-R1 = GTCTTGACCGGGGTCCGAGCTTCC1927 bpnad1nad1-F1 = AGGCCAGATCATGAGTAAATAAAA995 bpnad1-R1* = AAAGGTGACTAAAAGACCAGAAAC1000000000000000000000000000000000000		cox2-R1* = GTTTGGGGGGATTAATTGATTGG	
matRmatR-F1 = AGAGGCGATCAGAATGGTACTCGA1927 bpmatR-R1 = GTCTTGACCGGGTCCGAGCTTCCmatR-R1 = GTCTTGACCGGGTCCGAGCTTCC995 bpnad1nad1-F1 = AGGCCAGATCATGAGTAAATAAAA995 bpnad2nad2-F1 = GACCGTAACGTAACAGTGACTCAGTG1578 bpnad2-F1 = ACGGCCTACCCTTTCTTTGAAnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F3 = GGCAAACCCGTGTCAAAT414 bpnad3-R1 = CCCCCATTTTGGACTACC1479 bpnad4-R1* = TTGCCATGTTGGAATGCTATT1479 bpnad4-R1* = TTGCCCATGTTGGAATGCTATT1479 bpnad5-F1 = AGGGCAAACACCCTATC1937 bpnad5-F1 = TTTTCCGGCACTAACATTCTGGGA1937 bpnad5-F1 = TTTTCGGACGTTTTCGAGAC1937 bpnad5-F1 = TTTTCCGGCCTATCGGAAAACAA1937 bpnad5-F1 = TTTTCCGGCCTATCGGAAAACGA1937 bpnad5-F2 = GGTGATTTGGATTAGCTCTTG1179 bpnad6nad6-F1 = ATGATACTTTCTGTTTTGCGAG536 bpnad7-R1* = ATCACCCTCTCCAAAAAGGCAAAT1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-F1* = CAACCGGCGATTGGATG381 bporf114-F1* = CAACCGGCGATTGGATG381 bporf114-F1* = CGCCCCTCCTTGAATTTT179 bpnad7nafGGGGGGAGGAAAAGGAAACCGAGTAGnad7naf7-F2 = AAAGGATTGGGGATTCAGTGorf114-F1* = CAACCGGCGATTGGATG381 bporf114-F1* = CAACCGGCGATTGGATGnad7naf3-F2 = CGCCCCTCCTTGAATTCA		cox2-R2 = TCAGAATACTCATAAGTCCGATACC	
matR-R1 = GTCTTGACCGGGTCCGAGCTTCCnadlnadl-F1 = AGGCCAGATCATGAGTAAATAAAA995 bpnadl-R1* = AAAGGTGACTAAAAGACCAGAAAC995 bpnad2-R1 = GACCGTAACGTAAGTGACTCAGTG1578 bpnad2-R1 = ACGGCCTACCCTTTCTTGAA1578 bpnad2-F3 = GGCAAAGCCGTGTCAAAT414 bpnad3-R1 = CCCCCATTTTGTGCCCTATC414 bpnad3-R1 = CCCCCATTTTGTGCCCTATC1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACT1479 bpnad5-R1 = TTTTCCGGACATTTCTAGGA1937 bpnad5-R1 = TTTTCCGGACGTTTCTAGGA1937 bpnad5-R1 = TAAAACTACTCACTATCAAATGAAAG1937 bpnad5-R1 = TTTTCCGACGATTTCGAGAGCTTTAGAC1179 bpnad6nad6-F1 = ATGATACTTTCTGTTAGTAGCnad7nad7-F1 = ATCACCATCTAGCAAAAGCAAAT1179 bpnad7-R2 = TTCCCCGCTCTCCAAACACAATA1179 bpnad7-R1 = ATCCACCTCTCCAAACACAATA1179 bpnad7-R2 = TTCCCCGCTCCTCAAACACAATA381 bporf114-R1* = CAACCGGCATTGGAATGCTATG381 bporf114-R1* = AATGGAAAAGGAAACCGAGTAG381 bporf114-R1 = AATGGAAAAGGAAATCCGTATG381 bporf114-R1 = AATGGAAAAGGAAATCCGAATA381 bporf114-R1 = AATGGAAAAGGAAACCCGAGTAGG381 bporf114-R1 = AATGGAAAAGGAAACCCGAGTAGG381 bporf114-R1 = AATGGAAAAGGAAACCGAATTA381 bporf114-R1 = AATGGAAAAGGAAACCCGAGTAG381 bporf114-R1 = AATGGAAAAGGAAACCGAATTA381 bporf114-R1 = AATGGAAAAGGAAACCCGAGTAGG381 bporf114-R1 = AATGGAAAGGAAACCGAATTATA381 bporf114-R1 = AATGGAAAAGGAAATCCGTATA381 bporf114-R1 = AATG	matR	matR-F1 = AGAGGCGATCAGAATGGTACTCGA	1927 bp
nad1nad1-F1 = AGGCCAGATCATGAGTAAATAAAA995 bpnad1-R1* = AAAGGTGACTAAAAGACCAGAAACnad2-R1 = AAGGCCACCTACCTAAAGTGACTCAGTG1578 bpnad2-R1 = ACGGCCTACCCTTTCTTTGAAnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F2 = TTGATGCTTTTGACCCTATCnad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTGGCCCTATCnad4-F1 = ATGTTAGAACATTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-F1 = ATGTTAGAACATTCTGGGAAAACAA1479 bpnad5-F1 = CCCCCATTTCGGGAAAAACAnad5-F1 = TTTTTCCGAGGATTAGCTCTTG1937 bpnad5-F1 = TTTTTCCGACGTTTTCTAGGA1937 bp1937 bpnad5-F1 = GATGATTTTGGATTAGCTCTTGnad5-F3 = GAAGTCTCATTCGAAGCTTTAGAAC1179 bpnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad7nad7-F1 = ATGCACCTCTCCAAACACAATA1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1* = AATGGAAAAGGAACACCGAGTAG783-F3 = GGGTGGAGGAATCCGTATA		matR-R1 = GTCTTGACCGGGTCCGAGCTTCC	
nad1-R1* = AAAGGTGACTAAAAGACCAGAAACnad2nad2-F1 = GACCGTAACGTAAGTGACTCAGTG1578 bpnad2-R1 = ACGGCCTACCCTTCTTGAAnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F3 = GGCAAACCCGTGTCAAATnad3nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTGTGCCCTATCnad4-F1 = ATGTTAGAACATTCCTGTGAATGCTATTnad4nad4-F1 = ATGTTAGAACATTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad5nad5-F1 = TTTTTCGGACGTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F1 = TTTTCCGGACGTTTGCGAGnad6nad6-F1 = ATGATACTTCTGTTTTGTCGAG536 bpnad6nad6-F1 = ATGATACTTCTGTAGTAGTCCTATGC1179 bpnad7-R1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-F1 = ATGACCACTCTCCAAACACAAATAnad7-F2 = AAAGGATTGGGGATTCAGTGorf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114orf114-F1* = CGCCCCTCCTTGAATTTT181 bporf114sp3-F3 = GGGTGGAGGGAATCCGTATGC381 bp	nad1	nad1-F1 = AGGCCAGATCATGAGTAAATAAAA	995 bp
nad2nad2-F1 = GACCGTAACGTAACGTGACTCAGTG1578 bpnad2-R1 = ACGGCCTACCCTTTCTTTGAAnad2-R1 = ACGGCCTACCCTTTCTTTGAAnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F3 = GGCAAACCCGTGTCAAATnad3nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTGTGCCCTATC1479 bpnad4nad4-F1 = ATGTTAGAACATTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-R2 = CACGCTTTCGGGAAAAACAnad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTGnad6-F1 = ATGATACTTCTGTTTGTGCAG536 bpnad6-R1 = CTTTCACCTTAGTAGTCTATGC1179 bpnad7-R2 = TTCCCCGCTCCTCAAAA1179 bpnad7-R2 = TTCCCCGCTCCTCAAACACAATA1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114orf114-R1 = AATGGAAAAGGAACACCGAGTAG381 bporf114orf114-R1 = AATGGAAAAGGAACACCGAGTAG381 bp		nad1-R1* = AAAGGTGACTAAAAGACCAGAAAC	
nad2-R1 = ACGGCCTACCCTTTCTTTGAAnad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F3 = GGCAAACCCGTGTCAAATnad3nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTTGTGCCCTATCnad4nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACT1479 bpnad4-R2 = CACGCTTTCGGGAAAAACA1937 bpnad5-R1 = TATAAAACTACTCACTATCAAAATGAAAG1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAG1937 bpnad6-R1 = CTTTTCACCTATCGAAGCCTTTGG536 bpnad7-R1* = ATGCACCTCTCCAAACACAATA1179 bpnad7-R1* = ATCCACCTCTCCAAAAA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114.R1 = AATGGAAAAGGAAAAGGAAATCCGTATG381 bporf114.R1 = AATGGAAAAAGGAAAACCGAGTAG381 bp	nad2	nad2-F1 = GACCGTAACGTAAGTGACTCAGTG	1578 bp
nad2-F2 = TTGATGCTTTTGAATTCATTGTATnad2-F3 = GGCAAACCCGTGTCAAATnad3nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTGTGGCCCTATCnad4nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad5nad5-F1 = TTTTTCGGACGTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad6nad6-F1 = ATGATACTTCTGGTTTGCAGG536 bpnad7-R1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATAnad7-R2 = TTCTCCCGGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGATTCAGTGorf114orf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3rps3-F3 = GGGTGGAGGGAATCCGTATAG		nad2-R1 = ACGGCCTACCCTTTCTTTGAA	
nad2-F3 = GGCAAACCCGTGTCAAATnad3nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTTGTGCCCTATC1479 bpnad4nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-R1* = TTTGCGACGTTTCTGGGAAAAACAnad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTGnad6nad5-F3 = GAAGTCTCATTCGAAGCTTTAGAC536 bpnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad7-R1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGATTCAGTG381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAAACACCGAGTAG783 = GGGTGGAGGGAATCCGTATA		nad2-F2 = TTGATGCTTTTGAATTCATTGTAT	
nad3nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT414 bpnad3-R1 = CCCCCATTTTGTGGCCCTATCnad3-R1 = CCCCCATTTTGTGGCCCTATC1479 bpnad4nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-R2 = CACGCTTTCGGGAAAAACA1937 bpnad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5.R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTG1937 bpnad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad5-F3 = GAAGTCTCATTCGAAGCTTTAGAC536 bpnad6nad6-F1 = ATGATACTTTCTGTTTGTCGAG536 bpnad7-F1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGATTCAGTGorf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG781 bprps3rps3-F3 = GGGTGGAGGGAATCCGTATAG		nad2-F3 = GGCAAACCCGTGTCAAAT	
nad3-R1 = CCCCCATTTTGTGCCCTATCnad4nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-R2 = CACGCTTTCGGGAAAAACA1937 bpnad5-R1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTGnad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad6-F1 = ATGATACTTTCTGTTTGTCGAGnad6nad6-F1 = ATGATACTTTCTGTTTGTCGAG536 bpnad7-R1 = CTTTTCACCTTAGTAGTCCTATGC1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGATTCAGTG381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG381 bprps3rps3-F2 = CGCCCCTCCTTGAATTTTrps3-F3 = GGGTGGAGGGAATCCGTATA	nad3	nad3-F1 = AGCAAGGAGCGAGAAAACAAAGT	414 bp
nad4nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT1479 bpnad4-R1* = TTTGCCATGTTGCACTAAGTTACTnad4-R2 = CACGCTTTCGGGAAAAACAnad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTGnad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad6-F1 = ATGATACTTTCTGTTTTGTCGAGnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad7-R1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG381 bprps3rps3-F2 = CGCCCCTCCTTGAATTTT		nad3-R1 = CCCCCATTTTGTGCCCTATC	
nad4-R1* = TTTGCCATGTTGCACTAAGTTACT nad4-R2 = CACGCTTTCGGGAAAAACAnad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAG nad5-R1 = TAAAAACTACTCACTATCAAAATGAAAG1937 bpnad5-F3 = GGTGATTTTGGATTAGCTCTTG nad5-F3 = GAAGTCTCATTCGAAGCTTTAGAC536 bpnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad7nad7-F1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACACAATA1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG79s353-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAGGGAATCCGTATA	nad4	nad4-F1 = ATGTTAGAACATTTCTGTGAATGCTATT	1479 bp
nad4-R2 = CACGCTTTCGGGAAAAACAnad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTGnad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad6-R1 = CTTTTCACCTTAGTAGTCCTATGCnad6-R1 = CTTTTCACCTTAGGAAAAGGCAAATnad7nad7-F1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATAnad7-R2 = TTCTCCCGGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGATTCAGTG381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG1793rps3 = GGGTGGAGGGAATCCGTATA1179		nad4-R1* = TTTGCCATGTTGCACTAAGTTACT	
nad5nad5-F1 = TTTTTCGGACGTTTTCTAGGA1937 bpnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-R1 = TAAAAACTACTCACTATCAAAATGAAAGnad5-F2 = GGTGATTTTGGATTAGCTCTTGnad5-F3 = GAAGTCTCATTCGAAGCTTTAGACCnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad6-R1 = CTTTTCACCTTAGTAGTCCTATGCnad7-F1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAAATAnad7-R2 = TTCTCCCGGCTCCTCAAAA381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3-F2 = CGCCCCTCCTTGAATTTTrps3-F3 = GGGTGGAGGGAATCCGTATA		nad4-R2 = CACGCTTTCGGGAAAAACA	
nad5-R1 = TAAAAACTACTCACTATCAAAATGAAAG nad5-F2 = GGTGATTTTGGATTAGCTCTTG nad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG nad6-R1 = CTTTTCACCTTAGTAGTCCTATGC536 bpnad7nad7-F1 = ATGACGACTAGGAAAAGGCAAAT nad7-R1* = ATCCACCTCTCCAAACACAATA nad7-R2 = TTCTCCCGCTCCTCAAAA nad7-F2 = AAAGGATTGGGGATTCAGTG orf114-F1* = CAACCGGCGATTGGATG orf114-R1 = AATGGAAAAGGAACACCGAGTAG381 bprps3rps3-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAAGGGAATCCGTATA370	nad5	nad5-F1 = TTTTTCGGACGTTTTCTAGGA	1937 bp
nad5-F2 = GGTGATTTTGGATTAGCTCTTG nad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG nad6-R1 = CTTTTCACCTTAGTAGTCCTATGCnad7nad7-F1 = ATGACGACTAGGAAAAGGCAAAT nad7-R1* = ATCCACCTCTCCAAACACAATA nad7-R2 = TTCTCCCGGCTCCTCAAAA nad7-F2 = AAAGGATTGGGGATTCAGTGorf114orf114-F1* = CAACCGGCGATTGGATG orf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3rps3-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAGGGAATCCGTATA		nad5-R1 = TAAAAACTACTCACTATCAAAATGAAAG	
nad5-F3 = GAAGTCTCATTCGAAGCTTTAGACnad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad6-R1 = CTTTTCACCTTAGTAGTCCTATGC536 bpnad7-R1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATA1179 bpnad7-R2 = TTCTCCCGCTCCTCAAAA1179 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG381 bprps3rps3-F2 = CGCCCCTCCTTGAATTTTrps3-F3 = GGGTGGAGGGAATCCGTATA		nad5-F2 = GGTGATTTTGGATTAGCTCTTG	
nad6nad6-F1 = ATGATACTTTCTGTTTTGTCGAG536 bpnad6-R1 = CTTTTCACCTTAGTAGTCCTATGC1179 bpnad7-R1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATA1179 bpnad7-R2 = TTCTCCCGGCTCCTCAAAA1179 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG1114-F1*rps3rps3-F2 = CGCCCCTCCTTGAATTTT1179 bprps3-F3 = GGGTGGAAGGGAATCCGTATA1179 bp		nad5-F3 = GAAGTCTCATTCGAAGCTTTAGAC	
nad6-R1 = CTTTTCACCTTAGTAGTCCTATGCnad7nad7-F1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATAnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGGATTCAGTG381 bporf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3rps3-F2 = CGCCCCTCCTTGAATTTTrps3-F3 = GGGTGGAAGGGAATCCGTATArps3-F3 = GGGTGGAAGGGAATCCGTATA	nad6	nad6-F1 = ATGATACTTTCTGTTTTGTCGAG	536 bp
nad7nad7-F1 = ATGACGACTAGGAAAAGGCAAAT1179 bpnad7-R1* = ATCCACCTCTCCAAACACAATAnad7-R2 = TTCTCCCGGCTCCTCAAAAnad7-R2 = TTCTCCCGGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGGATTCAGTGorf114orf114-F1* = CAACCGGCGATTGGATGorf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3rps3-F2 = CGCCCCTCCTTGAATTTrps3-F3 = GGGTGGAAGGGAATCCGTATA		nad6-R1 = CTTTTCACCTTAGTAGTCCTATGC	
nad7-R1* = ATCCACCTCTCCAAACACAATAnad7-R2 = TTCTCCCGCTCCTCAAAAnad7-F2 = AAAGGATTGGGGGATTCAGTGorf114orf114-F1* = CAACCGGCGATTGGATGorf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3rps3-F2 = CGCCCCTCCTTGAATTTTrps3-F3 = GGGTGGAGGGAATCCGTATA	nad7	nad7-F1 = ATGACGACTAGGAAAAGGCAAAT	1179 bp
nad7-R2 = TTCTCCCGCTCCTCAAAA nad7-F2 = AAAGGATTGGGGGATTCAGTGorf114orf114-F1* = CAACCGGCGATTGGATGorf114-R1 = AATGGAAAAGGAACACCGAGTAGrps3rps3-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAGGGAATCCGTATA		nad7-R1* = ATCCACCTCTCCAAACACAATA	-
nad7-F2 = AAAGGATTGGGGGATTCAGTG orf114 orf114-F1* = CAACCGGCGATTGGATG 381 bp orf114-R1 = AATGGAAAAGGAACACCGAGTAG 7ps3 rps3-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAAGGGAATCCGTATA rps3-F3 = GGGTGGAGGGAATCCGTATA rps3-F3 = CGCCCCTCCTTGAATCCGTATA		nad7-R2 = TTCTCCCGCTCCTCAAAA	
orf114orf114-F1* = CAACCGGCGATTGGATG381 bporf114-R1 = AATGGAAAAGGAACACCGAGTAG7ps3rps3-F2 = CGCCCCTCCTTGAATTTTrps3-F3 = GGGTGGAGGGAATCCGTATA		nad7-F2 = AAAGGATTGGGGGATTCAGTG	
<pre>orf114-R1 = AATGGAAAAGGAACACCGAGTAG rps3 rps3-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAGGGAATCCGTATA</pre>	orf114	orf114-F1* = CAACCGGCGATTGGATG	381 bp
<i>rps3</i> rps3-F2 = CGCCCCTCCTTGAATTTT rps3-F3 = GGGTGGAGGGAATCCGTATA	-	orf114-R1 = AATGGAAAAGGAACACCGAGTAG	-
rps3-F3 = GGGTGGAGGGAATCCGTATA	rps3	rps3-F2 = CGCCCCTCCTTGAATTTT	
	-	rps3-F3 = GGGTGGAGGGAATCCGTATA	

Supplemental Table 2. New primers used in mitochondrial genes screening

rps4

rps3-R2 = AGAATAATACACCTACCGAGACGA rps4-F2 = AAGTTTGGATCCGAAAAAGTATG rps4-R2 = CACAAACCCTTCGATGACTTAT

UTR and introns

cox3-trailer	cox3-tr-F1 = TTCGGCAATATCTTGGTCATCTG	341 bp
	cox3-tr-R1 = TTCGGGTCATTTCTTGGTGAAC	
nad5 -intron	nad5-intr-F1 = GTGGGGGCAGAGGGGCTCGTAGTACC	406 bp
	nad5-intr-R1 = CGGTCGGGCTATCGAACACAGAGT	
nad6-leader	nad6-ld-F1 = AAGGGCTTGGAAGAAGAAAATG	375 bp
	nad6-ld-R1 = ATAGCGCAATACTTCTTCGTGAAT	
nad7-leader	nad7-ld-F1: GGAGATGCATTTCTGGTACAAGTG	203 bp
	nad7-ld-R1: GTTCCGCACGTTCCACCAC	
nad7-2nd-intron	nad7-2ndintr-F1 = CTCCGCCCGGTGACTAAGAAAG	343 bp
	nad7-2ndintr-R1 = AGCGTGTTCTTGGGCCATCATAG	
rpl5 -trailer	rpl5-tr-F1 = TATTCGGGGGGTTCAATGTGACTATTA	338 bp
	rpl5-tr-R1 = TTCGAAAGAACTCAGATACAGAACGA	
rpl16 -trailer	rpl16-tr-F1= ATTAGCGGCTCATAAACCATGTTC	469 bp
	rpl16-tr-R1= CCATACATATCGAGGGCTTTATCA	
rps7-leader	rps7-ld-F1 = ACAAACTCGACTAAAAGAAGAGGT	410 bp
	rps7-ld-R1 = CGTCAACCATAAGTTTGATTACAT	

^aPrimers in bold are internal primers used for bulk-sequencing the RT-PCR products. Primers with * have already been reported (Bentolila et al., 2008)

Gene	Position of differentially edited site ^a	PPE primers ^b
Coding seque	nces	
ccb206	80	ccb206-80 = CACCAATCACGAGTTTTTCTTTATTCCTC
ccb256	184	ccb256-184R = GCCGTGGCGATATAAACAATAATACTCATC
	421	ccb256-421 = CATCTTGTTTTTATTTACCTGGGTGCAC
ccb382	709	ccb382-709 = GCTCTTGGCATTGCTTTGTTTTCTC
ccb452	1246	ccb452-1246 = ATAATGAAACTGCCTTTTATTTTTTTAT
	1280	ccb452-1280 = ATTTATGTTGGCTTCGTTGGGAGGC
matR	374	matR-374R = GATCGAGCTTGTGTAGGTAGATGTTGCC
	461	matR-461 = CGAAATTCCGATTGTTCAGAGAGTCAG
	1771	matR-1771R = CGTTCGGACTTGATAAAGGTTGTCG
	1807	matR-1807R = GAATATTGCAGACCAGCGAATCTGG
orf114	327	orf114-327 = GGATCAAAGAAACTAGCAGACTAATCACTAAATAG
orfX	552	orfX-552 = CGTATTTTGTTCATTCATCGGTATGCTC
rpl5	92	rpl5-92R = GGGTGCCTTTGGTACTACTCTTATTTCAC
	329	rpl5-329R = CGACCGGAAAATCTAATAGAGACATTACTG
rps14	194	rps14-194 = CCTTTGCACGAGTCAGAAACCGATG
rps4	967	rps4-967 = GGATCGAACTACCTACTCATTATTTGGAGG
UTR and intr	rons	· · · ·
nad5-intron	803	nad5-int-803R = CTCGTAATTCACTTTTGACTCTGTGTTCG
nad7-leader	-39	nad7-ld-39 = TTTCTGGTACAAGTGGTATTGGACAAGATC
rol5-trailer	195	rol5-tr-195 = CCTTTGCACGAGTCAGAAACCGATG

Supplemental Table 3. New PPE primers used in this report

^aSite differentially edited in silenced vs. control plants was detected by bulk sequencing. Position of the edited C in the coding sequence is given relative to the start codon. In leader and trailer, the position of the edited C is relative to the start codon and stop codon respectively.

^bddCTP is used with gene-site, ddGTP is used with gene-siteR.

Product	Primers	Amplicon length
nad2-intron2 spliced	nad2-ex2-F1 = AATATTTGATCTTAGGTGCATTTTC nad2-ex3-R1 = AAAGGAACTGCAGTGATCTTGA	205 bp
nad2-intron2 adjoining	nad2-ex2-F1 = AATATTTGATCTTAGGTGCATTTTC nad2-int2-R1 = CCCGATCCGATAGTTTACAA	145 bp
nad2-intron3 spliced	nad2-ex3-F1 = CTATGGGTCTACTGGAGCTACCC nad2-ex4-R1 = GCGCAATAGAAAGGAATGCT	215 bp
nad2-intron3 adjoining	nad2-int3-F1 = GGCGAATTTCAAACTTGTGG nad2-ex4-R1 = GCGCAATAGAAAGGAATGCT	156 bp

Supplemental Table 4. Primers used for semi quantitative RT-PCR