Table S2. Functions of transcripts edited in the Arabidopsis plastid. PE – partially edited (<50%) in the wild type.

Gene function	Biological process	Gene	Edited	Affected
		abbreviation	nucleotide	in
β subunit of acetyl-CoA carboxylase	Fatty acid synthesis	accD	C794	orrm6
		accD	C1568	orrm1
F subunit of ATP synthase	Photophosphorylation	atpF	C92	
Caseinolytic protease P1	Protein quality control	clpP	C559	orrm1
Maturase K	RNA splicing	matK	C706	orrm1
B subunit of NADH dehydrogenase	Cyclic electron flow	ndhB	C149	
		ndhB	C467	orrm1
		ndhB	C586	orrm1
		ndhB	C746	orrm1
		ndhB	C830	orrm1
		ndhB	C836	orrm1
		ndhB	C872	orrm1
		ndhB	C1255	orrm1
		ndhB	C1481	
D subunit of NADH dehydrogenase	Cyclic electron flow	ndhD	$C2^{PE}$	orrm1
		ndhD	C383	
		ndhD	C674	orrm1
		ndhD	C878	orrm1
		ndhD	C887	orrm1
F subunit of NADH dehydrogenase	Cyclic electron flow	ndhF	C290 ^{PE}	
G subunit of NADH dehydrogenase	Cyclic electron flow	ndhG	C50	orrm1
L subunit of cytochrome $b_6 f$	Photosynthetic electron transport	petL	C5 ^{PE}	
α subunit of cytochrome b_{559} in PSII		psbE	C214	
$β$ subunit of cytochrome b_{559} in PSII	Photosynthetic electron transport	psbF	C77	orrm6
Z subunit of PSII	Photosynthetic electron transport	psbZ	C50	
Ribosomal protein L23	Translation	rpl23	C89	
α subunit of RNA polymerase	Transcription	rpoA	C200	orrm1
β subunit of RNA polymerase	Transcription	rpoB	C338	orrm1
		rpoB	C551 ^{PE}	orrm1
		rpoB	C2432	orrm1
β' subunit of RNA polymerase	Transcription	rpoC1	C488 ^{PE}	
Ribosomal protein S12	Translation	rps12-intron	C(i1 58)PE	orrm1
Ribosomal protein S14	Translation	rps14	C80	
•		rps14	C149	orrm1