

Table S4. RNA editing sites in the *C. monnieri* chloroplast genome.

Gene	Nt Pos	AA Pos	Align Col	Effect	Score
accD (C = 0.8)	74	25	26	ACG (T) => ATG (M)	1
	154	52	53	CGG (R) => TGG (W)	1
	785	262	288	TCG (S) => TTG (L)	0.8
	1186	396	434	CCA (P) => TCA (S)	1
	1394	465	503	CCT (P) => CTT (L)	1
atpA (C = 0.8)	791	264	264	CCC (P) => CTC (L)	1
atpB (C = 0.8)	35	12	12	GCT (A) => GTT (V)	0.86
	1487	496	496	TCG (S) => TTG (L)	1
ccsA (C = 0.8)	556	186	191	CTT (L) => TTT (F)	0.8
matK (C = 0.8)	640	214	229	CAT (H) => TAT (Y)	1
	991	331	346	CTT (L) => TTT (F)	0.86
	1013	338	353	GCG (A) => GTG (V)	0.86
ndhA (C = 0.8)	107	36	36	CCT (P) => CTT (L)	1
	320	107	107	ACA (T) => ATA (I)	1
	566	189	189	TCA (S) => TTA (L)	1
	1073	358	358	TCC (S) => TTC (F)	1
ndhB (C = 0.8)	149	50	50	TCA (S) => TTA (L)	1
	467	156	156	CCA (P) => CTA (L)	1
	586	196	196	CAT (H) => TAT (Y)	1
	611	204	204	TCA (S) => TTA (L)	0.8
	737	246	246	CCA (P) => CTA (L)	1
	746	249	249	TCT (S) => TTT (F)	1
	830	277	277	TCA (S) => TTA (L)	1
	836	279	279	TCA (S) => TTA (L)	1
	902	301	301	ACC (T) => ATC (I)	1
	1481	494	494	CCA (P) => CTA (L)	1
ndhD (C = 0.8)	29	10	10	ACG (T) => ATG (M)	1
	701	234	234	TCG (S) => TTG (L)	1
	905	302	302	TCA (S) => TTA (L)	1
	1325	442	442	TCA (S) => TTA (L)	0.8
ndhF (C = 0.8)	290	97	97	TCA (S) => TTA (L)	1
	2209	737	743	CTT (L) => TTT (F)	1
ndhG (C = 0.8)	166	56	56	CAT (H) => TAT (Y)	0.8
	314	105	105	ACA (T) => ATA (I)	0.8
petB (C = 0.8)	418	140	140	CGG (R) => TGG (W)	1
	611	204	204	CCA (P) => CTA (L)	1
petG (C = 0.8)	104	35	35	TCG (S) => TTG (L)	1
psaI (C = 0.8)	104	35	51	ACT (T) => ATT (I)	1
psbF (C = 0.8)	77	26	26	TCT (S) => TTT (F)	1
psbE (C = 0.8)	214	72	72	CCT (P) => TCT (S)	1
rpl20 (C = 0.8)	308	103	103	TCA (S) => TTA (L)	0.86
rpoC2 (C = 0.8)	842	281	292	ACT (T) => ATT (I)	0.86

	1529	510	529	ACG (T) => ATG (M)	0.86
	1933	645	829	CCA (P) => TCA (S)	1
	2314	772	960	CGG (R) => TGG (W)	1
	2993	998	1198	ACT (T) => ATT (I)	1
	3560	1187	1397	ACT (T) => ATT (I)	0.86
rpoC1 (C = 0.8)	41	14	14	TCA (S) => TTA (L)	1
	1447	483	500	CGT (R) => TGT (C)	1
rpoB (C = 0.8)	338	113	113	TCT (S) => TTT (F)	1
	551	184	185	TCA (S) => TTA (L)	1
	566	189	190	TCG (S) => TTG (L)	1
	1450	484	501	CCT (P) => TCT (S)	1
	2000	667	684	TCT (S) => TTT (F)	1
	2426	809	827	TCA (S) => TTA (L)	0.86
rpoA (C = 0.8)	833	278	280	TCA (S) => TTA (L)	1
rps2 (C = 0.8)	676	226	226	CGT (R) => TGT (C)	0.86
rps14 (C = 0.8)	80	27	27	TCA (S) => TTA (L)	1
	149	50	53	CCG (P) => CTG (L)	1