

Table S3. The list of the full names of the analyzed genes and appropriate abbreviations.

Organelles	Category	Genes
Mitochondrion	<b>Respiration chain complexes</b>	
	Complex I (Nicotinamide adenine dinucleotide dehydrogenase, NADH)	<i>nad1, nad2, nad3, nad4, nad4L, nad5, nad6, nad7, nad9</i>
	Complex II (Succinate dehydrogenase, SDH)	<i>sdh3, sdh4</i>
	Complex III (Cytochrome b)	<i>cob</i>
	Complex IV (Cytochrome c oxidase)	<i>cox1, cox2, cox3</i>
	Complex V (ATP synthase)	<i>atp1, atp4, atp6, atp8,</i>
	Cytochrome c biogenesis	<i>ccmB, ccmC, ccmFC, ccmFN</i>
	<b>Translation</b>	
	Ribosomal proteins (Large subunit, LSU)	<i>rpl2, rpl5, rpl6, rpl10, rpl16</i>
	Ribosomal proteins (Small subunit, SSU)	<i>rps1, rps2, rps3, rps4, rps7, rps10, rps11, rps12, rps13, rps14, rps19</i>
	<b>Others</b>	
	Intron maturase	<i>matR, mttB</i>
Chloroplast	<b>Photosynthesis</b>	
	RuBisCO large subunit	<i>rbcL</i>
	Photosystem I	<i>psaA, psaB, psaC, psaI, psaJ</i>
	Assembly/stability of photosystem I	<i>ycf3, ycf4</i>
	Photosystem II	<i>psbA, psbB, psbC, psbD, psbE, psbF, psbH, psbI, psbJ, psbK, psbL, psbM, psbN, psbT, psbZ</i>
	Cytochrome b/f complex	<i>petA, petB, petD, petG, petL, petN</i>
	C-type cytochrome	<i>ccsA</i>
	ATP synthase	<i>atpA, atpB, atpE, atpF, atpH, atpI</i>
	Nicotinamide adenine dinucleotide dehydrogenase (NADH)	<i>ndhA, ndhB, ndhC, ndhD, ndhE, ndhF, ndhG, ndhH, ndhI, ndhJ, ndhK</i>
	<b>Transcription and translation</b>	
	Ribonucleic acid (RNA ) polymerase	<i>rpoA, rpoB, rpoC1, rpoC2</i>
	Ribosomal protein (Large subunit, LSU)	<i>rpl2, rpl14, rpl16, rpl20, rpl22, rpl23, rpl32, rpl33, rpl36</i>
	Ribosomal proteins (Small subunit, SSU)	<i>rps2, rps3, rps4, rps7, rps8, rps11, rps12, rps14, rps15, rps16, rps18, rps19</i>
	Transfer ribonucleic acid (tRNA)	<i>trnA, trnC, trnD, trnE, trnF, trnG, trnH, trnI, trnK, trnL, trnM, trnN, trnP, trnQ, trnR, trnS, trnT, trnV, trnW, trnY</i>
	<b>Others</b>	
	Maturase	<i>matK</i>
	Acetyl-CoA carboxylase subunit	<i>accD</i>
	Inorganic carbon uptake	<i>cemA</i>
	ATP-dependent protease subunit	<i>clpP</i>
	Conserved reading frames (ycfs)	<i>ycf1, ycf2</i>