

Sequence	SV	Region	Primer sequence (5'–3')	Tm (°C)	%GC	Product size (bp)
<i>ndhF-rpl32</i>	41.19	SSC	ndhF* – CCAATATCCCTYYTTTCCAA [79]	60.0	52.4	-
			rpl32R* – CCAATATCCCTYYTTTCCAA [79]	60.0	50.0	
			Pleuro_ndhF-F – AGGAAAGCCCACATACGACG	60.1	55.0	1068
			Pleuro_rpl32-R – ACTTTTTGACTGTCCGGTGGA	59.8	47.6	
<i>petN-psbM</i>	29.78	LSC	Pleuro_petN-F – TCTCGCTTGGGCTGCTTTAA	60.0	50.0	973
			Pleuro_psbM-R – GCGCTTATTGCTACTACGCTG	59.8	52.4	
<i>psbB-psbT</i>	38.76	LSC	Pleuro_psbB-F – TGCTACGTTTGCTTTGCTCT	58.4	45.0	800
			Pleuro_psbT-R – GGGGGCTCATTACTTCAATGGA	60.1	50.0	
<i>psbI-trnS^{GCU}</i>	41.67	LSC	Pleuro_psbI-F – AATGATCCGGGGCGTAATCC	60.0	55.0	266
			Pleuro_trnS(GCU)-R – ATGGGGAGAGATGGCTGAGT	60.0	55.0	
<i>psbK-psbI</i>	29.63	LSC	psbK – TTAGCCTTTGTTTGCAAG [80]	54.1	42.1	-
			psbI* – AGAGTTTGAGAGTAAGCAT [80]	50.7	36.8	
			Pleuro_psbI-R – GGATTACGCCCCGGATCATT	60.0	55.0	699
<i>rpl16-rps3</i>	29.68	LSC	Pleuro_rpl16-F – AACGAGTCACACTGAGCA	59.5	50.0	337
			Pleuro_rps3-R – TGCACGTGTCGAATGGATCA	60.0	50.0	
<i>rpl32-trnL^{UAG}</i>	33.88	SSC	rpl32-F – CAGTTCCAAAAAACGTACTTC [79]	54.5	36.4	-
			trnL_UAG – CTGCTTCCTAAGAGCAGCGT [79]	60.1	55.0	
<i>trnR^{UCU}-atpA</i>	31.94	LSC	Pleuro_trnR(UCU)-F – AATGAAGGGCGTCCATTGTCT	60.0	47.6	700
			Pleuro_atpA-R – GCTGGAATCAGGCCTGCTAT	59.9	55.0	
<i>trnS^{GCU}-trnG^{UCC}</i>	36.19	LSC	trnS* – AGATAGGGATTCTGAACCCTCG [79]	58.8	52.4	-
			trnG* – GTAGCGGGAATCGAACCCGCATC [79]	65.8	60.9	
			Pleuro_trnS(GCU)-F – ACGCTTTAGTCCACTCAGCC	60.0	55.0	1336
Pleuro_trnG(UCC)-R – AGCCGAGGGTTCTAGTAAACC	59.2	52.4				
<i>trnW^{CCA}-trnP^{UGG}</i>	30.61	LSC	Pleuro_trnW(CCA)-F – ATTTGAACCTACGACATCGGGT	59.8	45.5	300
			Pleuro_trnP(UGG)-R – GCTTGGTAGCGCGTTTGT	60.3	50.0	