

	No.	Sequence	No.	Sequence		
<i>trnF</i> ^{GAA} - <i>ndhJ</i> - <i>ndhK</i> - <i>ndhC</i>	529	TCGGGATAGCTCAGTTGGTA	107	GTTTCTGCTTCACGAATATG	47650	1379
<i>trnR</i> ^{ACG} - <i>trnN</i> ^{GUU} - <i>ndhF</i> - <i>rpl32</i>	554	ATTCGAACCTACGACCAGTCA	1038	TCCCTTTTTTCTGACGAATTA	107482	2196
<i>ccsA</i> - <i>ndhD</i>	549	TGAAATTGGTAGACACGCTGC	1059	ACCGAAGATTGTGTAGGTTG	109333	2728
<i>psaC</i> - <i>ndhE</i> - <i>ndhG</i>	1058	TGCTCGGGAGAAGAATAATA	1061	TTTGTGGGAACCATAAAATGT	111849	1697
<i>ndhG</i> - <i>ndhI</i> - <i>ndhA</i> - <i>ndhH</i>	1095	TGAATACCAATTTGTTGAACG	1917	TCAAGTATTCCATTTACCA	113319	1828
<i>ndhB</i>	437	TGATCTGGCATGTACAGAATG	438	AAAGAGGGTATCCTGAGCAA	133250	2221
<i>trnH</i> ^{GUG} - <i>psbA</i>	460	AAGCGTCCTGTAGTAAGAGGA	476	GGGAAACCACTGAAAATGAG	145562	1426
<i>accD</i>	423	TGGTTCAATTCAATGTTGTCT	424	ATTCAAGGGAAGGAAACCGT	55778	1430
<i>matK</i>	1785	TCTAGCACACGAAAGTCGAAGT	1784	CGATCTATTCATTCAATATTTTC	1931	936
<i>trnF</i> ^{GAA} - <i>ndhJ</i>	529	TCGGGATAGCTCAGTTGGTA	107	GTTTCTGCTTCACGAATATG	47650	1379
<i>rbcL</i>	505	AGGGAGGGACTTATGTCACCA	506	TATCTGGCTTATCCACTGGGT	53528	1443
<i>rpoB</i>	986	GCCTCTTGCTCATATCTCTC	1078	TGTGGAGCAATGAGGCATAA	23345	1666
<i>rpoC1</i>	893	TATGAGTAGGCCCCGCCAAA	895	TGGTCCTTACTGGGAAC TTGA	19989	1785
<i>ycf1</i> - <i>trnR</i> ^{ACG}	461	AAAATCTTCGTAAACCGGGC	554	ATTCGAACCTACGACCAGTCA	119956	1370