

Table S5. Length of analyzed plastid DNA and nrITS (nuclear ITS: ITS1-5.8S rDNA-ITS2) data sets.

Data set	sequence length		
<i>rpl32-trnL</i>	<i>rpl32</i> gene 18 bp	<i>rpl32-trnL</i> spacer 436 - 874 bp	-
<i>rps16-trnK</i>	-	<i>rps16-trnK</i> spacer 664 – 729 bp	-
<i>psbD-trnT</i>	-	<i>trnT-psbD</i> spacer 760 – 1076 bp	-
<i>ndhC-trnV</i>	-	<i>ndhC-trnV</i> spacer 734 – 988 bp	-
nrITS	ITS1 167 - 180 bp	5,8S rDNA 158 bp	ITS2 201 – 262 bp