

Table S1.

Selected sequences of the primer used in this study based on [28] Rønsted *et al.* (2012).

Primer name	Target	Sequence (5'–3')	Direction	Reference
ITS5	ITS region	GGA AGT AAA AGT CGT AAC AAG G	Forward	[27]
ITS4	ITS region	TCC TCC GCT TAT TGA TAT GC	Reverse	[27]
trnC	<i>trnL-trnF</i> region	CGA AAT CGG TAG ACG CTA CG	Forward	[33]
trnF	<i>trnL-trnF</i> region	ATT TGA ACT GGT GAC ACG AG	Reverse	[33]
19F	<i>matK</i> gene	CGT TCT GAC CAT ATT GCA CTA TG	Forward	[31]
2R	<i>matK</i> gene	AAC TAG TCG GAT GGA GTA G	Reverse	[32]
KatF*	<i>matK</i> gene	ATC GGG CCA TCC TAT TAG	Forward	[35]
1326R*	<i>matK</i> gene	TCT AGC ACA CGA AAG TCG AAG T	Reverse	[36]
nad1eB	<i>nadl</i> gene	GCA TTA CGA TCT GCA GCT CA	Forward	[34]
nad1eCR	<i>nadl</i> gene	GGA GCT CGA TTA GTT TCT GC	Reverse	[34]
nad1iB2*	<i>nadl</i> gene	GTC GAG CAT ACG ACG ATG CCG C	Forward	[37]
nad1iB2R*	<i>nadl</i> gene	GGC GGC ATC GTC GTA TGC TCG	Reverse	[37]

* Primers used for Sanger sequencing.

References

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