

**Table S3.**

1. Arct\_1F (5'-3') TAAACTATTCCCTGGTACA  
H16299 (5'-3') GGAGCGAGAAGAGGTACACGT  
(ca 270 bp exl. primers)
2. Arct\_1F (5'-3') TAAACTATTCCCTGGTACA  
OP55R (5'-3') CTCCCGGACTAAGTGAAATACATG  
(ca 209 bp exl. primers)
3. Arct\_1F (5'-3') TAAACTATTCCCTGGTACA  
Arct\_1R (5'-3') AGGTATTCGAGGACATAC  
(61 bp exl. primers)
4. L16164 (5'-3') GCCCATGCATATAAGCATG  
H16299 (5'-3') GGAGCGAGAAGAGGTACACGT  
(132 bp exl. primers)
5. OP55F (5'-3') CCCCATGCATATAAGCATGTACAT  
OP55R (5'-3') CTCCCGGACTAAGTGAAATACATG  
(67 bp exl. primers)
6. OP54F (5'-3') GTATAGTCTGTAACGATGTAT  
OP54R (5'-3') AGGTACACGTAACGATGTAT  
(55 bp exl. primers)

Figure below shows coverage of primer pairs: Black bar at bottom represents entire fragment of the mitochondrial control region, spanning ca. 270 bp. Dotted line indicates a ca. 70 bp stretch of repetitions and extremely slight variation, which was not included.

