

Gene	Primer sequence	Fragment size (bp)
16S ribosomal RNA	F: AAG ACG AGA AGA CCC TAT GGA R: CCG GTC TGA ACT CAG ATC ACG T	~350
12S ribosomal RNA	F: ATA CCG CCA TCT TCA GCA AA R: CAT AGG TTA CAC CTT GAC CTA ACG	89
Control region, fragment 1	F: AGC GGG TTG TTG GTT TCA C R: CCG TTC TAG TCA ACA TGC GTA	78
Control region, fragment 2	F: CGC ATG TTG ACT AGA ACG GAT T R: CCA TGC ATA TAA GCA AGC ACA	104
Control region, fragment 3	F: ACA TTC AAG GTG GAC ATA GGG R: ACA CGG ACT TTC CAC TCC AC	117