

Table S2 Primers and thermal cycling conditions used in amplification and sequencing of genes.

Gene	Primer name	Primer sequence	Reference	PCR conditions			
				Denature	Annealing	Extension	Cycles
ND3	L10755	5'-GACTTCCAATCTTTAAAATCTG-3'	Chesser (1999)	94°C	58°C	72°C	35
	H11151	5'-GATTTGTTGAGCCGAAATCAAC-3'		30s	40s	40s	
Cyt-b	CytbF25	5' GGCTCTCAATCTTCGTAAAAACC-3'	This study	94°C	60°C	72°C	35
	CytbR649	5'-GGGTGGAATGGGATTTTGTC-3'	This study	30s	60s	90s	
	CytbF409	5'-GTAGGCTACGTCCTACCCTGAG-3'	This study				
	CytbH16065	5'-GAGTCTTCAGTCTCTGGTTTACAAGAC-3	Helm-Bychowski & Cracraft (1993)				
TFGβ2	TGF5	5'-GAAGCGTGCTCTAGATGCTG-3'	Primmer <i>et al.</i> (2002)	94°C	56°C	72°C	35
	TGF6	5'-AGGCAGCAATTATCCTGCAC-3'		30s	30s	45s	

PCR amplifications were performed in 15µl volumes with 2µl total genomic DNA, 9.7µl ddH₂O, 1.5µl 10X PCR buffer, 0.75µl MgCl₂ (50mM), 0.15µl dNTP (2.5 mM of each), 0.45µl of each primer (10mM), and 0.15µl Taq DNA polymerase (5 units/µl).

References

- Chesser RT (1999) Molecular systematics of the Rhinocryptid genus *Pteroptochos*. *The Condor*, **101**, 439-446.
- Helm-Bychowski K, Cracraft J (1993) Recovering phylogenetic signal from DNA sequences: relationships within the Corvine assemblage (Class Aves) as inferred from complete sequences of the mitochondrial DNA cytochrome-*b* gene. *Molecular Biology and Evolution*, **10**, 1196-1214.
- Primmer CR, Borge T, Lindell J, Setre GP (2002) Single-nucleotide polymorphism characterization in species with limited available sequence information: high nucleotide diversity revealed in the avian genome. *Molecular Ecology*, **11**, 603-612.