

Additional file 3 – Sequences produced for this study.

Table S1. GenBank accession numbers for the *Ericabatrachus baleensis* sequences generated in the present study.

No./ Molecular Accession	Voucher and Field Number	Locality	12S	16S	28S	H3A	RAG1
T880	ZNHM-AAU-A2013-003 SL 065	Fute, Harena Forest, Bale Mountains	KF938362	KF938365	KF938368	KF938369	KF938370
T1083	ZNHM-AAU-A2013-001 AK 2020	Fute, Harena Forest, Bale Mountains	KF938363	KF938366	-	-	KF938371
T1084	ZNHM-AAU-A2013-002 AK 2022	Fute, Harena Forest, Bale Mountains	KF938364	KF938367	-	-	KF938372

Table S2. Primers used in this study.

Refer to text for abbreviations.

Gene	Primer
<i>12S</i>	12S A-L: AACTGGGATTAGATACCCCACTAT 12S F-H: CTTGGCTCGTAGTCCCTGGCG
<i>16S</i>	16AR: CGCCTGTTTATCAAAAACAT 16Br: CCGGTCTGAACTCAGATCACGT
<i>RAG1</i>	RAG 1c: GGAGATGTTAGTGAGAARCA YGG RAG 1e: TCCGCTGCATTTCCRATGTCRCA
<i>28S</i>	28Sv: AAGGTAGCCAAATGCCTCATC 28Sjj: AGTAGGGTAAAACCT
<i>H3A</i>	h3F: ATGGCTCGTACCAAGCAGACVGC h3R: TATCCTTRGGCATRATRGTGAC

Note on intraspecific variation

The three samples of *Ericabatrachus baleensis* presented no variation at the nuclear RAG1 and mitochondrial 12S and 16S.