

Appendix III(c). Table 1C. Diagnostic nucleotide sites from the D-loop fragment Pairwise Aggregate Analysis (PAA) for Lepilemur.

	11111111111111111111111111222222222222222333333333344445555
	11222333355589001112222222333345689234456777888001112357977933335
	091450347458783701701345670279090751129360235678120477077656356784
ANAL5	TAAAC-CTA--CAACTTCGGAGCCACAA-AAGTTAAACGATTTTTTAATCATAACCCAAAACTT
DAR4.7	...T-...--T...C.....G.....C.....
LOK04.2	...T-...--T.....-.....C.....
FARY5.1	...T-GCCCC-.....A.....-.....G.....C.....A.....A.....
ANT5.2	...T-...--.....A...T.T...-...A.....C.....A.....GAC.
LAZA5.1	...T-.C.CC-.....AAG.....AC.....A.....T.C.
BEZ14-.....A...T-T.....TT.....C.....A.....C.
AND65	.G.....-.....AA..T-T.....TT.....G.....C.....
MOR117-C-.C-.G...A...GGTC.GG-.A...CC...C...GCA...A...GC.A.
ZOMB8-C-.....A...-C.GG-.....GTT.G...C.GT...G.....T.A.
BEMA5-T-.C-.....AA...TC.GG-.....CT...A..C.....G.....A.
MAR1	...C...C-.C...A..AG.G-.CC-.AC.....A.....G.
HIH21	...C...C-T...C.A..AG.-.CT-.AG.....CC.....T.....C.
KIBO22	...T-...C-.....C.AAGA.....AC.....A.....C.
JAM4.27	.G...C-.C-.....A...-TT.GG-.A...T...ACC..G.....A.....A.
TAD25	..GTTAC.ACA...C...AA...G...AC..CT...AA...C...ATA.C..T.A.
M83B	..GTTAC.ACAC...C...AA...A...AC..TT...AA...C.T...A.A.C..T.A.
FAN10	..GTTAC.ACA...C...AA...G...AC..CT...CCAG...C...A.A.C..T.A.
AND20	C..C.TACGACAT...C...AG...G...AC...T...CACG.C.AG.CA.A.C..T.A.
RANO234	..GAC.G.-CT..TG.....C.....C.....A.T.....CA
KALA24	..GTTA..CACT...C...TAA-.G...AC..T...GC..TC.T...G.ACTC.C.A.
JAR2TAC.GCTT...C...T.GT..G.-T.ACC.CT...CA...T...A.ACC..T.A.
NARA4.20TAC.ACTG...C...TAAT..G...GAA.TCT...A...C.A.ATC..T.A.
MAS6.12	...T.AAC.ACTT...C...TAAT..G..A..TC..CT.A...A...A.ACC..T.A.
MAS6.12	...T.AAC.ACTT...C...TAAT..G..A..TC..CT.A...A...A.ACC..T.A.