

Table S3. Contaminant sequences nucleotide misincorporation rate within each mitochondrial fragment.

Mitochondrial fragment	Number of clones	Length of fragment	Number of nucleotide misincorporations	<i>m</i>
L16518/H47	19	97	5	0.00271
L1177/H1238	14	61	3	0.00351
L1720/H1833	14	112	0	0
L11261/H11325	13	63	8	0.00977
L11456/H11536	29	79	0	0
L14097/H14204	22	106	5	0.00214
L16303/H16407	29	103	6	0.00201

NOTE. — Mitochondrial fragments are numbered with reference to the Cambridge reference sequence (rCRS) [28].