Supplementary Material 6. Mitochondrial *cox1* gene and genome sequence divergence between African savanna elephant (*Loxodonta africana*, GenBank accession number NC_000934) and Asian elephant (*Elephas maximus*, NC_005129)

	Pairwise alignment length	Sequence divergence
coxI	1,548 bp	4.52%
Entire mt genome (D-loop excluded)	15,419 bp	5.03%
Entire mt genome (D-loop included)	16,905 bp	5.16%