

Table 1. Molecular diversity statistics for the seven maximally differentiated mitochondrial haplogroups defined by SAMOVA

Haplogroup (<i>k</i> = 7)	Sites (<i>n</i> = 20)	Average pairwise diffs.	Nucleotide diversity π	Coalescent θ 3 runs, \pm SD	Growth parameter estimate (<i>g</i>)	Tajima's <i>D</i>	Fu's <i>F</i>
A (<i>n</i> = 12)	28, 29	1.894 \pm 1.160	0.008 \pm 0.005	(0.084) – (0.099) \pm 0.016 - 0.019	(-2.941) – (-3.773) \pm 0.490 – 0.665	-1.138 (<i>P</i> = 0.150)	-1.475 (<i>P</i> = 0.130)
B (<i>n</i> = 16)	25-27	1.492 \pm 0.949	0.006 \pm 0.005	(0.067) – (0.092) \pm 0.017 - 0.029	(544.6) – (630.3) [†] \pm 52.21 – 137.0	-1.032 (<i>P</i> = 0.165)	-2.659 (<i>P</i> = 0.027)*
C1/2 (<i>n</i> = 33)	1-4	5.915 \pm 2.897	0.028 \pm 0.015	(0.053) – (0.058) \pm 0.008 - 0.009	(-1.811) – (0.323) \pm 8.523 – 8.622	-1.608 (<i>P</i> = 0.032)*	-2.789 (<i>P</i> = 0.159)
D1 (<i>n</i> = 15)	5, 6	3.790 \pm 2.021	0.016 \pm 0.010	(0.014) – (0.016) \pm 0.004	(-9.431) – (3.076) \pm 41.82 – 45.10	0.464 (<i>P</i> = 0.740)	1.992 (<i>P</i> = 0.839)
D2 (<i>n</i> = 19)	23	1.743 \pm 1.059	0.008 \pm 0.005	(0.013) – (0.018) \pm 0.004 - 0.006	(216.8) – (327.4) \pm 141.9 – 157.0	-0.434 (<i>P</i> = 0.389)	-0.742 (<i>P</i> = 0.338)
D3 (<i>n</i> = 44)	8, 12-14, 16,18, 20, 24	0.984 \pm 0.679	0.004 \pm 0.003	(0.025) – (0.180) \pm 0.005 - 0.019	(389.8) – (1722) [†] \pm 63.47 – 188.2	-1.820 (<i>P</i> = 0.015)*	-4.490 (<i>P</i> = 0.006)*
Ivindo (<i>n</i> = 8)	19	13.821 \pm 6.956	0.059 \pm 0.034	(0.027) \pm 0.007	(-20.40) – (-21.13) \pm 12.46 – 12.68	-1.129 (<i>P</i> = 0.918)	7.986 (<i>P</i> = 0.995)

*Value significant at $P \leq 0.05$.

† Standard deviation is greater than three times the magnitude of the growth parameter estimate.