

Table S5. Summary of variants identified on mitochondrial and Y chromosome

Variant Type	No. of SNPs (% novel)	No. of Indels and block substitutions (% of novel)
Mitochondria	285 (53.7%)	21 (90.5%)
Noncoding	90 (54.4%)	20 (90.0%)
Coding	195 (53.3%)	1 (100%)
Y Chromosome	8,673 (54.3%)	2,348 (65.2%)
Intergenic	7,915 (54.0%)	2,170 (63.8%)
Intragenic	758 (57.5%)	178 (82.3%)
Intron	578 (62.6%)	144 (84.0%)
UTR	139 (43.9%)	28 (82.1%)
Splicing acceptor site	1 (0%)	1 (100%)
Splicing donor site	0	0
Coding domain	40 (32.5%)	5 (40.0%)