

Pos. ^a	Primer pair (5'->3')	Background ^b	Overlap ^c
1174	CGGGCACGCGTTTTAATTACTTTGTAAAGC GTAATTAACGCGTGCCCGAAAAGGCGC	Cos.	partial
1174	GGGCAAGCGTTTTAATTACTTTGTAAAGC GTAATTAACGCTTGCCCAAAAAGGCGC	Afr.	partial
1155	CGGGCAAGCGTTTTAATTACTTTGTAAAGCTGCATTTTTG CAAAAATGCAGCTTAACAAAGTAATTAACGCTTGCCCG	Cos.	full
1063	GCCGTCTTAATGTGTGTTTGTGTGCGAGCCAAGTGC CGACACAAACACACATTAAGACGGCAAAAAAATC	Cos.	partial
1063	CTTAATGTTTGTGTTTGTGTGCGAGCCAAGTGC CGACACAAACAAACATTAAGACAGC	Afr.	partial
765	CAATTTTGTATTTTTTAAATCTATGCTTTGATTTTAG GCATAGATTTAAAAATAACAAAATTGTTTTAAATTTTATAAC	Cos.	partial
67	CGCGACTGGGCCTCAGAGTCAAATAG CTGAGGCCAGTCGCGGCTGAAGATTCGC	Afr.	partial

^aPosition (pos.) in bp before *CG9509* start codon

^bMutations were introduced into either a Cosmopolitan (Cos.) or sub-Saharan African background (Afr.).

^cOverlap indicates degree of primer pair overlap.

