Site 23731, Region: S Inferred minimum changes on tree: 2 Consistency: 0.500, Nearest homoplasic site is 22bp away Prop. nearest neighbour has homoplasy: 0.987 (76/77) Seen from 16 originating labs across 10 countries. 6000 -4000 density 2000 -

0.00050

cophenetic distance

0.00075

0.00100

0.00125

0 -

0.00000

0.00025