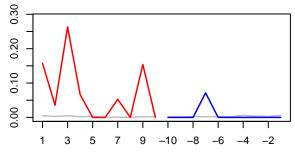


edit distance: default

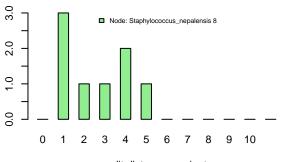


C2T/G2A rate

Read Position

Staphylococcus_nepalensis;_NZ_CP017460.1 100%
NA;_NZ_CP065425.1 001%
NA%

Pool-3-038-AZP-223_S42.unmapped.rma6



read count

edit distance: ancient

out distance. ancient	
Node	Staphylococcus_nepalensis
Top Reference	Staphylococcus_nepalensis
all reads	92
nonDup	76
readDis	0.97
nonStacked	74
destacking?	Off
C>T_1	0.1579
G>A1	0
mean length (sd)	63 (19.98)