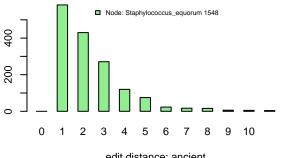


C2T/G2A rate

Read Position
Staphylococcus_equorum;_NZ_CP068576.1 100%
NA;_NZ_CP051881.1 000%
NA;_NZ_CP065795.1 000%
NA;_NZ_CP051882.1 000%
Staphylococcus;_NZ_CP033460.1 000%
NA;_NZ_CP041880.1 000%
NA;_NZ_CP069389.1 000%
NA;_NZ_CP080326.1 000%
NA;_NZ_CP068479.1 000%
NA;_NZ_CP065792.1 000%

Pool-4-046-AZP-283_S47.unmapped.rma6



read count

edit distance: ancient	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	6701
nonDup	5788
readDis	0.669
nonStacked	2875
destacking?	Off
C>T_1	0.3392
G>A1	0.0045
mean length (sd)	61 (18.359)