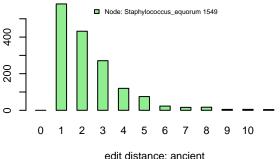


C2T/G2A rate

Read Fosition		
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

ean	distance: ancient
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	6703
nonDup	5796
readDis	0.667
nonStacked	2868
destacking?	Off
C>T_1	0.3396
G>A1	0.0045
mean length (sd)	61 (18.36)