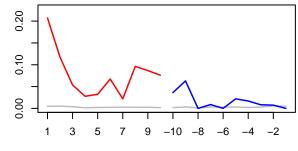


read count

C2T/G2A rate

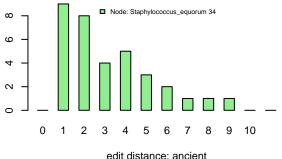
Damage plot for Staphylococcus_equorum node



Read Position

Staphylococcus_equorum;_NZ_CP068063.1 100%
NA;_NZ_CP065792.1 000%
NA;_NZ_CP080323.1 000%
NA;_NZ_CP041879.1 000%
NA;_NZ_CP065795.1 000%
NA;_NZ_CP070605.1 000%
NA;_NZ_CP070607.1 000%
NA;_NZ_CP065960.1 000%
NA;_NZ_CP077960.1 000%
NA;_NZ_CP069389.1 000%

Pool-3-016-AZP-201_S20.unmapped.rma6



read count

edit distance, ancient	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	184
nonDup	161
readDis	0.986
nonStacked	159
destacking?	Off
C>T_1	0.2072
G>A1	0
mean length (sd)	57 (17.037)