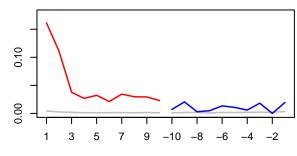


read count

C2T/G2A rate

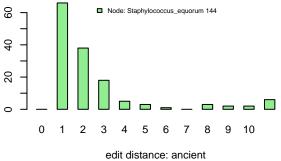
Damage plot for Staphylococcus_equorum node



Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
NA;_NZ_CP059679.1 000%
NA;_NZ_CP065795.1 000%
NA;_NZ_CP051882.1 000%
NA;_NZ_CP065793.1 000%
NA;_NZ_CP075505.1 000%
NA;_NZ_CP075504.1 000%
NA;_NZ_CP075503.1 000%
NA;_NZ_CP069390.1 000%
Staphylococcus;_NZ_CP033460.1 000%

Pool-4-027-AZP-264_S28.unmapped.rma6



read count

edit distance, ancient	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	1277
nonDup	1103
readDis	0.902
nonStacked	920
destacking?	Off
C>T_1	0.1612
G>A1	0.0194
mean length (sd)	65 (16.764)