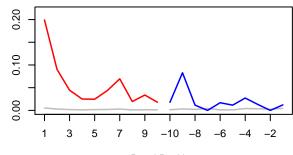


read count

C2T/G2A rate

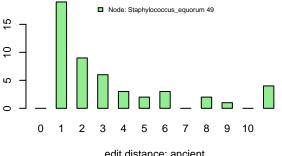
Damage plot for Staphylococcus_equorum node



Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
Staphylococcus;_NZ_JAESMX010000012.1 005%
NA;_NZ_JAJNIG010000028.1 003%
NA;_NZ_JAJNIH010000031.1 003%
Staphylococcus;_NZ_JAESJX010000019.1 003%
NA;_NZ_JAJNIV010000033.1 002%
Staphylococcus;_NZ_JAESMS010000002.1 002%
Staphylococcus;_NZ_JAFENQ010000028.1 002%
Staphylococcus;_NZ_JAESMK010000020.1 002%
Staphylococcus;_NZ_JAESLM010000022.1 002%

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



read count

edit distance: ancient		
Node	Staphylococcus_equorum	
Top Reference	Staphylococcus_equorum	
all reads	219	
nonDup	192	
readDis	0.937	
nonStacked	164	
destacking?	Off	
C>T_1	0.1991	
G>A1	0.0121	
mean length (sd)	63 (16.012)	