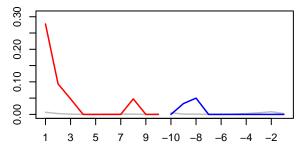


Damage plot for Staphylococcus_equorum node

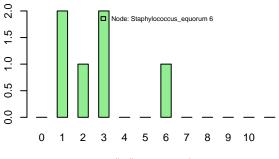


C2T/G2A rate

Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%	
NA%	

Pool-4-000-UDI_54_S1.unmapped.rma6



edit distance: ancient

cuit distance, anoiem	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	33
nonDup	29
readDis	1
nonStacked	29
destacking?	Off
C>T_1	0.2778
G>A1	0
mean length (sd)	61 (16.506)