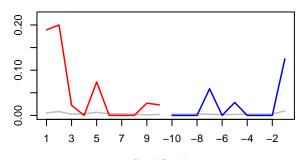


## Damage plot for Staphylococcus\_equorum node

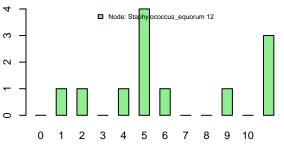


C2T/G2A rate

Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
NA;_NZ_CP075505.1 002%
NA;_NZ_CP075504.1 002%
NA;_NZ_CP075503.1 002%
NA;_NZ_CP022983.1 002%
NA%

## Pool-3-052-AZP-237\_S56.unmapped.rma6



read count

edit distance: ancient

cuit distance, ancient	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	55
nonDup	50
readDis	1
nonStacked	50
destacking?	Off
C>T_1	0.1892
G>A1	0.125
mean length (sd)	63 (17.729)