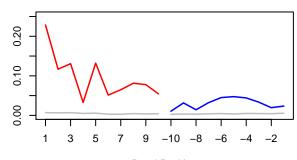


## Damage plot for Staphylococcus\_equorum node

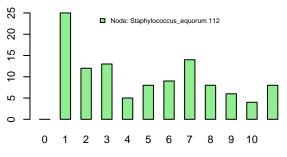


C2T/G2A rate

Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
Staphylococcus;_NZ_CP068241.1 001%
NA;_NZ_CP041879.1 000%
NA;_NZ_CP059679.1 000%
NA;_NZ_CP075505.1 000%
NA;_NZ_CP047394.1 000%
NA;_NZ_CP075504.1 000%
NA;_NZ_LT906462.1 000%
NA;_NZ_CP075503.1 000%
NA%

## Pool-3-024-AZP-209\_S28.unmapped.rma6



read count

edit distance: ancient

Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	389
nonDup	327
readDis	0.917
nonStacked	284
destacking?	Off
C>T_1	0.2286
G>A1	0.0236
mean length (sd)	65 (16.385)