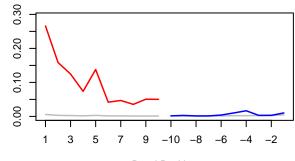


Damage plot for Staphylococcus_equorum node

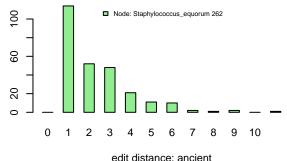


C2T/G2A rate

Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%		
NA;_NZ_CP059679.1 000%		
NA;_NZ_CP065792.1 000%		
NA;_NZ_CP077960.1 000%		
NA;_NZ_CP068479.1 000%		
Staphylococcus;_NZ_CP064056.1 000%		
NA%		

Pool-4-042-AZP-279_S43.unmapped.rma6



read count

cuit distance, anoiem	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	1369
nonDup	1162
readDis	0.897
nonStacked	962
destacking?	Off
C>T_1	0.2659
G>A1	0.0101
mean length (sd)	63 (17.306)