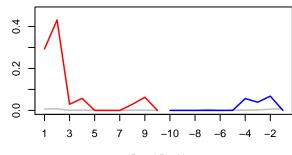


Damage plot for Staphylococcus_simulans node

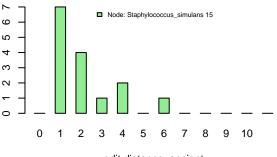


C2T/G2A rate

Read Position

NA;_NZ_CP059679.1 100%		
Staphylococcus_simulans;_NZ_CP014016.2 011%		
Staphylococcus_simulans;_NZ_CP023497.1 009%		
Staphylococcus_simulans;_NZ_LT963435.1 004%		
Staphylococcus_simulans;_NZ_CP017428.1 004%		
Staphylococcus_simulans;_NZ_LS483313.1 002%		
Staphylococcus_simulans;_NZ_LR134264.1 002%		
NA%		
NA%		
NA%		

Pool-3-045-AZP-230_S49.unmapped.rma6



read count

edit distance: ancient

Node	Staphylococcus_simulans
Top Reference	NA
all reads	82
nonDup	68
readDis	0.734
nonStacked	44
destacking?	Off
C>T_1	0.2927
G>A1	0
mean length (sd)	59 (16.235)