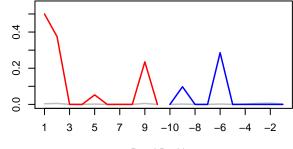


Damage plot for Staphylococcus_hominis node



C2T/G2A rate

Read Position

NA;_NZ_CP075505.1 100%	
Staphylococcus_hominis;_NZ_CP014107.1 022%	
Staphylococcus_hominis;_NZ_CP054883.1 016%	
Staphylococcus_hominis;_NZ_LT963438.1 013%	
Staphylococcus_hominis;_NZ_CP031277.1 008%	
NA%	

read count

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

edit distance: ancient

Node	Staphylococcus_hominis
Top Reference	NA
all reads	45
nonDup	40
readDis	0.901
nonStacked	36
destacking?	Off
C>T_1	0.5
G>A1	0
mean length (sd)	61 (14.257)