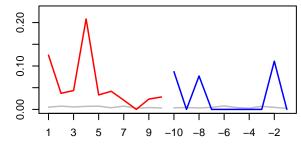


read count

C2T/G2A rate

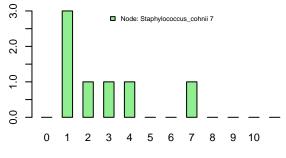
Damage plot for Staphylococcus_cohnii node



Read Position

Staphylococcus_cohnii;_NZ_LT963440.1 100%		
Staphylococcus;_NZ_CP068241.1 064%		
NA;_NZ_CP046640.1 012%		
NA%		

Pool-3-041-AZP-226_S45.unmapped.rma6



read count

edit distance: ancient

Node	Staphylococcus_cohnii
Top Reference	Staphylococcus_cohnii
all reads	33
nonDup	27
readDis	0.932
nonStacked	25
destacking?	Off
C>T_1	0.125
G>A1	0
mean length (sd)	62 (15.124)