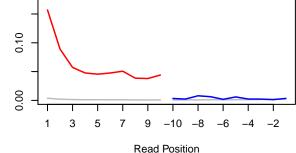




edit distance: default

0

C2T/G2A rate



Staphylococcus_nepalensis;_NZ_CP017460.1 100% Staphylococcus;_NZ_JAESKI010000003.1 030% Staphylococcus;_NZ_JAESKF010000002.1 026% Staphylococcus;_NZ_JAESKI010000008.1 020% Staphylococcus;_NZ_JAESKI010000007.1 020% Staphylococcus;_NZ_JAESKF010000006.1 019% Staphylococcus;_NZ_JAESKI010000013.1 017% Staphylococcus;_NZ_JAESKI010000002.1 016% Staphylococcus; NZ_JAESKF010000008.1 015%

Staphylococcus; NZ_JAESKF010000020.1 012%

250 Node: Staphylococcus_nepalensis 571 20 50

Pool-4-031-AZP-268_S32.unmapped.rma6

10

edit distance:	ancient

ead count

0

10

edit distance: ancient		
Node	Staphylococcus_nepalensis	
Top Reference	Staphylococcus_nepalensis	
all reads	6954	
nonDup	6021	
readDis	0.816	
nonStacked	4038	
destacking?	Off	
C>T_1	0.1569	
G>A1	0.0034	
mean length (sd)	63 (17.439)	