

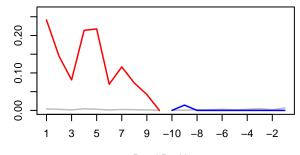
0

C2T/G2A rate

2 3

edit distance: default

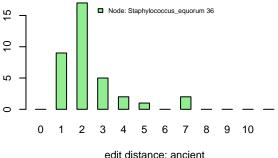
Damage plot for Staphylococcus_equorum node



Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
NA%

Pool-4-029-AZP-266_S30.unmapped.rma6



read count

10

euit distance, ancient	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	179
nonDup	149
readDis	0.933
nonStacked	129
destacking?	Off
C>T_1	0.2414
G>A1	0
mean length (sd)	64 (18.763)