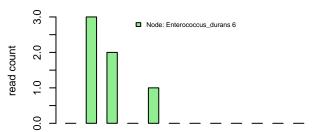


C2T/G2A rate

Read Position

Enterococcus_durans;_NZ_LR607372.1 100%
NA;_NZ_AP024685.1 002%
Enterococcus;_NZ_AP025635.1 002%
NA%



Pool-3-038-AZP-223_S42.unmapped.rma6

edit distance: ancient

10

0

1 2 3

Node	Enterococcus_durans
Top Reference	Enterococcus_durans
all reads	61
nonDup	51
readDis	0.799
nonStacked	39
destacking?	Off
C>T_1	0.2258
G>A1	0
mean length (sd)	63 (15.011)