

read count

C2T/G2A rate

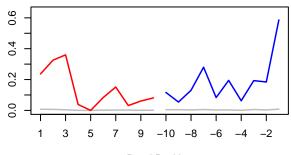
15

2

0 1 2 3



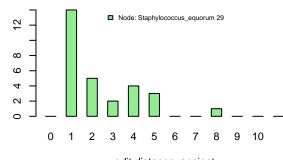
Damage plot for Staphylococcus_equorum node



Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
NA;_NZ_CP059679.1 008%
NA;_NZ_CP075505.1 004%
NA;_NZ_CP075504.1 004%
NA;_NZ_CP075503.1 004%
NA;_NZ_CP080323.1 002%
NA;_NZ_CP077960.1 002%
NA;_NZ_CP069389.1 002%
NA%
NA%

EASI273.unmapped.rma6



read count

10

edit distance: ancient

Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	49
nonDup	48
readDis	1
nonStacked	48
destacking?	Off
C>T_1	0.2364
G>A1	0.5862
mean length (sd)	49 (16.529)