EASI273.unmapped.rma6

Node: Staphylococcus_succinus 168

read count

edit distance: default

EASI273.unmapped.rma6

Node: Staphylococcus_succinus 45

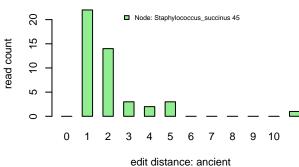read count

edit distance: ancient

Damage plot for Staphylococcus_succinus node

C2T/G2A rate

Read Position

Node	Staphylococcus_succinus
Top Reference	Staphylococcus_succinus
all reads	168
nonDup	166
readDis	0.988
nonStacked	162
destacking?	Off
C>T_1	0.2778
G>A_−1	0.5294
mean length (sd)	47 (14.587)

Staphylococcus_succinus;_NZ_CP018199.1 100%
NA;_NZ_CP022047.2 000%
NA%
NA%
NA%
NA%
NA%
NA%
NA%