Pool−4−046−AZP−283_S47.unmapped.rma6

□ Node: Staphylococcus_cohnii 290

read count

edit distance: default

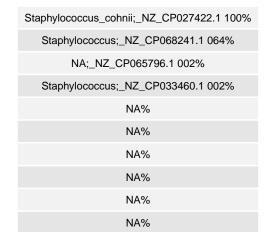## Pool−4−046−AZP−283_S47.unmapped.rma6

□ Node: Staphylococcus_cohnii 36

read count

edit distance: ancient

Damage plot for Staphylococcus_cohnii node
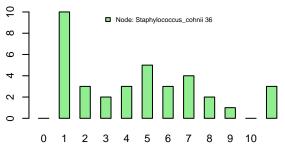
C2T/G2A rate

Read Position

Node	Staphylococcus_cohnii
Top Reference	Staphylococcus_cohnii
all reads	81
nonDup	74
readDis	0.677
nonStacked	42
destacking?	Off
C>T_1	0.1552
G>A_−1	0.0392
mean length (sd)	60 (17.302)

Staphylococcus_cohnii;_NZ_CP027422.1 100%
Staphylococcus;_NZ_CP068241.1 064%
NA;_NZ_CP065796.1 002%
Staphylococcus;_NZ_CP033460.1 002%
NA%
NA%
NA%
NA%
NA%
NA%