AT+Mock−1:50−3.rma6

read count

Node: Staphylococcus_aureus 971

edit distance: default

AT+Mock−1:50−3.rma6

read count

Node: Staphylococcus_aureus 0

edit distance: ancient

Node	Staphylococcus_aureus
Top Reference	Staphylococcus_aureus
assigned reads	1528
all reads	1176
nonDup	1176
readDis	0.784
nonStacked	893
destacking?	On
C>T_1	0
G>A_−1	0
mean length (sd)	75 (1.697)

Damage plot for Staphylococcus_aureus node

C2T/G2A rate

Read Position

Staphylococcus_aureus;_CP007690.1 100%

hylococcus_aureus_subsp._aureus_USA300_TCH1516;_CP000730.1 099%

hylococcus_aureus_subsp._aureus_USA300_FPR3757;_CP000255.1 099%

Staphylococcus_aureus_USA300−ISMMS1;_CP007176.1 098%

Staphylococcus_aureus_subsp._aureus;_CP012119.1 098%

Staphylococcus_aureus_subsp._aureus;_CP012120.1 097%

taphylococcus_aureus_subsp._aureus_NCTC_8325;_CP000253.1 090%

Staphylococcus_aureus_subsp._aureus_COL;_CP000046.1 089%

taphylococcus_aureus_subsp._aureus_str._Newman;_AP009351.1 089%

taphylococcus_aureus_subsp._aureus_DSM_20231;_CP011526.1 087%