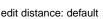


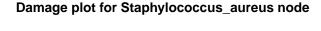
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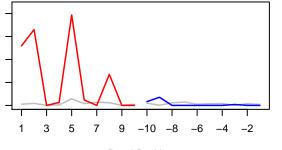
0.20

0.00

S2T/G2A rate 0.10







Read Position

aphylococcus_aureus_subsp._aureus_NCTC_8325;_NC_007795.1 035%

Staphylococcus_aureus_RF122;_NC_007622.1 029%

------ 04057 NO 000440 4 0470/

Staphylococcus_aureus_subsp._aureus_SA957;_NC_022442.1 017%

Staphylococcus_aureus_subsp._aureus_SA268;_NZ_CP006630.1 023%

Staphylococcus_aureus;_NZ_CP060603.1 100%

Staphylococcus_aureus_subsp._aureus_LGA251;_NC_017349.1 017%

Staphylococcus_aureus_subsp._aureus_TW20;_NC_017331.1 017%

Staphylococcus_aureus_subsp._aureus_6850;_NC_022222.1 017%

Staphylococcus_aureus_subsp._aureus_JKD6159;_NC_017338.1 017%

Staphylococcus_aureus_subsp._aureus_SA40;_NC_022443.1 017%

read count

0.0

Pool-3-024-AZP-209_S28.unmapped.rma6

3

Node	Staphylococcus_aureus
Top Reference	Staphylococcus_aureus
all reads	24
nonDup	22
readDis	0.785
nonStacked	17
destacking?	Off
C>T_1	0.1296
G>A1	0
mean length (sd)	61 (19.13)