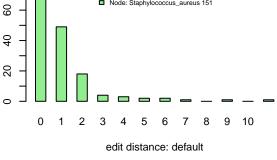
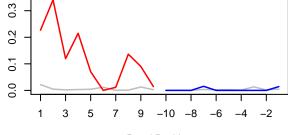


read count

C2T/G2A rate



## Damage plot for Staphylococcus\_aureus node



Read Position

NA;_NZ_CP059679.1 100%
Staphylococcus_aureus;_NZ_CP047833.1 004%
Staphylococcus_aureus;_NZ_CP047798.1 003%
Staphylococcus_aureus;_NZ_CP032160.1 003%
Staphylococcus_aureus;_NZ_CP083259.1 003%
Staphylococcus_aureus;_NZ_CP047828.1 003%
Staphylococcus_aureus;_NZ_CP076660.1 003%
Staphylococcus_aureus;_NZ_CP047837.1 003%
Staphylococcus_aureus;_NZ_CP077931.1 003%
Staphylococcus_aureus;_NZ_CP077894.1 003%

## Node: Staphylococcus\_aureus 26 Node: Staphylococcus\_aureus 26 Node: Staphylococcus\_aureus 26 Node: Staphylococcus\_aureus 26

edit distance: ancient

read count

Pool-3-045-AZP-230\_S49.unmapped.rma6

Node	Staphylococcus_aureus
Top Reference	NA
all reads	124
nonDup	99
readDis	0.923
nonStacked	89
destacking?	Off
C>T_1	0.2267
G>A1	0.0143

60 (16.384)

mean length (sd)