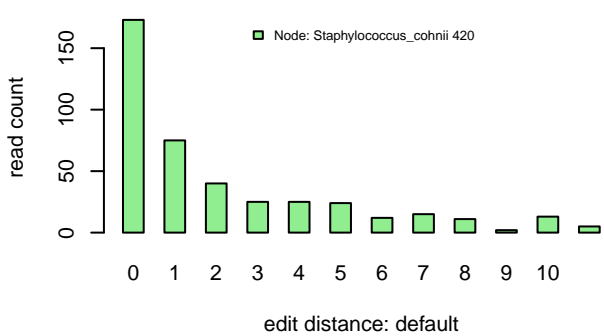
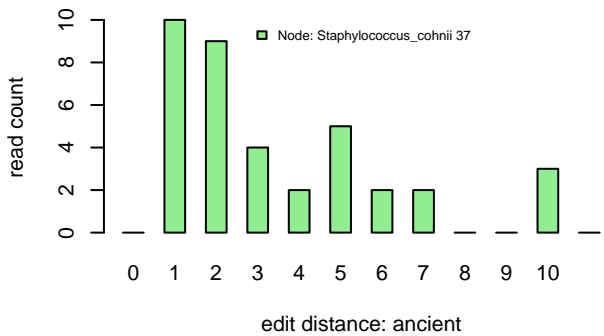


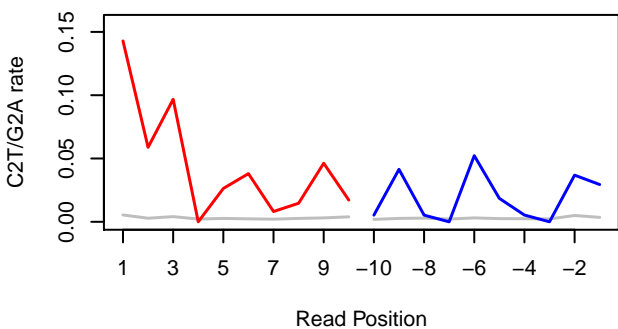
Pool-4-031-AZP-268_S32.unmapped.rma6



Pool-4-031-AZP-268_S32.unmapped.rma6



Damage plot for Staphylococcus_cohnii node



Node	Staphylococcus_cohnii
Top Reference	NA
all reads	189
nonDup	163
readDis	0.676
nonStacked	98
destacking?	Off
C>T_1	0.1429
G>A_-1	0.0294
mean length (sd)	63 (17.841)

NA;_NZ_CP059679.1	100%
Staphylococcus_cohnii;_NZ_CP054807.1	094%
Staphylococcus_cohnii;_NZ_CP033735.1	077%
Staphylococcus_cohnii;_NZ_LT963440.1	075%
Staphylococcus_cohnii;_NZ_CP027422.1	025%
Staphylococcus_cohnii;_NZ_CP054808.1	012%
Staphylococcus;_NZ_CP068241.1	012%
Staphylococcus;_NZ_CP033460.1	004%
Staphylococcus_cohnii;_NZ_CP027420.1	003%
Staphylococcus_cohnii;_NZ_CP027421.1	003%