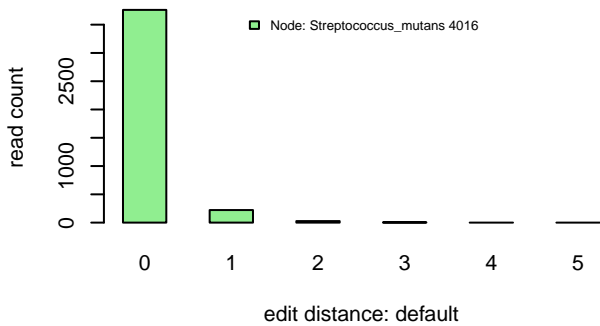
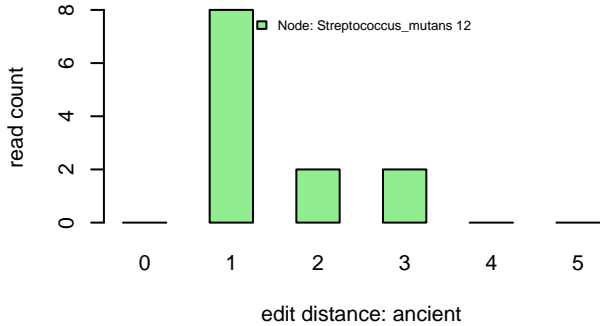


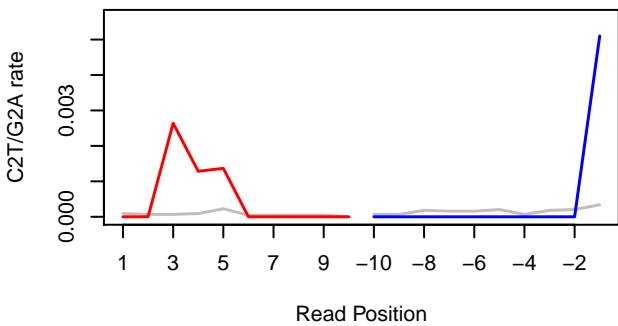
AT+Mock-1:50-1.rma6



AT+Mock-1:50-1.rma6



Damage plot for Streptococcus\_mutans node



Node	Streptococcus_mutans
Top Reference	Streptococcus_mutans_UA159-FR
assigned reads	10890
all reads	10859
nonDup	10859
readDis	0.538
nonStacked	3989
destacking?	On
C>T_1	0
G>A_-1	0.0051
mean length (sd)	75 (2.248)

Streptococcus_mutans_UA159-FR;_CP007016.1	100%
Streptococcus_mutans_UA159;_AE014133.2	100%
Streptococcus_mutans_GS-5;_CP003686.1	077%
Streptococcus_mutans;_CP013237.1	076%
Streptococcus_mutans_NN2025;_AP010655.1	069%
Streptococcus_mutans_LJ23;_AP012336.1	068%
Streptococcus_troglodytae;_AP014612.1	012%
Streptococcus_intermedius_B196;_CP003857.1	000%
Streptococcus_intermedius_JTH08;_AP010969.1	000%
Streptococcus_intermedius;_AP014880.1	000%