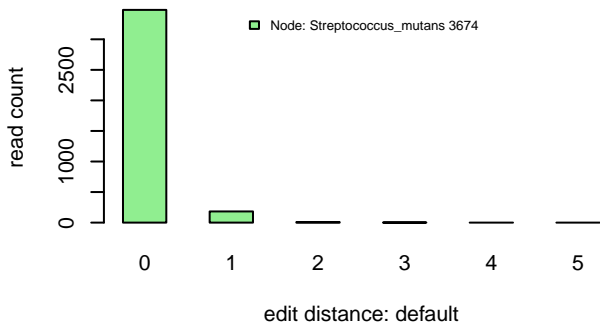
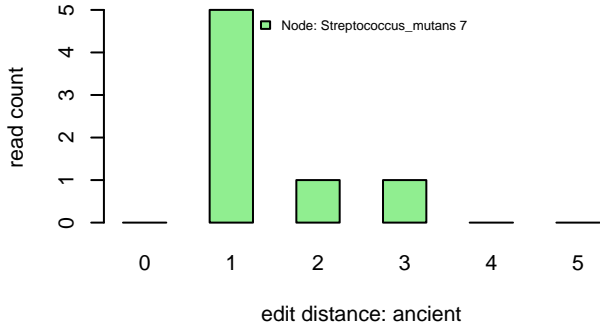


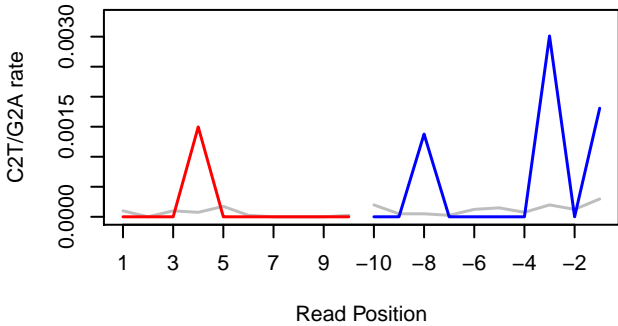
AT+Mock-1:50-3.rma6



AT+Mock-1:50-3.rma6



Damage plot for Streptococcus\_mutans node



Node	Streptococcus_mutans
Top Reference	Streptococcus_mutans_UA159
assigned reads	8475
all reads	8451
nonDup	8451
readDis	0.581
nonStacked	3661
destacking?	On
C>T_1	0
G>A_-1	0.0018
mean length (sd)	75 (2.223)

Streptococcus_mutans_UA159;_AE014133.2	100%
Streptococcus_mutans_UA159-FR;_CP007016.1	100%
Streptococcus_mutans_GS-5;_CP003686.1	075%
Streptococcus_mutans;_CP013237.1	074%
Streptococcus_mutans_NN2025;_AP010655.1	067%
Streptococcus_mutans_LJ23;_AP012336.1	066%
Streptococcus_troglodytae;_AP014612.1	011%
Streptococcus_intermedius_B196;_CP003857.1	000%
Streptococcus_intermedius;_AP014880.1	000%
Paenibacillus_yonginensis;_CP014167.1	000%