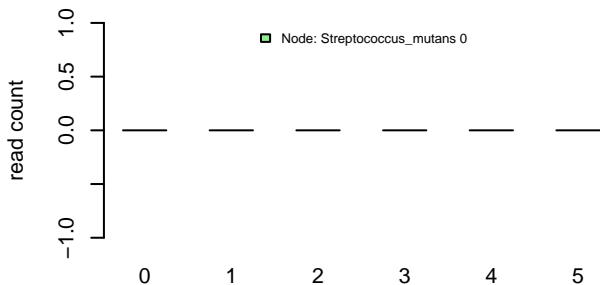
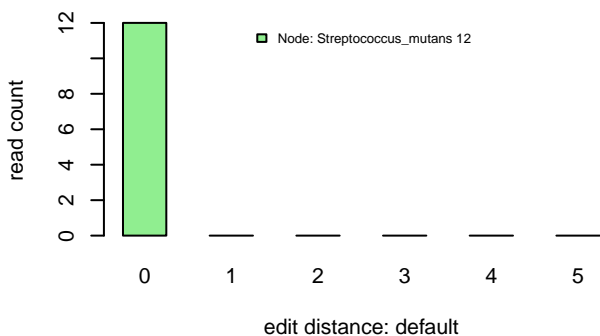
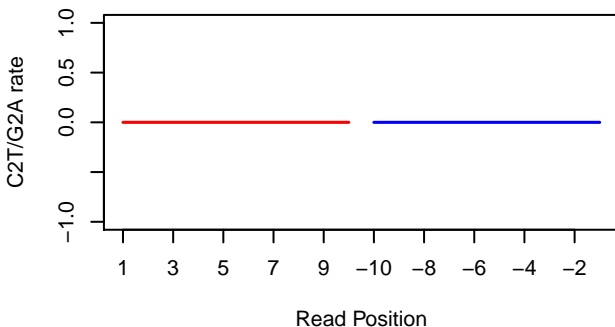


XXX001.A17841.SG1.1rc\_S0\_L006\_R1\_001.fastq.gz.rXXXX001.A17841.SG1.1rc\_S0\_L006\_R1\_001.fastq.gz.r



### Damage plot for Streptococcus\_mutans node



Node	Streptococcus_mutans
Top Reference	Streptococcus_mutans_UA159-FR
assigned reads	12
reads ME	12
all reads	12
nonDup	12
readDis	1
nonStacked	12
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (2.275)

Streptococcus_mutans_UA159-FR;_CP007016.1	100%
Streptococcus_mutans_UA159;_AE014133.2	100%
Streptococcus_mutans_GS-5;_CP003686.1	083%
Streptococcus_mutans_NN2025;_AP010655.1	083%
Streptococcus_mutans;_CP013237.1	083%
Streptococcus_mutans_LJ23;_AP012336.1	058%
Streptococcus_troglodytae;_AP014612.1	008%
	NA%
	NA%
	NA%