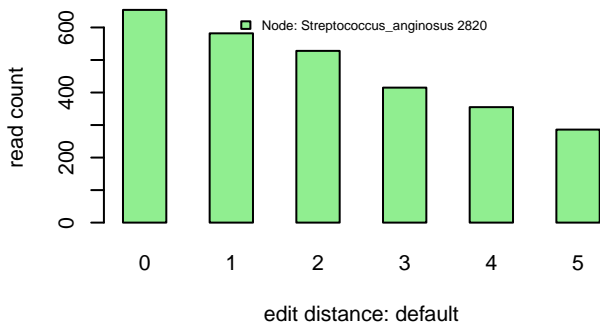
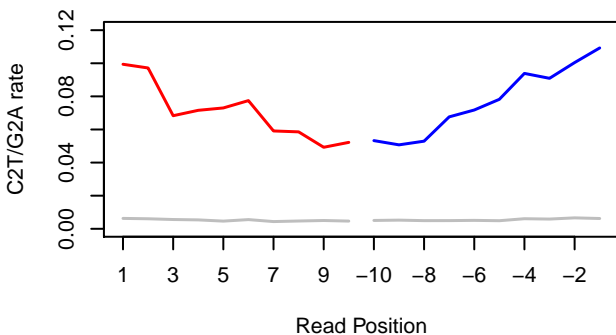


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for Streptococcus\_anginosus node



Node	Streptococcus_anginosus
Top Reference	Streptococcus_anginosus
assigned reads	3696
all reads	2432
nonDup	2379
readDis	0.034
nonStacked	51
destacking?	Off
C>T_1	0.0994
G>A_-1	0.1093
mean length (sd)	67 (24.102)

Streptococcus_anginosus;_CP007573.1 100%
Streptococcus_anginosus_subsp._whileyi_MAS624;_AP013072.1 100%
Streptococcus_anginosus_C238;_CP003861.1 100%
Streptococcus_anginosus_C1051;_CP003860.1 100%
Streptococcus_intermedius;_AP014880.1 052%
Streptococcus_mutans_GS-5;_CP003686.1 011%
streptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1 009%
Pasteurella_multocida_OH1905;_CP004392.1 003%
Clostridioides_difficile_BI9;_FN668944.1 003%
Streptococcus_agalactiae;_CP011329.1 003%