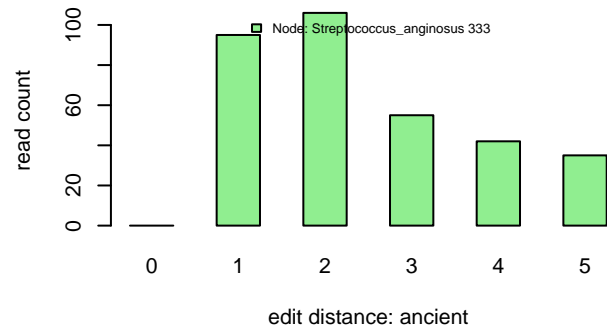
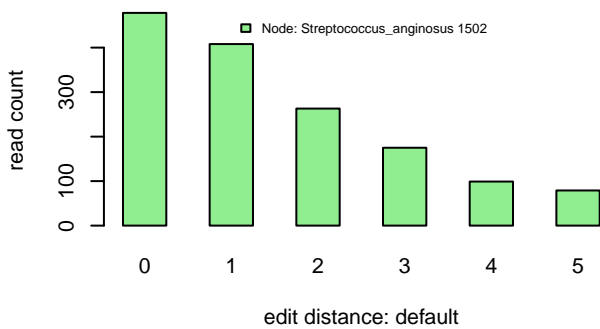
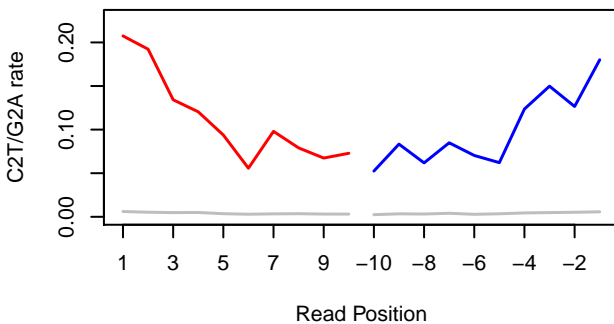


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for Streptococcus\_anginosus node



Node	Streptococcus_anginosus
Top Reference	Streptococcus_anginosus_C238
assigned reads	1662
all reads	804
nonDup	785
readDis	0.703
nonStacked	518
destacking?	Off
C>T_1	0.2074
G>A_1	0.1801
mean length (sd)	50 (16.423)

Streptococcus_anginosus_C238;_CP003861.1 100%
Streptococcus_anginosus_subsp._whileyi_MAS624;_AP013072.1 100%
Streptococcus_anginosus_C1051;_CP003860.1 100%
Streptococcus_anginosus;_CP012719.1 100%
Streptococcus_anginosus;_CP012805.1 098%
Streptococcus_anginosus;_CP007573.1 098%
Streptococcus_anginosus_SK52;=_DSM_20563;_AB643814.1 002%
Streptococcus_anginosus;_KY315465.1 001%
Streptococcus_anginosus;_KC617870.1 000%
Streptococcus_mutans_GS-5;_CP003686.1 000%