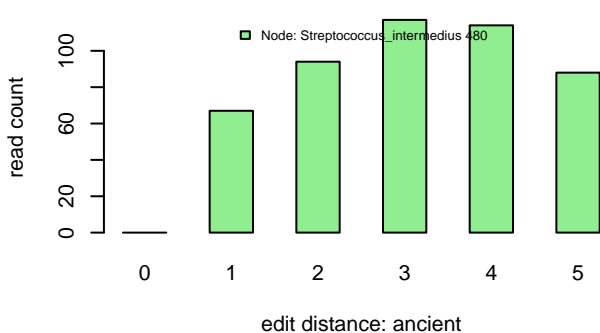
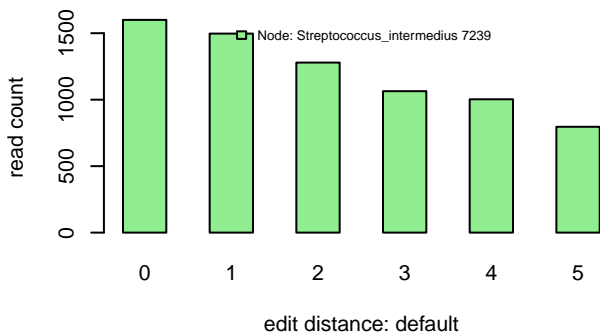
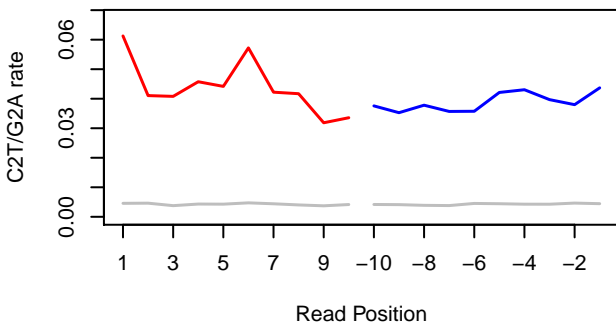


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius_C270
assigned reads	16361
all reads	4483
nonDup	4417
readDis	0.24
nonStacked	746
destacking?	Off
C>T_1	0.0613
G>A_-1	0.0437
mean length (sd)	88 (26.267)

Streptococcus_intermedius_C270;_CP003858.1	100%
Streptococcus_intermedius;_CP020433.1	100%
Streptococcus_intermedius;_CP012718.1	100%
Streptococcus_intermedius_JTH08;_AP010969.1	100%
Streptococcus_intermedius;_AP014880.1	100%
Streptococcus_intermedius_B196;_CP003857.1	100%
Streptococcus_sanguinis_SK36;_CP000387.1	035%
Streptococcus_gordonii;_CP020450.1	005%
Streptococcus_cristatus_AS_1.3089;_CP004409.1	005%
Streptococcus_intermedius;_AB608790.1	005%