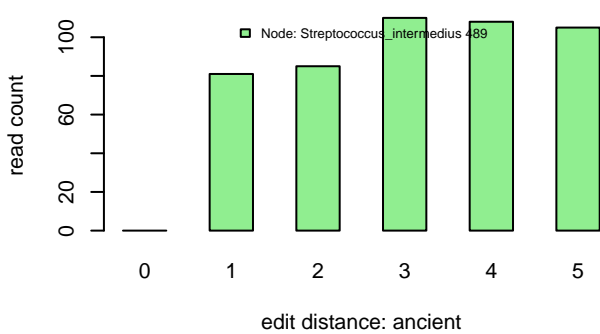
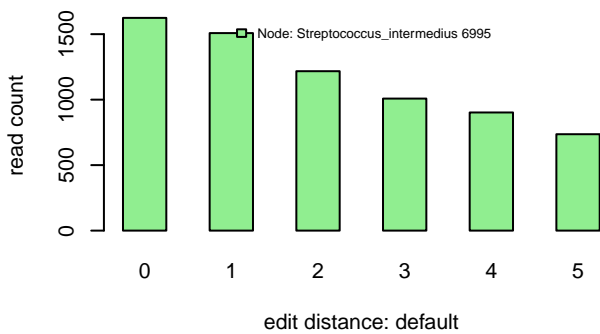
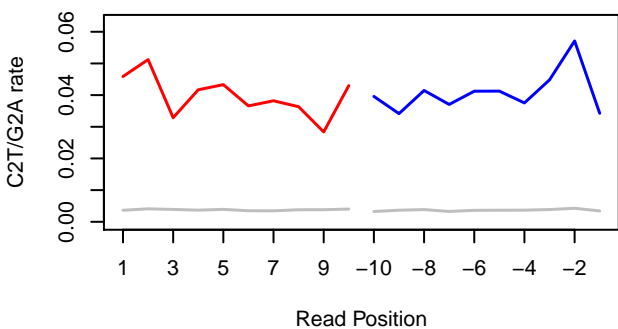


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	17312
all reads	5854
nonDup	5722
readDis	0.146
nonStacked	515
destacking?	Off
C>T_1	0.0459
G>A_-1	0.0343
mean length (sd)	103 (39.982)

Streptococcus_intermedius;_CP020433.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_sanguinis_SK36;_CP000387.1 091%
Streptococcus_gordonii;_CP020450.1 014%
Streptococcus_cristatus_AS_1.3089;_CP004409.1 009%
Streptococcus_gordonii_str._Challis_substr._CH1;_CP000725.1 009%
Streptococcus_anginosus_C238;_CP003861.1 004%
Streptococcus_gordonii;_CP012648.1 004%