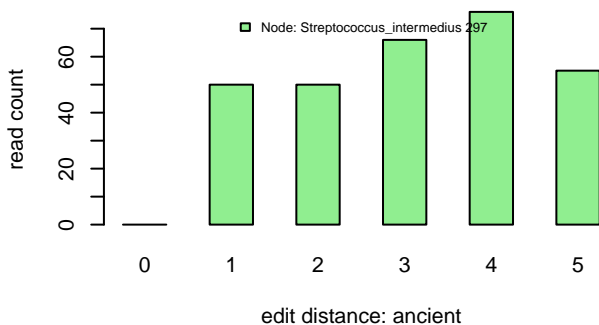
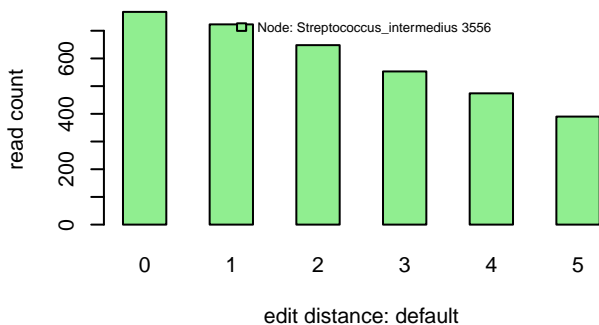
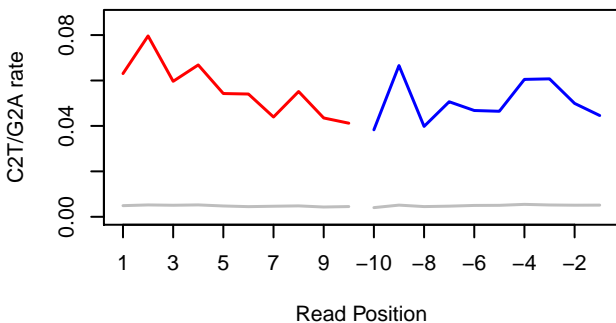


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius_C270
assigned reads	5126
all reads	3015
nonDup	2940
readDis	0.186
nonStacked	379
destacking?	Off
C>T_1	0.0631
G>A_-1	0.0446
mean length (sd)	83 (22.69)

Streptococcus_intermedius_C270;_CP003858.1	100%
Streptococcus_intermedius;_CP020433.1	100%
Streptococcus_intermedius;_CP012718.1	100%
Streptococcus_intermedius;_AP014880.1	100%
Streptococcus_intermedius_JTH08;_AP010969.1	100%
Streptococcus_intermedius_B196;_CP003857.1	100%
Streptococcus_sanguinis_SK36;_CP000387.1	016%
Streptococcus_anginosus_C1051;_CP003860.1	010%
Streptococcus_gordonii;_CP020450.1	006%
Streptococcus_gordonii_str._Challis_substr._CH1;_CP000725.1	006%