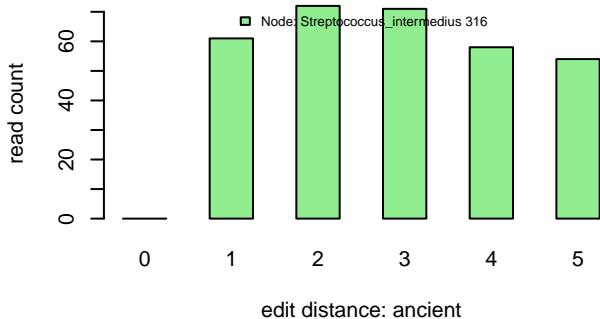
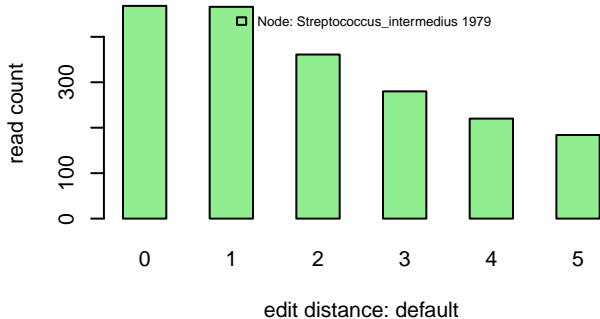
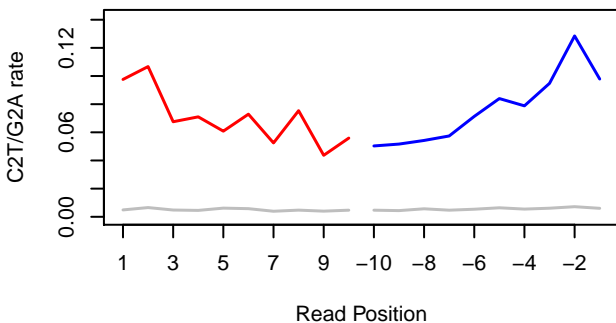


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	2566
all reads	1675
nonDup	1637
readDis	0.308
nonStacked	401
destacking?	Off
C>T_1	0.0976
G>A_-1	0.0979
mean length (sd)	62 (21.528)

Streptococcus_intermedius;_CP012718.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_sanguinis_SK36;_CP000387.1 020%
Streptococcus_cristatus_AS_1.3089;_CP004409.1 014%
Streptococcus_anginosus_C1051;_CP003860.1 006%
Streptococcus_gordonii;_CP020450.1 003%
Streptococcus_gordonii;_CP017295.1 003%
Streptococcus_gordonii_str._Challis_substr._CH1;_CP000725.1 003%