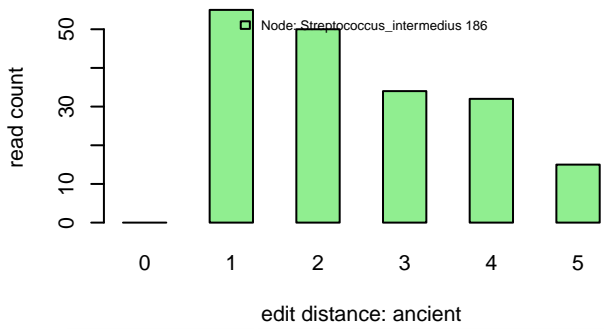
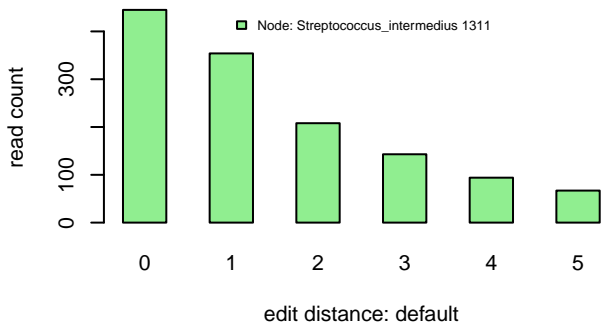
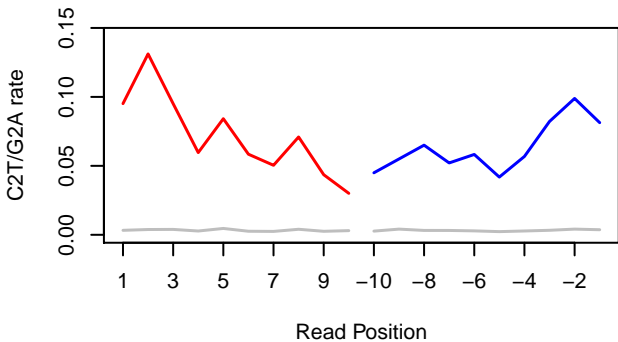


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius_C270
assigned reads	1467
all reads	862
nonDup	839
readDis	0.581
nonStacked	423
destacking?	Off
C>T_1	0.0951
G>A_-1	0.0813
mean length (sd)	71 (26.32)

Streptococcus_intermedius_C270;_CP003858.1	100%
Streptococcus_intermedius_B196;_CP003857.1	100%
Streptococcus_intermedius;_CP012718.1	100%
Streptococcus_intermedius;_CP020433.1	100%
Streptococcus_intermedius_JTH08;_AP010969.1	100%
Streptococcus_intermedius;_AP014880.1	100%
Streptococcus_anginosus_C238;_CP003861.1	001%
Streptococcus_anginosus_C1051;_CP003860.1	000%
Streptococcus_anginosus_subsp._whileyi_MAS624;_AP013072.1	000%
Streptococcus_intermedius;_KY315473.1	000%