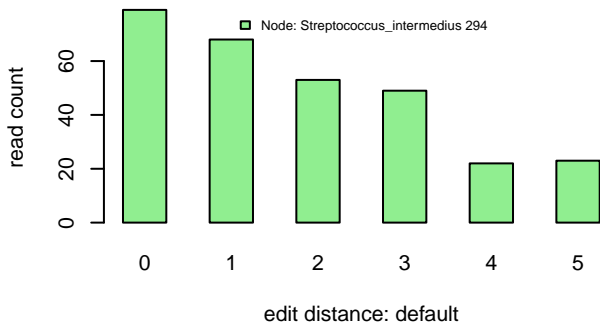
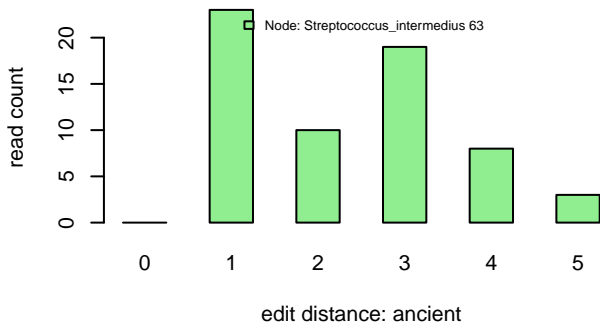


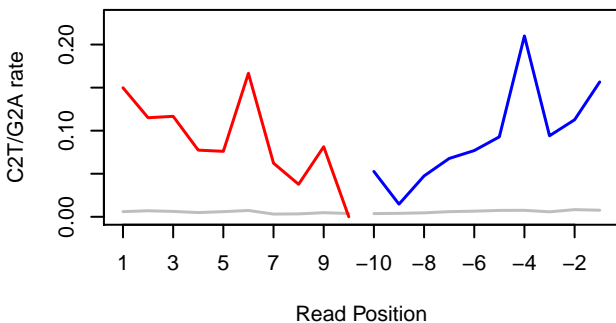
101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	403
all reads	217
nonDup	215
readDis	0.852
nonStacked	174
destacking?	Off
C>T_1	0.1498
G>A_-1	0.1566
mean length (sd)	56 (20.762)

Streptococcus_intermedius;_CP020433.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_angermanii;_KY315478.1 001%
Streptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 001%
Streptococcus_trogolodytae;_AP014612.1 000%
Enterococcus;_CP021874.1 000%
Streptococcus_pneumoniae_Hungary19A-6;_CP000936.1 000%
Streptococcus_haltotolerans;_CP014835.1 000%