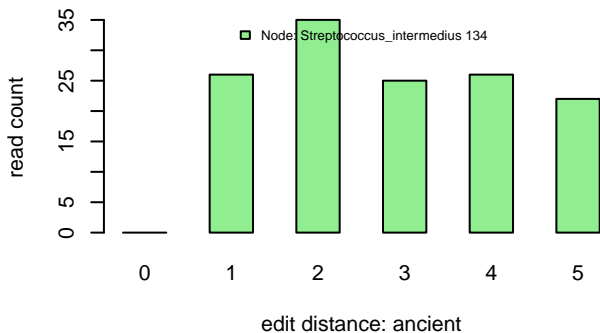
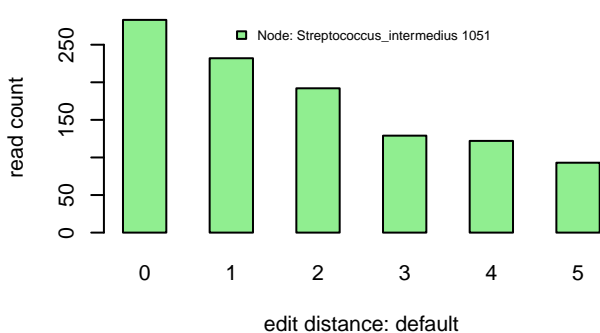
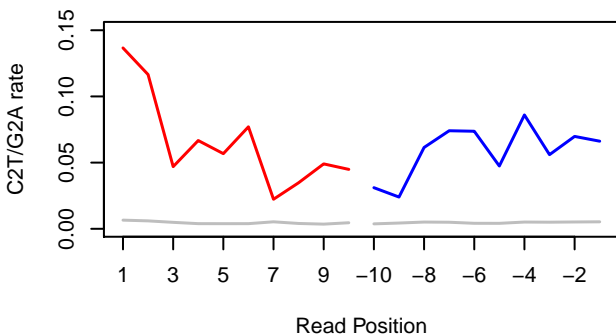


101_S0_L000_R1_000.fastq.merged.prefixed.hg19un101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	1321
all reads	755
nonDup	742
readDis	0.604
nonStacked	389
destacking?	Off
C>T_1	0.1366
G>A_-1	0.0662
mean length (sd)	69 (24.405)

Streptococcus_intermedius;_CP012718.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_sanguinis_SK36;_CP000387.1 005%
Streptococcus_cristatus_AS_1.3089;_CP004409.1 002%
Streptococcus_gordonii;_CP017295.1 001%
Streptococcus_anginosus_subsp._whileyi_MAS624;_AP013072.1 000%
Streptococcus_anginosus;_CP012719.1 000%
Streptococcus_anginosus_C238;_CP003861.1 000%