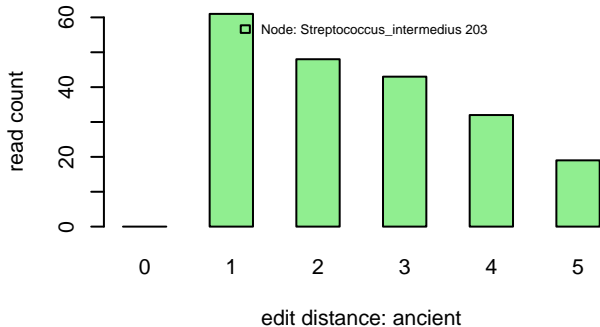
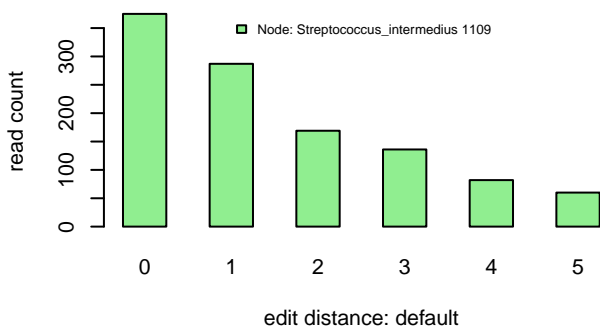
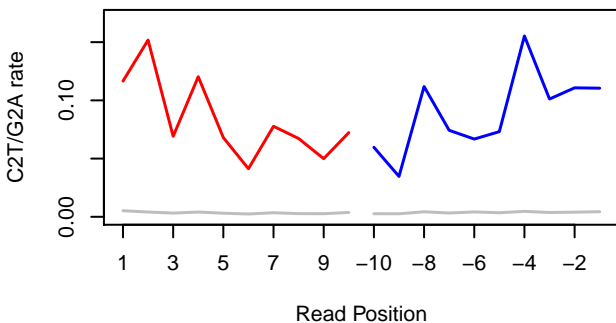


101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un)101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un)



### Damage plot for Streptococcus\_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	1226
all reads	762
nonDup	749
readDis	0.745
nonStacked	524
destacking?	Off
C>T_1	0.1166
G>A_-1	0.1105
mean length (sd)	56 (20.054)

Streptococcus_intermedius;_CP012718.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_angermanus_C238;_CP003861.1 001%
Streptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 000%
Streptococcus_sanguinis_SK36;_CP000387.1 000%
Streptococcus_angermanus_subsp._whitleyi_MAS624;_AP013072.1 000%
Streptococcus_angermanus;_CP012719.1 000%
Streptococcus_angermanus;_KY315478.1 000%