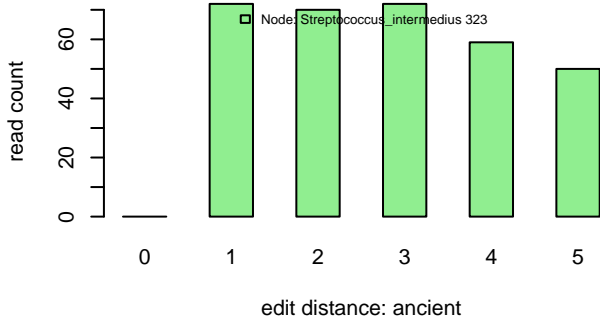
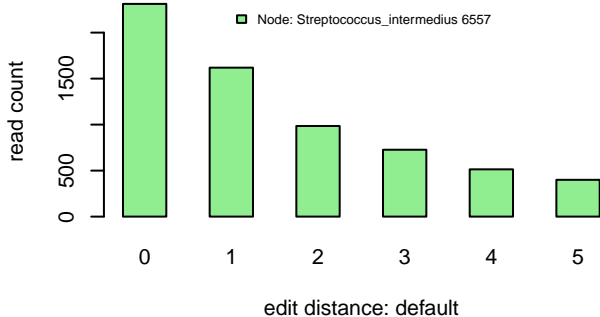
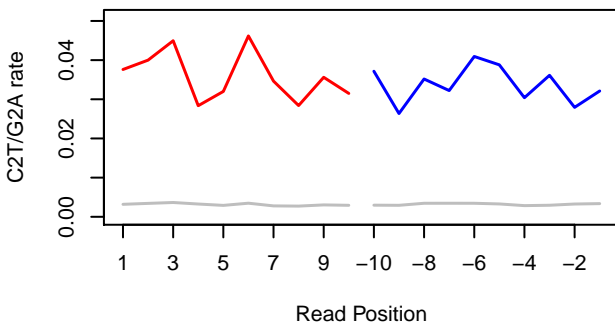


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius_C270
assigned reads	8149
all reads	5373
nonDup	5296
readDis	0.253
nonStacked	1019
destacking?	Off
C>T_1	0.0376
G>A_-1	0.0321
mean length (sd)	83 (21.876)

Streptococcus_intermedius_C270;_CP003858.1	100%
Streptococcus_intermedius_JTH08;_AP010969.1	100%
Streptococcus_intermedius;_CP020433.1	100%
Streptococcus_intermedius;_CP012718.1	100%
Streptococcus_intermedius;_AP014880.1	100%
Streptococcus_intermedius_B196;_CP003857.1	100%
Streptococcus_sanguinis_SK36;_CP000387.1	007%
Streptococcus_intermedius;_AB608790.1	003%
Streptococcus_anginosus_C238;_CP003861.1	002%
Streptococcus_intermedius;_KY315471.1	001%