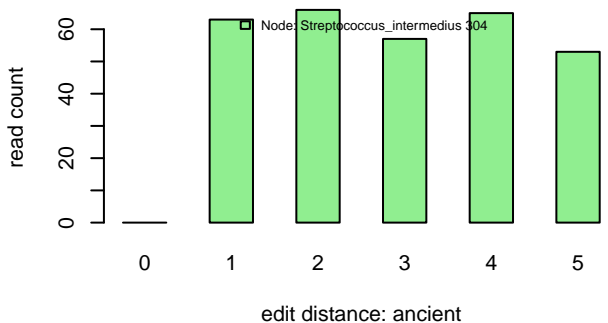
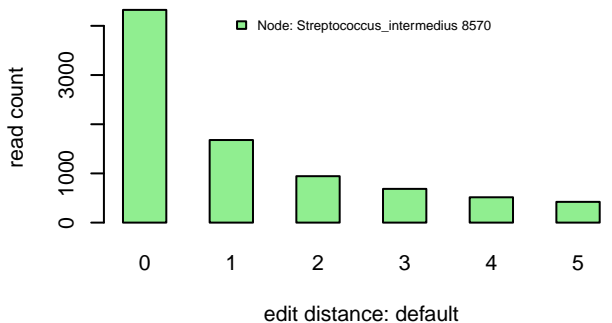
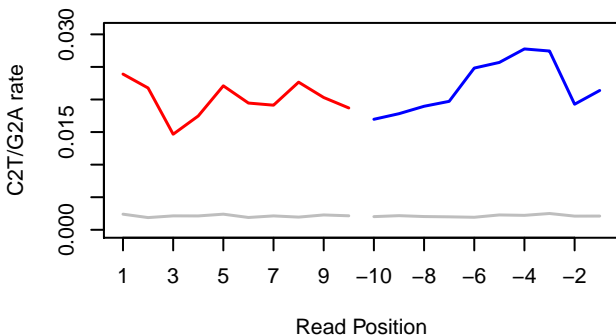


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius_JTH08
assigned reads	28643
all reads	5826
nonDup	5734
readDis	0.477
nonStacked	2082
destacking?	Off
C>T_1	0.0239
G>A_-1	0.0214
mean length (sd)	89 (27.915)

Streptococcus_intermedius_JTH08;_AP010969.1 100%

Streptococcus_intermedius;_AP014880.1 100%

Streptococcus_intermedius;_CP020433.1 100%

Streptococcus_intermedius;_CP012718.1 100%

Streptococcus_intermedius_C270;_CP003858.1 100%

Streptococcus_intermedius_B196;_CP003857.1 100%

Streptococcus_sanguinis_SK36;_CP000387.1 008%

Streptococcus_intermedius;_AB608790.1 002%

Streptococcus_anginosus_C1051;_CP003860.1 002%

Streptococcus_gordonii;_CP020450.1 001%