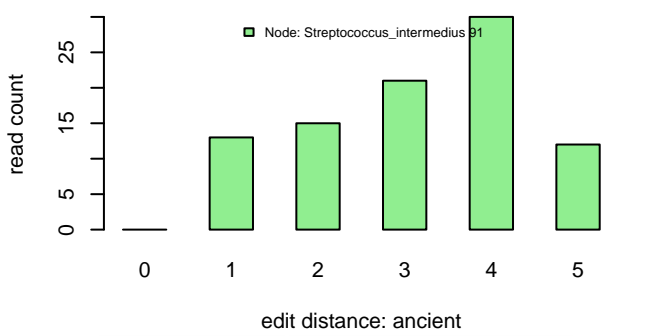
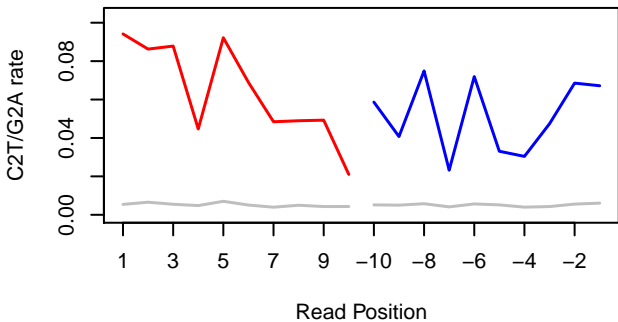


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus\_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	1081
all reads	623
nonDup	609
readDis	0.492
nonStacked	231
destacking?	Off
C>T_1	0.0942
G>A_-1	0.0672
mean length (sd)	64 (22.453)

Streptococcus_intermedius;_CP020433.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_sanguinis_SK36;_CP000387.1 013%
Streptococcus_cristatus_AS_1.3089;_CP004409.1 007%
Streptococcus_anginosus_C1051;_CP003860.1 004%
Streptococcus_gordonii;_CP020450.1 003%
Streptococcus_gordonii;_CP017295.1 002%
Streptococcus_gordonii_str._Challis_substr._CH1;_CP000725.1 002%