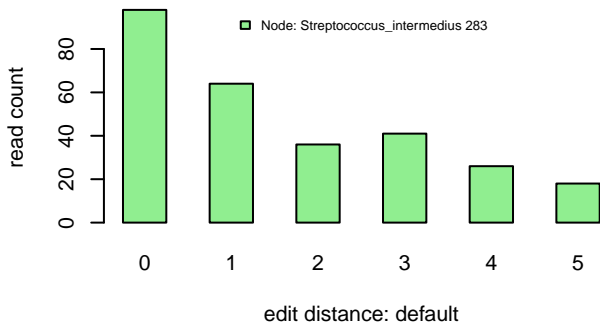
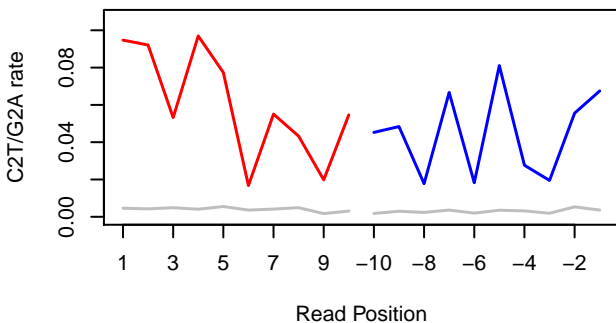


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius_JTH08
assigned reads	334
all reads	183
nonDup	180
readDis	0.759
nonStacked	131
destacking?	Off
C>T_1	0.0947
G>A_-1	0.0675
mean length (sd)	55 (19.105)

Streptococcus_intermedius_JTH08;_AP010969.1	100%
Streptococcus_intermedius_C270;_CP003858.1	100%
Streptococcus_intermedius_B196;_CP003857.1	100%
Streptococcus_intermedius;_CP012718.1	100%
Streptococcus_intermedius;_CP020433.1	100%
Streptococcus_intermedius;_AP014880.1	100%
Streptococcus_intermedius;_KY315470.1	003%
Streptococcus_intermedius;_KY315460.1	003%
Streptococcus_intermedius;_KY315461.1	003%
Streptococcus_intermedius;_KY315471.1	003%