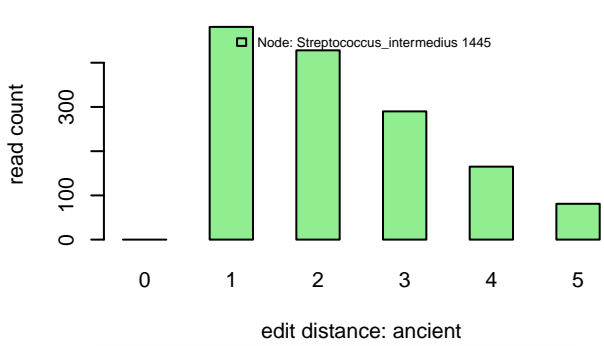
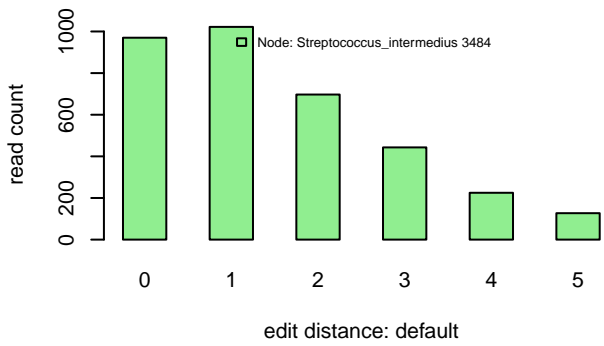
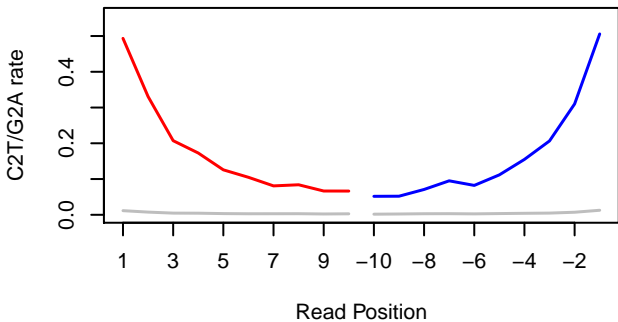


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_intermedius node



Node	Streptococcus_intermedius
Top Reference	Streptococcus_intermedius
assigned reads	3686
all reads	2590
nonDup	2547
readDis	0.552
nonStacked	1032
destacking?	Off
C>T_1	0.4935
G>A_-1	0.5058
mean length (sd)	50 (16.057)

Streptococcus_intermedius;_CP020433.1 100%
Streptococcus_intermedius_JTH08;_AP010969.1 100%
Streptococcus_intermedius_B196;_CP003857.1 100%
Streptococcus_intermedius_C270;_CP003858.1 100%
Streptococcus_angermanii_C238;_CP003861.1 003%
Streptococcus_angermanii_C1051;_CP003860.1 001%
Streptococcus_angermanii;_CP012719.1 001%
Streptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 001%
Streptococcus_angermanii_subsp._whitleyi_MAS624;_AP013072.1 001%
Streptococcus_constellatus_subsp._pharyngis_C232;_CP003800.1 000%