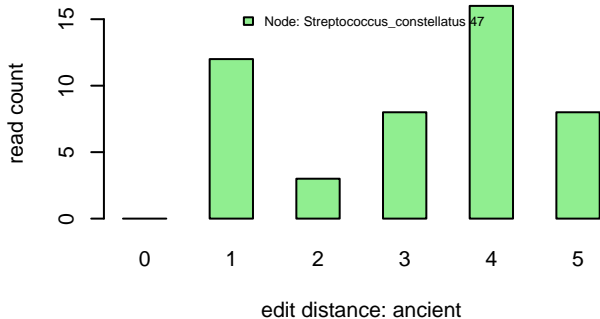
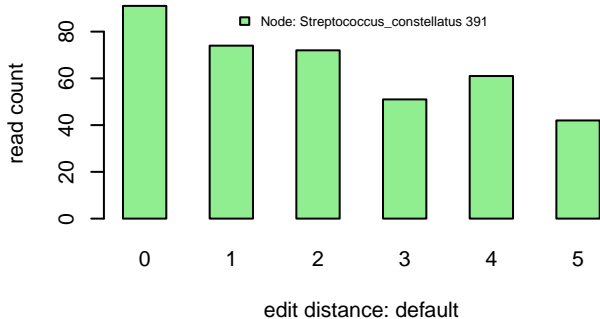
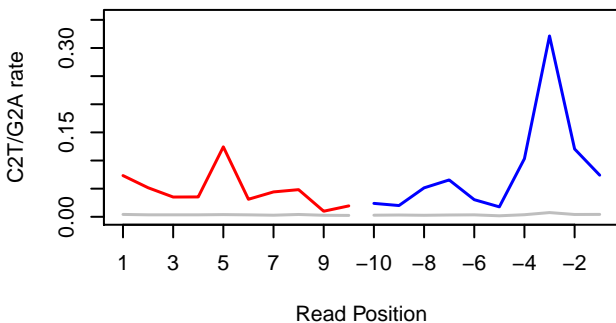


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_constellatus node



Node	Streptococcus_constellatus
Top Reference	Streptococcus_constellatus
assigned reads	539
all reads	425
nonDup	413
readDis	0.139
nonStacked	34
destacking?	Off
C>T_1	0.0733
G>A_-1	0.074
mean length (sd)	68 (23.203)

Streptococcus_constellatus;_KY315462.1 100%
reptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 100%
reptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1 100%
reptococcus_constellatus_subsp._pharyngis_C232;_CP003800.1 100%
Streptococcus_constellatus_subsp._constellatus;_KY315463.1 058%
Streptococcus_constellatus_subsp._constellatus;_KY315464.1 058%
Streptococcus_intermedius;_KY315447.1 032%
Streptococcus_constellatus_subsp._constellatus;_AF385683.1 029%
Streptococcus_angeris_group;_JN787547.1 020%
Streptococcus_constellatus_subsp._constellatus;_EU156798.1 020%