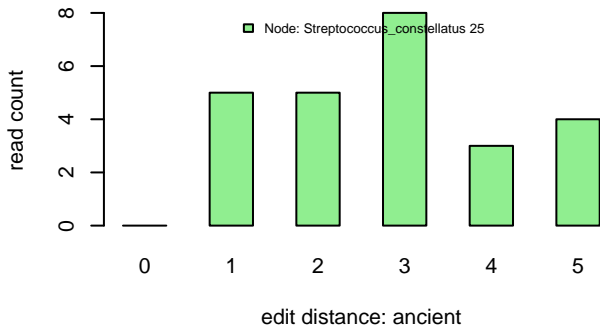
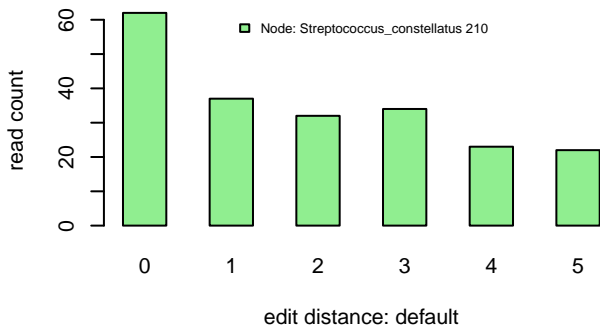
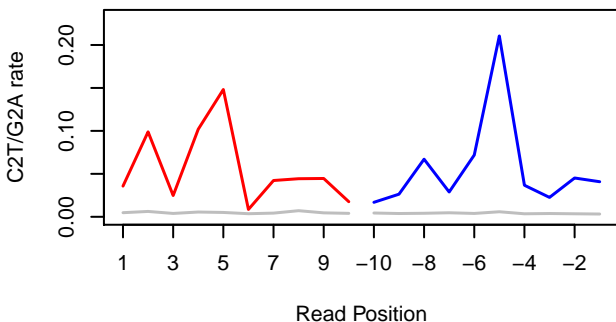


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Streptococcus_constellatus node



Node	Streptococcus_constellatus
Top Reference	Streptococcus_constellatus
assigned reads	434
all reads	304
nonDup	301
readDis	0.203
nonStacked	34
destacking?	Off
C>T_1	0.0357
G>A_-1	0.0408
mean length (sd)	83 (22.823)

Streptococcus_constellatus;_KY315462.1	100%
reptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1	100%
treptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1	100%
treptococcus_constellatus_subsp._pharyngis_C232;_CP003800.1	100%
Streptococcus_constellatus_subsp._constellatus;_AB839362.1	055%
Streptococcus_constellatus_subsp._constellatus;_KY315463.1	032%
Streptococcus_constellatus_subsp._constellatus;_KY315464.1	032%
Streptococcus_constellatus_subsp._pharyngis;_AB839365.1	032%
eptococcus_constellatus_subsp._viborgensis_SK1359;_AB860419.1	032%
Streptococcus_constellatus_subsp._pharyngis;_KY315452.1	023%