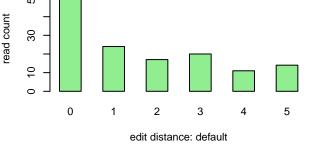
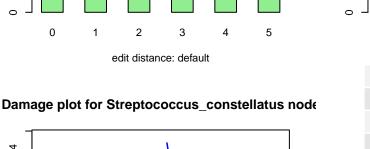
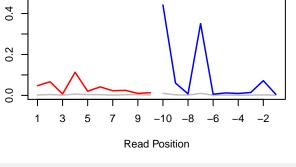


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur







C2T/G2A rate

treptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1 061% reptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 059%

Streptococcus constellatus subsp. constellatus; KY315463.1 035%

Streptococcus_constellatus;_KY315462.1 100% treptococcus_constellatus_subsp._pharyngis_C232;_CP003800.1 061%

Streptococcus_constellatus_subsp._constellatus;_KY315464.1 035%

Streptococcus_intermedius;_KY315447.1 024%

Streptococcus_constellatus_subsp._pharyngis;_KY315452.1 015%

Streptococcus_constellatus_subsp._pharyngis;_KY315453.1 015%

Streptococcus_constellatus_subsp._pharyngis;_KY315450.1 015%

0 2 3

edit distance: ancient Node Streptococcus_constellatus Top Reference Streptococcus_constellatus assigned reads 207 all reads 144 nonDup 141 readDis 0.535 nonStacked 57 Off destacking? C>T_1 0.0473 G>A_-1 0.005

68 (25.377)

mean length (sd)