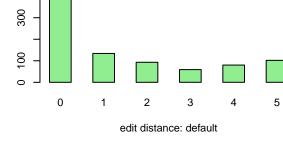


 \sim

0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un

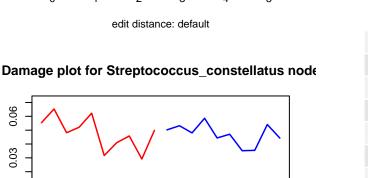


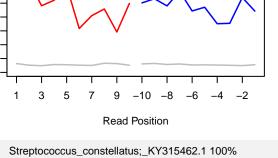
90.0

0.03

0.00

C2T/G2A rate





Streptococcus_constellatus_subsp._constellatus;_AF385683.1 016% reptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 012%

treptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1 012%

treptococcus_constellatus_subsp._pharyngis_C232;_CP003800.1 011%

Streptococcus_constellatus_subsp._constellatus;_KY315463.1 005%

Streptococcus_constellatus_subsp._constellatus;_KY315464.1 005%

Streptococcus_constellatus_subsp._pharyngis;_NR_121935.1 005%

Streptococcus_constellatus_subsp._constellatus;_AM286938.1 004%

Streptococcus_merionis;_LT906439.1 002%

0 2 3 edit distance: ancient Node Streptococcus_constellatus Top Reference Streptococcus_constellatus assigned reads 1604 all reads 1536 nonDup 1517 readDis 0.098 nonStacked 72 Off destacking? C>T_1 0.0554 G>A_-1 0.0443

84 (21.713)

mean length (sd)