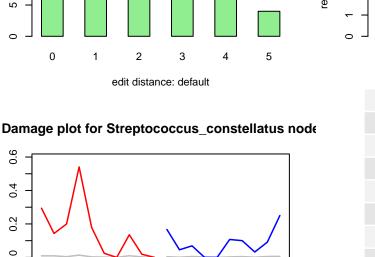
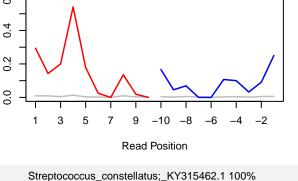


0

S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu





C2T/G2A rate

Streptococcus_constellatus_subsp._constellatus;_AF385683.1 100%

reptococcus_constellatus_subsp._pharyngis_C1050;_CP003859.1 052% treptococcus_constellatus_subsp._pharyngis_C232;_CP003800.1 042% Streptococcus_constellatus_subsp._pharyngis;_NR_121935.1 036%

treptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1 057%

Node Streptococcus_constellatus Top Reference Streptococcus_constellatus assigned reads 87 all reads 34 nonDup 32 readDis 0.568 nonStacked 19 Off destacking? C>T_1 0.2941 G>A_-1 0.25 mean length (sd) 54 (19.172)

2

edit distance: ancient

3

Streptococcus_merionis;_LT906439.1 021%

Streptococcus_constellatus_subsp._constellatus;_HM751893.1 010%

Streptococcus_constellatus_subsp._constellatus;_JN787245.1 005%

Streptococcus_constellatus_subsp._constellatus;_KP215270.1 005%