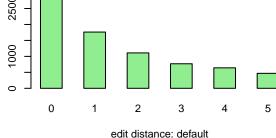
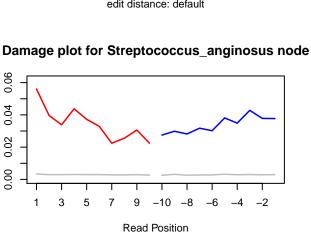


20

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





C2T/G2A rate



Streptococcus_anginosus;_CP007573.1 100%

Streptococcus_anginosus_subsp._whileyi_MAS624;_AP013072.1 100%

Streptococcus_anginosus_C238;_CP003861.1 100%

Streptococcus_anginosus_SK52_=_DSM_20563;_AB643814.1 002% Streptococcus_anginosus_subsp._whileyi;_KY315457.1 001%

Streptococcus_anginosus_SK52_=_DSM_20563;_AF535183.1 000%

Streptococcus_anginosus_subsp._whileyi;_JN787718.1 000% Streptococcus_anginosus_subsp._whileyi;_KY274990.1 000%

Streptococcus_anginosus_subsp._whileyi;_JX993493.1 000%

0 2 3 edit distance: ancient Node Streptococcus_anginosus Top Reference Streptococcus_anginosus assigned reads 29541 all reads 4309 nonDup 4242 readDis 0.303 nonStacked 1083 Off destacking?

0.0561

0.0377

89 (27.425)

C>T_1

G>A_-1

mean length (sd)