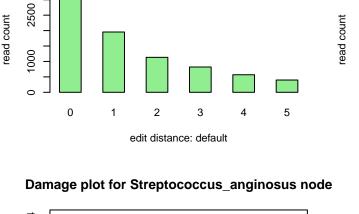
■ Node: Streptococcus_anginosus 8321 Node: Streptococcus_anginosus 357 80

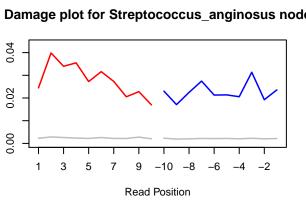
9 40

20

0

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





C2T/G2A rate

Streptococcus_anginosus;_CP012719.1 100% Streptococcus_anginosus_C1051;_CP003860.1 100%

Streptococcus_anginosus_subsp._whileyi_MAS624;_AP013072.1 100%

Streptococcus_anginosus_C238;_CP003861.1 100%

Streptococcus_anginosus_SK52_=_DSM_20563;_AB643814.1 002%

Streptococcus_anginosus_SK52_=_DSM_20563;_KY315441.1 000% Streptococcus_sanguinis_SK36;_CP000387.1 000%

Streptococcus_intermedius_C270;_CP003858.1 000%

Streptococcus_intermedius_B196;_CP003857.1 000%

Streptococcus_anginosus_SK52_=_DSM_20563;_AF535183.1 000%

edit distance: ancient Node Streptococcus_anginosus Top Reference Streptococcus_anginosus assigned reads 18878 all reads 5548 nonDup 5418 readDis 0.529 nonStacked 1804 Off destacking? C>T_1 0.0245 G>A_-1 0.0236 mean length (sd) 113 (40.979)

2

3