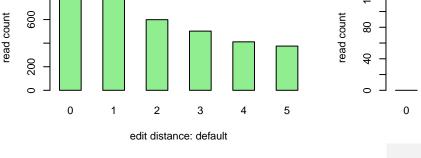
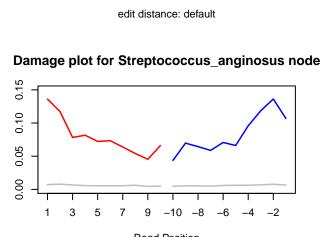
120 Node: Streptococcus_anginosus 574 Node: Streptococcus_anginosus 3602

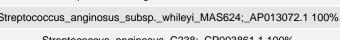
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur





C2T/G2A rate

Read Position Streptococcus_anginosus;_CP012719.1 100%



Streptococcus_anginosus_C238;_CP003861.1 100% Streptococcus_anginosus_C1051;_CP003860.1 100%

Streptococcus_cristatus_AS_1.3089;_CP004409.1 002% Streptococcus_anginosus_subsp._whileyi;_KY315457.1 001%

Streptococcus_sanguinis_SK36;_CP000387.1 002%

Streptococcus_anginosus_SK52_=_DSM_20563;_AB643814.1 001%

2 3 edit distance: ancient Node Streptococcus_anginosus Top Reference Streptococcus_anginosus assigned reads 4841 all reads 2111 nonDup 2078 readDis 0.421 nonStacked 682 Off destacking?

0.1361

0.1069

65 (23.196)

C>T_1

G>A_-1

mean length (sd)

Streptococcus_intermedius;_AP014880.1 001%

treptococcus_constellatus_subsp._pharyngis_C818;_CP003840.1 001%