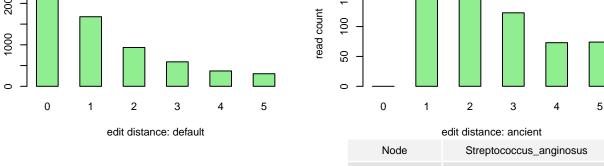
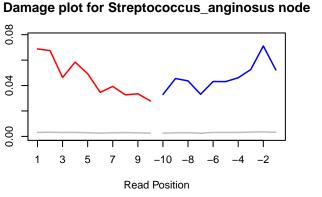
■ Node: Streptococcus_anginosus 6442 Streptococcus_anginosus 621 150 2000

0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un

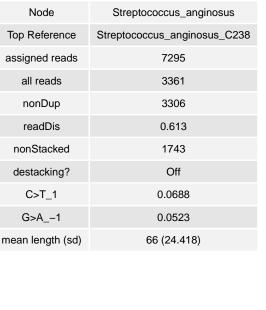




C2T/G2A rate

Streptoco

Streptococcus_anginosus_C238;_CP003861.1 100%
occus anginosus subsp. whileyi MAS624; AP013072.1 100%



Streptococcus_anginosus;_CP012805.1 100%

Streptococcus_anginosus_C1051;_CP003860.1 100%

Streptococcus_anginosus;_CP012719.1 100%

Streptococcus_anginosus;_CP007573.1 100%

Streptococcus_anginosus;_KY315465.1 004%

Streptococcus_anginosus_SK52_=_DSM_20563;_AB643814.1 002%

Streptococcus_anginosus;_KC617870.1 000%

Streptococcus_anginosus_subsp._whileyi;_KY315457.1 000%