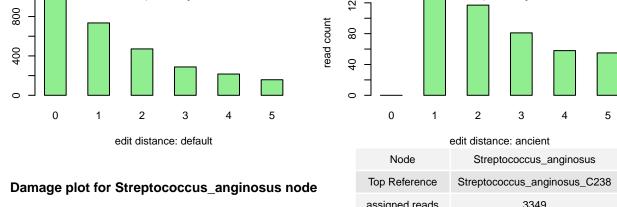
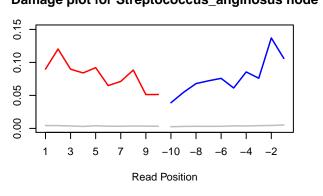
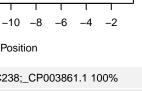
## Node: Streptococcus\_anginosus 449 Node: Streptococcus\_anginosus 2948 120 800

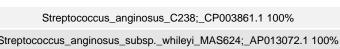
0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un)101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un





C2T/G2A rate





Streptococcus\_anginosus;\_CP012719.1 100% Streptococcus\_anginosus;\_CP007573.1 100%

Streptococcus\_anginosus\_C1051;\_CP003860.1 100%

Streptococcus\_anginosus;\_CP012805.1 100% Streptococcus\_anginosus;\_KY315465.1 012%

Streptococcus\_anginosus\_SK52\_=\_DSM\_20563;\_AB643814.1 002%

Streptococcus\_anginosus;\_KY315455.1 000%

Streptococcus\_anginosus;\_KY315458.1 000%

assigned reads 3349 all reads 1366 nonDup 1337 readDis 0.589 nonStacked 739 destacking? Off C>T\_1 0.0898 G>A\_-1 0.1059 mean length (sd) 58 (20.356)