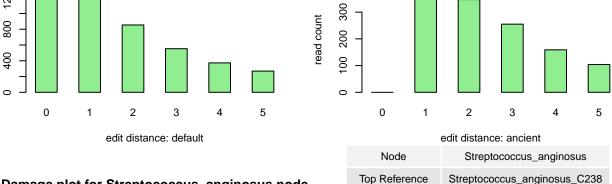
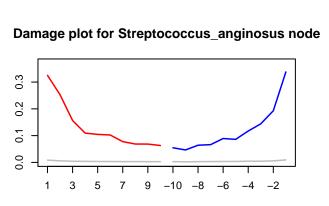
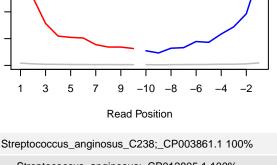
## 1200 Node: Streptococcus\_anginosus 1263 Node: Streptococcus\_anginosus 4679 read count

0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur





C2T/G2A rate



Streptococcus\_anginosus;\_CP012805.1 100%

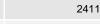
Streptococcus\_anginosus\_C1051;\_CP003860.1 100% Streptococcus\_anginosus;\_CP007573.1 100%

Streptococcus\_anginosus\_subsp.\_whileyi\_MAS624;\_AP013072.1 100%

assigned reads

all reads

nonDup



Streptococcus\_anginosus\_C238

5262

2361

readDis 0.545 nonStacked 1198

destacking? Off C>T\_1 0.325

G>A\_-1 0.3377

mean length (sd) 65 (23.587)

Streptococcus\_anginosus;\_CP012719.1 100% Streptococcus\_anginosus;\_KY315465.1 002%

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

Streptococcus\_intermedius\_C270;\_CP003858.1 000%

Streptococcus\_anginosus;\_KC617870.1 000%