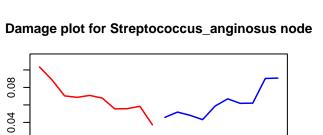


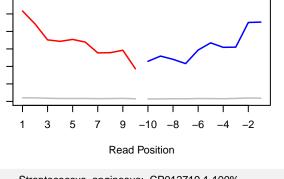
S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu

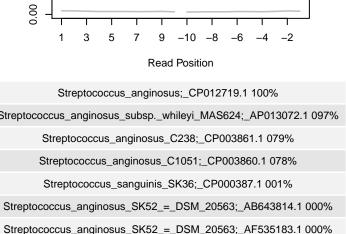


edit distance: default

0

C2T/G2A rate





Streptococcus\_intermedius\_C270;\_CP003858.1 000% Streptococcus\_anginosus\_SK52\_=\_DSM\_20563;\_KY315441.1 000% Streptococcus\_cristatus\_AS\_1.3089;\_CP004409.1 000%

