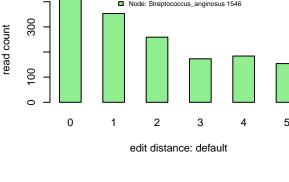
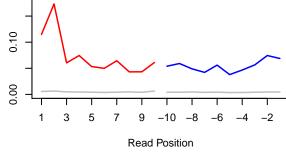
Node: Streptococcus_anginosus 1 ■ Node: Streptococcus_anginosus 1546

read count

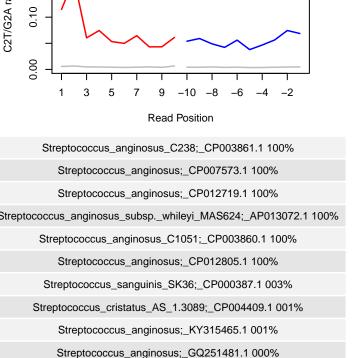
S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu

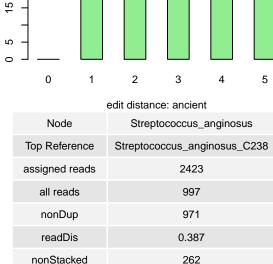






C2T/G2A rate





Off

0.1149

0.0688

106 (40.029)

destacking?

C>T_1

G>A_-1

mean length (sd)