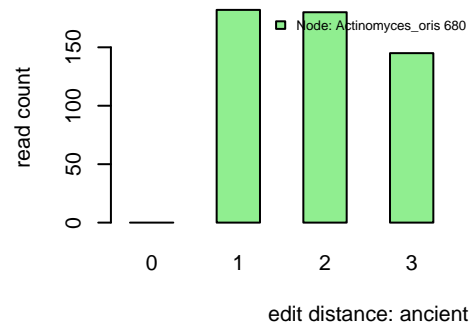
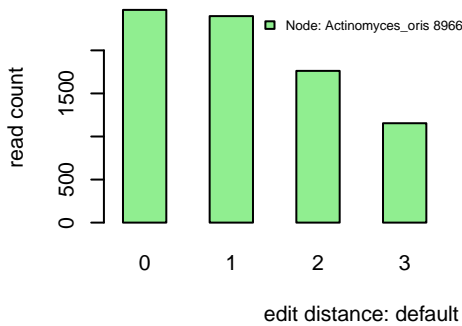
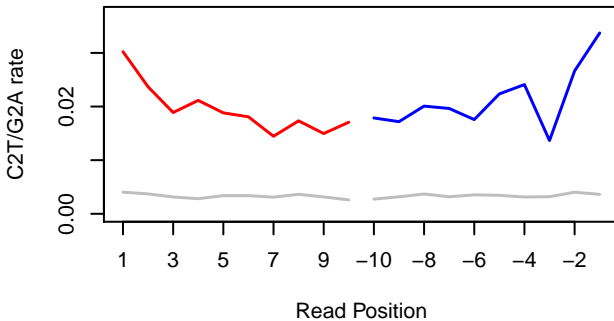


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Actinomyces_oris node



Node	Actinomyces_oris
Top Reference	Actinomyces_oris
assigned reads	27258
all reads	9943
nonDup	9772
readDis	0.611
nonStacked	4134
destacking?	Off
C>T_1	0.0302
G>A_1	0.0337
mean length (sd)	61 (23.171)

Actinomyces_oris;_CP014232.1	100%
Actinomyces_naeslundii;_AP017894.1	0 000%
Actinomyces;_JF825158.1	0 000%
Actinomyces_naeslundii;_DQ425099.1	0 000%
Actinomyces_naeslundii;_AF019629.1	0 000%
Actinomyces_naeslundii;_AF106035.1	0 000%
Actinomyces_naeslundii;_AF107020.1	0 000%
Actinomyces_naeslundii;_DQ425102.1	0 000%
Actinomyces_viscosus;_L06898.1	0 000%
Actinomyces_viscosus;_X62276.1	0 000%