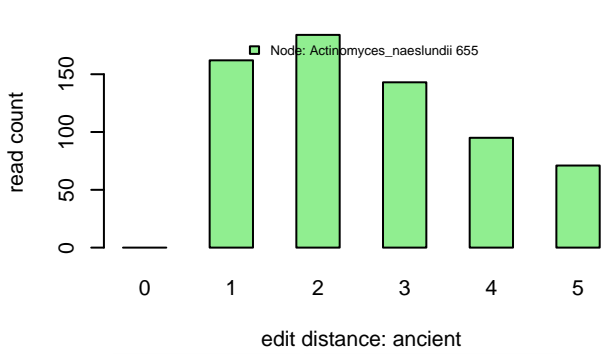
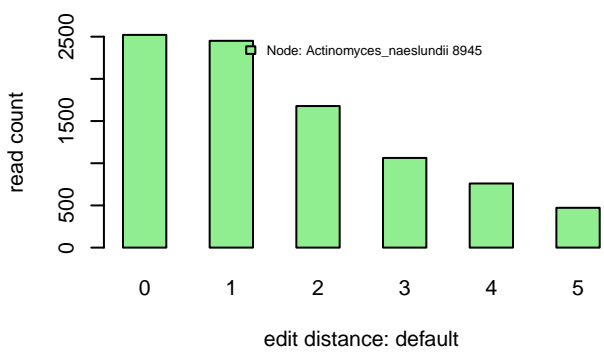
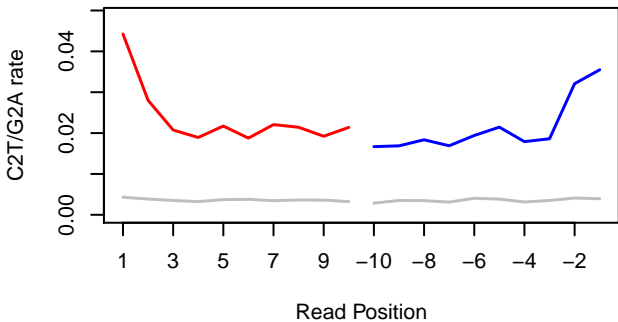


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Actinomyces_naeslundii node



Actinomyces_naeslundii;_AP017894.1	100%
Actinomyces_oris;_CP014232.1	0.000%
Actinomyces_oris;_EU621066.1	0.000%
NA	%
NA	%
NA	%
NA	%
NA	%
NA	%
NA	%

Node	Actinomyces_naeslundii
Top Reference	Actinomyces_naeslundii
assigned reads	24710
all reads	9957
nonDup	9760
readDis	0.592
nonStacked	3941
destacking?	Off
C>T_1	0.0443
G>A_-1	0.0355
mean length (sd)	60 (23.088)