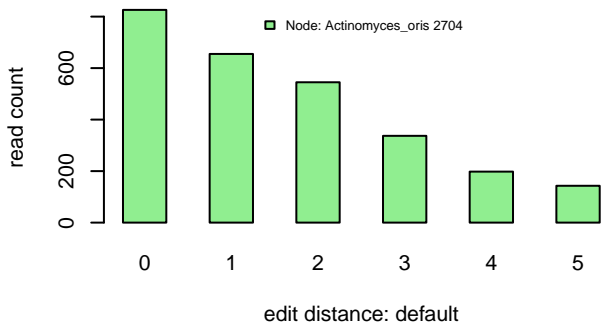
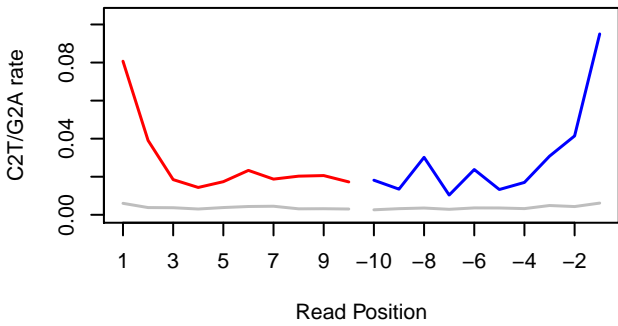


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



### Damage plot for Actinomyces\_oris node



Node	Actinomyces_oris
Top Reference	Actinomyces_oris
assigned reads	3060
all reads	3034
nonDup	2085
readDis	0.723
nonStacked	1478
destacking?	Off
C>T_1	0.0807
G>A_-1	0.095
mean length (sd)	60 (25.185)

Actinomyces_oris;_CP014232.1	100%
Actinomyces_naeslundii;_AP017894.1	0.000%
Actinomyces_oris_K20;_AB573870.1	0.000%
Actinomyces_naeslundii;_DQ658412.1	0.000%
Actinomyces_naeslundii;_AF107019.1	0.000%
Actinomyces_viscosus;_M32067.1	0.000%
Actinomyces_viscosus;_EU621125.1	0.000%
Actinomyces_odontolyticus;_EU647590.1	0.000%
NA	
NA	