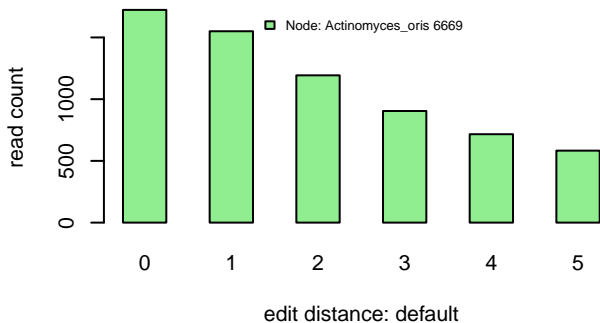
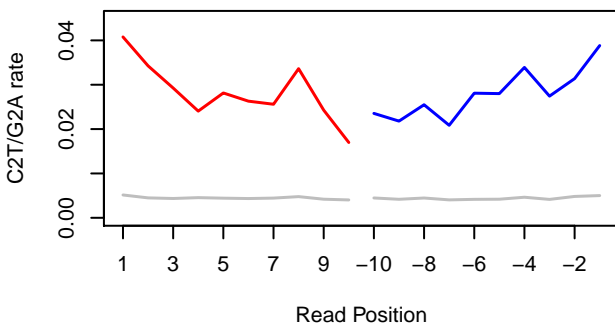


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for Actinomyces\_oris node



Actinomyces_oris;_CP014232.1	100%
Actinomyces_naeslundii;_AP017894.1	0.000%
Actinomyces_oris_K20;_AB573870.1	0.000%
Actinomyces_naeslundii;_EU603199.1	0.000%
Actinomyces_naeslundii;_EU805620.1	0.000%
Actinomyces_naeslundii;_EU805612.1	0.000%
Actinomyces_viscosus;_L06898.1	0.000%
Actinomyces_sp._Chiba101;_AP017896.1	0.000%
Actinomyces_sp._oral_taxon_414;_CP012590.1	0.000%
Actinomyces_naeslundii;_EU621259.1	0.000%

Node	Actinomyces_oris
Top Reference	Actinomyces_oris
assigned reads	643089
all reads	9943
nonDup	9885
readDis	0.565
nonStacked	3672
destacking?	Off
C>T_1	0.0408
G>A_-1	0.0388
mean length (sd)	92 (27.982)