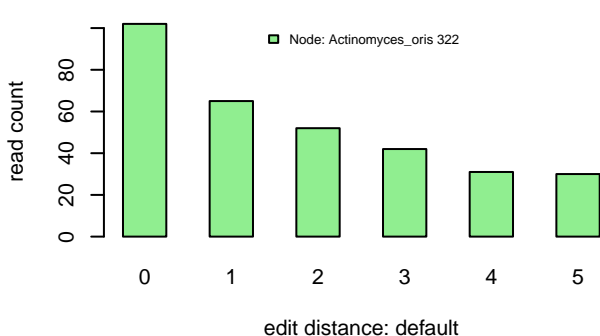
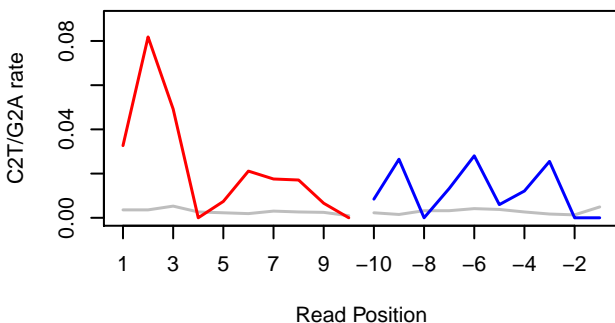


0401\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0401\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for Actinomyces\_oris node



Actinomyces_oris;_CP014232.1	100%
Actinomyces_naeslundii;_AP017894.1	001%
Actinomyces_viscosus;_EU620893.1	001%
Actinomyces_oris_K20;_AB573870.1	001%
NA	%
NA	%
NA	%
NA	%
NA	%
NA	%

Node	Actinomyces_oris
Top Reference	Actinomyces_oris
assigned reads	406
all reads	399
nonDup	230
readDis	0.731
nonStacked	166
destacking?	Off
C>T_1	0.0327
G>A_-1	0
mean length (sd)	84 (21.32)