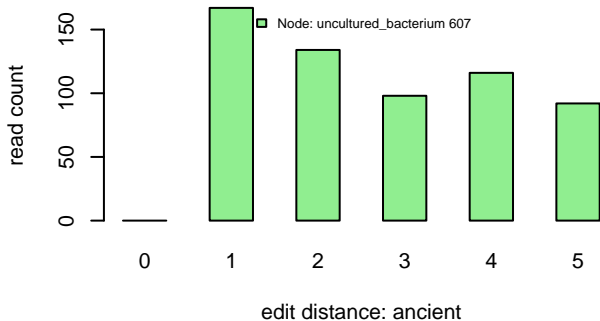
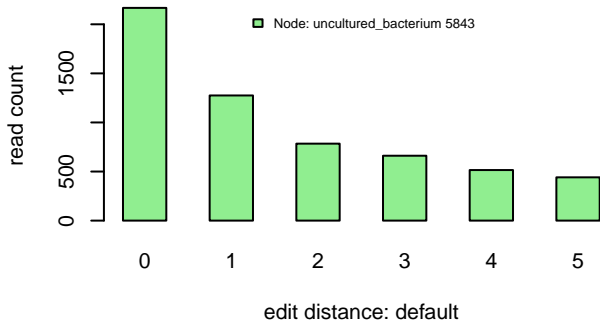
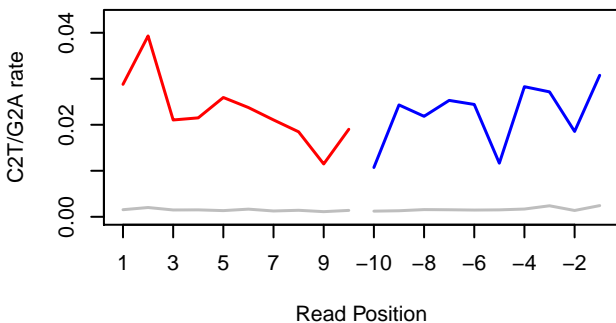


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for uncultured\_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	7434
all reads	375
nonDup	350
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.0288
G>A_-1	0.0308
mean length (sd)	67 (20.465)

uncultured_bacterium;_KU545305.1 100%
unidentified_proteobacterium;_X73223.1 100%
Lautropia_mirabilis;_GU397891.1 100%
Actinomyces_sp._oral_clone_EP011;_AY008315.1 100%
Actinomyces_sp._CM84;_HQ610192.1 100%
uncultured_Lautropia_sp.;_AY807005.1 100%
Actinomyces_naeslundii;_GU561309.1 100%
Actinomyces_naeslundii;_AB062278.1 100%
uncultured_Lautropia_sp.;_HQ616276.1 100%
Actinomyces_naeslundii;_AJ234045.1 100%