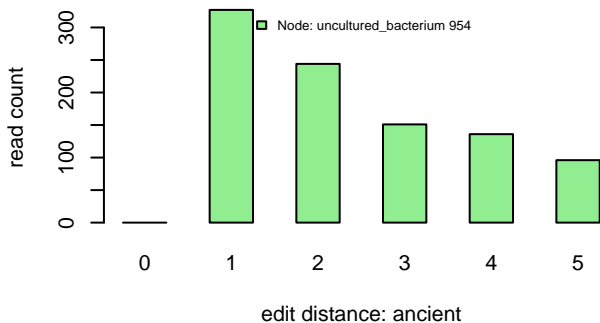
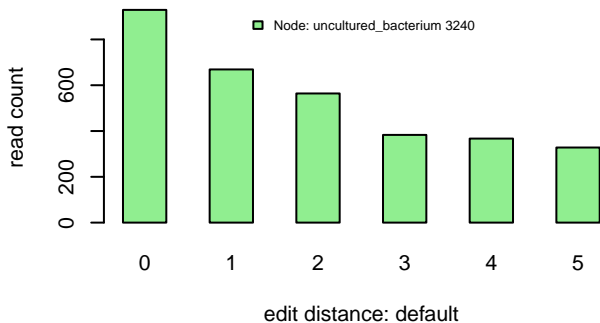
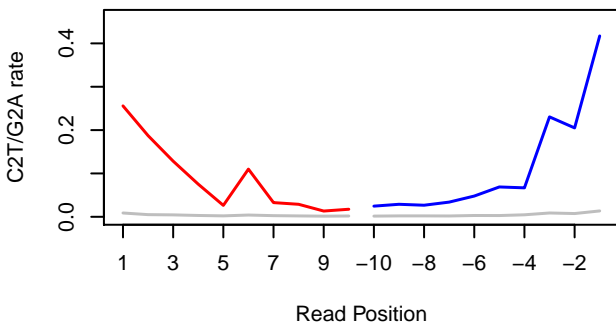


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



### Damage plot for uncultured\_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	8487
all reads	153
nonDup	41
readDis	0.154
nonStacked	9
destacking?	Off
C>T_1	0.2558
G>A_-1	0.4172
mean length (sd)	65 (28.676)

uncultured_bacterium;_KF835381.1 100%
uncultured_Chlorobiaceae_bacterium;_AM749878.1 100%
Chlorobium_sp._ShCl03;_EF153291.1 100%
Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%
bacterium;_MF314746.1 100%
Bacteria;_EU888384.1 100%
uncultured_Propionibacterium_sp.;_AM420135.1 100%
uncultured_actinobacterium;_EF999054.1 100%
Pseudopropionibacterium_propionicum;_HQ413290.1 100%
uncultured_Tessaracoccus_sp.;_KC675457.1 100%