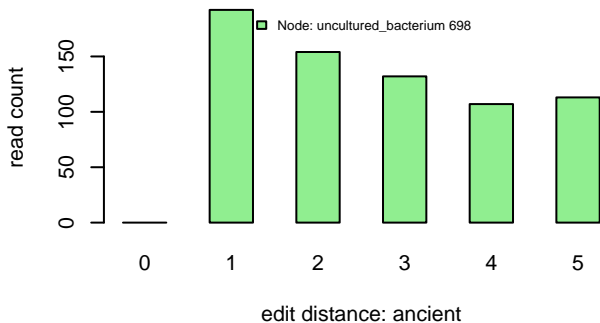
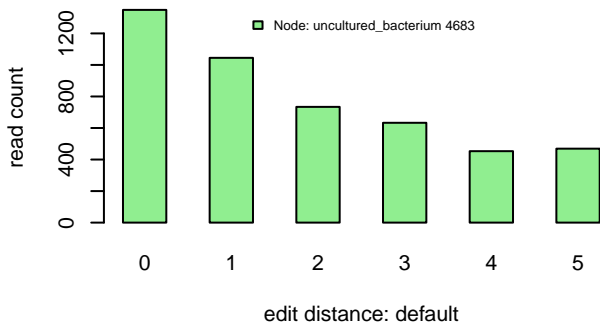
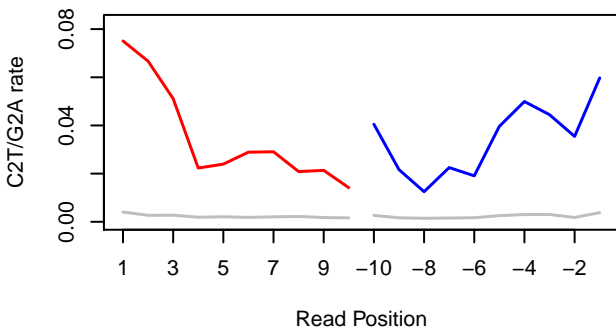


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



### Damage plot for uncultured\_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	5681
all reads	155
nonDup	150
readDis	0.002
nonStacked	0
destacking?	Off
C>T_1	0.0751
G>A_-1	0.0597
mean length (sd)	51 (19.247)

uncultured\_bacterium;\_JQ696889.1 100%

Bacteria;\_MF322780.1 100%

Olsenella\_sp.\_F0004;\_EU592964.1 100%

Actinomyces\_israelii;\_AF479270.1 100%

uncultured\_Coriobacteriaceae\_bacterium;\_JQ083420.1 100%

Olsenella\_sp.\_F0206;\_EU592965.1 100%

uncultured\_actinobacterium;\_JN314630.1 100%

Senegalimassilia\_anaerobia\_JC110;\_NR\_125595.1 100%

uncultured\_Olsenella\_sp.;\_EU669568.1 100%

Slackia\_piriformis\_YIT\_12062;\_NR\_112898.1 100%