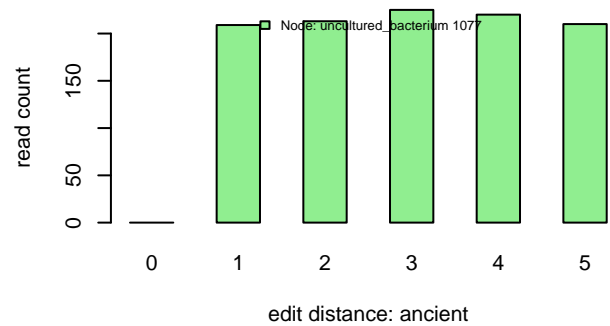
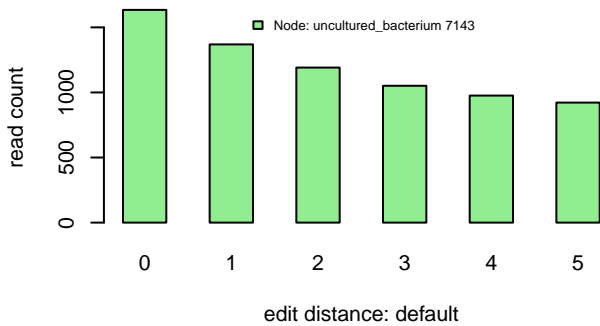
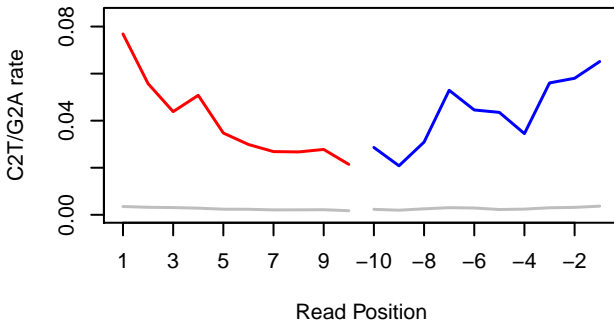


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	10677
all reads	845
nonDup	833
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.0769
G>A_-1	0.0651
mean length (sd)	57 (21.383)

uncultured_bacterium;_JQ696889.1 100%

uncultured_Actinomyces_sp.;_AM420166.1 100%

Actinomyces;_NR_025633.1 100%

uncultured_Actinomyces_sp.;_FJ976258.1 100%

Actinomyces_israelii;_AF479270.1 100%

uncultured_human_oral_bacterium_A15;_AF202026.1 100%

Actinomyces_sp._oral_taxon_448_str._F0400;_HM596273.1 100%

uncultured_human_oral_bacterium_A16;_AF202027.1 100%

Actinomyces_genomosp._C1;_AY278610.1 100%

uncultured_Actinomyces_sp.;_GU227173.1 100%