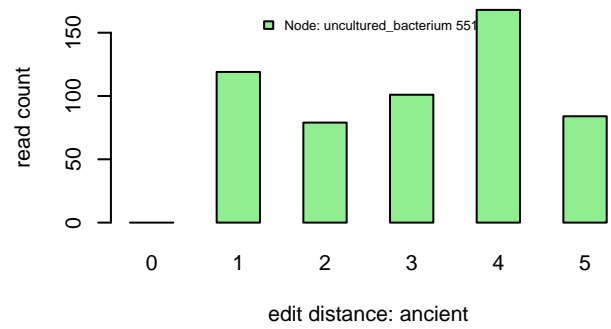
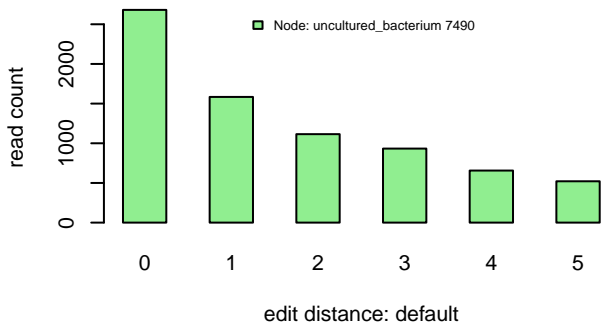
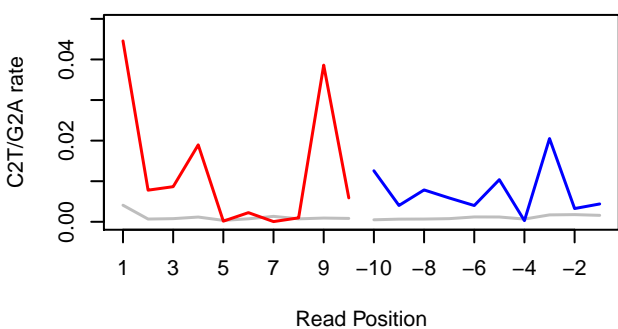


0519_S0_L000_R1_000.fastq.merged.prefixed.hg19un0519_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	13196
all reads	852
nonDup	113
readDis	0.004
nonStacked	0
destacking?	Off
C>T_1	0.0446
G>A_-1	0.0044
mean length (sd)	60 (20.871)

uncultured_bacterium;_HE681331.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
Actinomyces_genomosp._C1;_AY278610.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
Actinomyces_sp._oral_taxon_414;_CP012590.1 100%
uncultured_Actinomyces_sp.;_FJ976446.1 100%
Bacteria;_FJ976368.1 100%
Bacteria;_NR_036819.1 100%
Atopobium_rimae;_X67149.1 100%