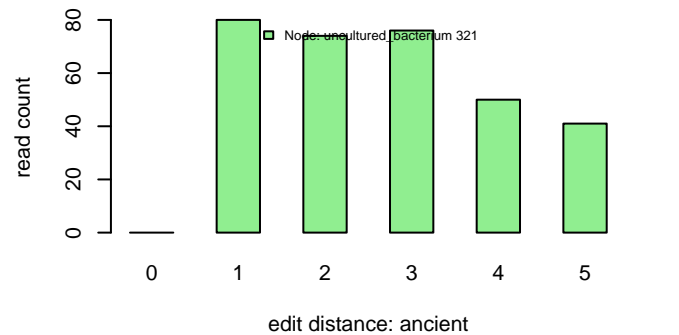
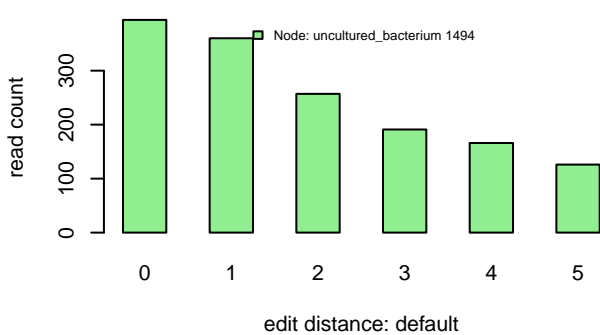
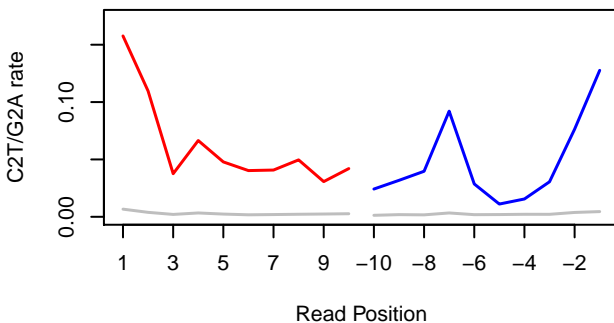


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	1854
all reads	55
nonDup	55
readDis	0.617
nonStacked	13
destacking?	Off
C>T_1	0.1576
G>A_-1	0.1277
mean length (sd)	55 (19.62)

uncultured_bacterium;_JQ972699.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
uncultured_Actinomyces_sp.;_FJ976446.1 100%
Actinomyces_genomosp._C1;_AY278610.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
Actinomyces_sp._oral_taxon_414;_CP012590.1 100%
Bacteria;_FJ976368.1 100%