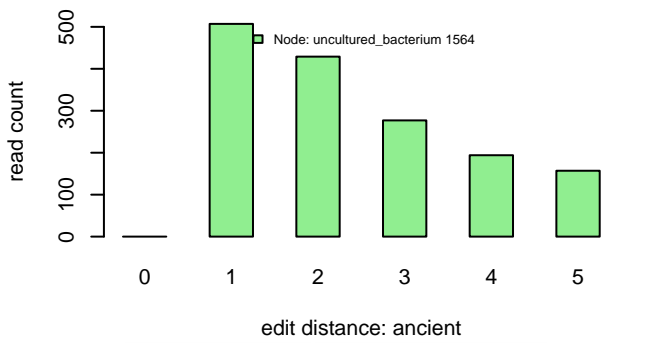
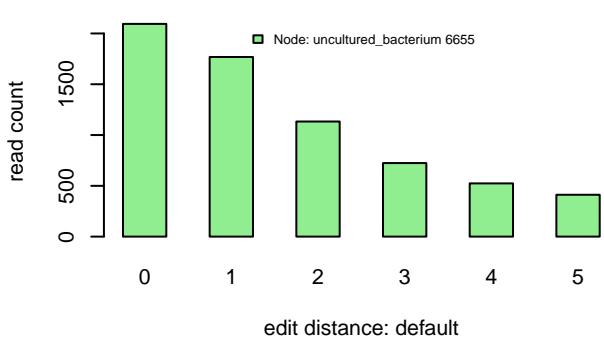
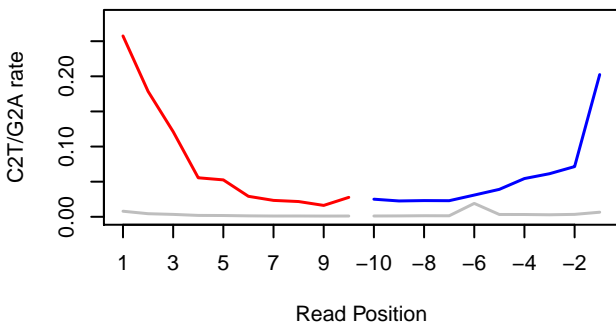


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	8100
all reads	376
nonDup	350
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.2575
G>A_-1	0.2025
mean length (sd)	65 (13.491)

uncultured_bacterium;_JQ696889.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
Actinomyces;_NR_025633.1 100%
Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%
Actinomyces_massiliensis_4401292;_NR_044288.1 100%
Streptococcus_pneumoniae;_AJ608938.1 100%
Actinomyces_sp._oral_taxon_414;_CP012590.1 100%