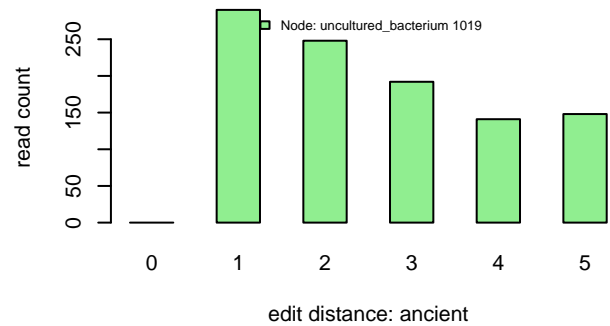
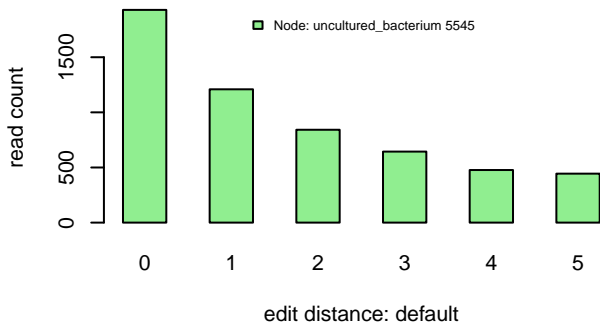
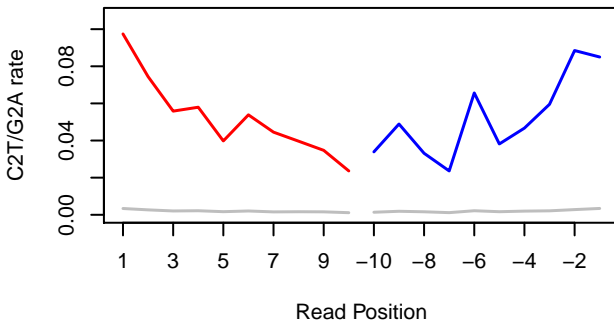


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	7092
all reads	306
nonDup	298
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.0974
G>A_-1	0.085
mean length (sd)	66 (24.971)

uncultured_bacterium;_FM873742.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_AM420132.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976255.1 100%
Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_AY144358.1 100%