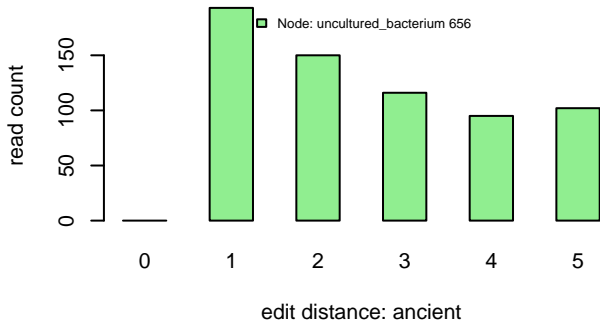
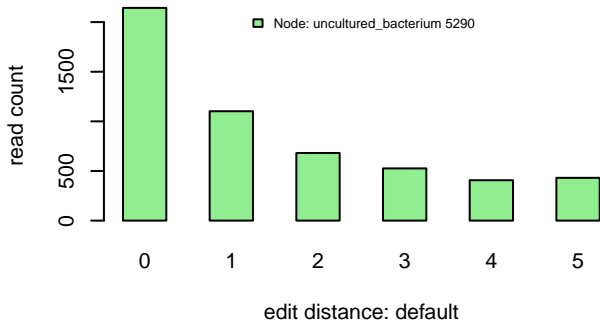
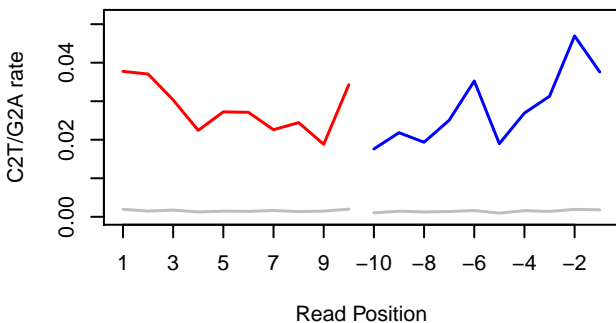


101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for uncultured_bacterium node



uncultured_bacterium;_JF119278.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
Actinomyces_genomosp._C1;_AY278610.1 100%
Actinomyces_sp._oral_taxon_414;_CP012590.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_AM420132.1 100%
uncultured_Actinomyces_sp.;_FJ976446.1 100%

edit distance: ancient	
Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	6650
all reads	433
nonDup	425
readDis	0.002
nonStacked	1
destacking?	Off
C>T_1	0.0378
G>A_-1	0.0376
mean length (sd)	66 (25.52)