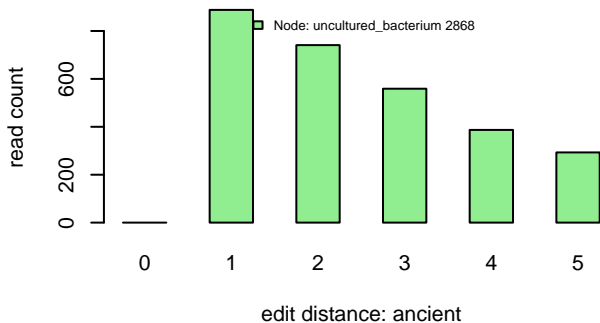
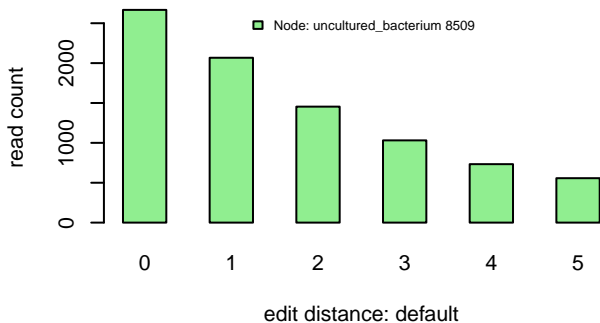
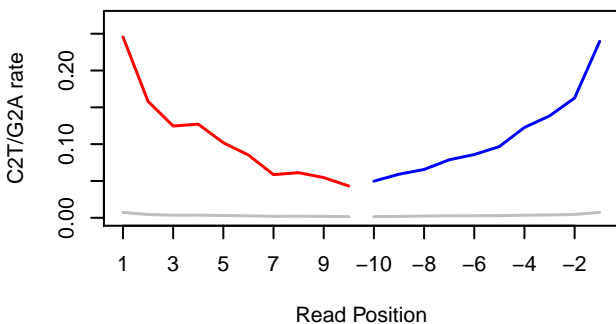


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	10769
all reads	1074
nonDup	1053
readDis	0.001
nonStacked	1
destacking?	Off
C>T_1	0.2456
G>A_1	0.2396
mean length (sd)	59 (22.738)

uncultured_bacterium;_FM873742.1 100%
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%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976257.1 100%
Actinomyces_genomosp._C1;_AY278610.1 100%
uncultured_Actinomyces_sp.;_FJ976446.1 100%
Actinomyces_sp._2002-42188;_AY244765.1 100%