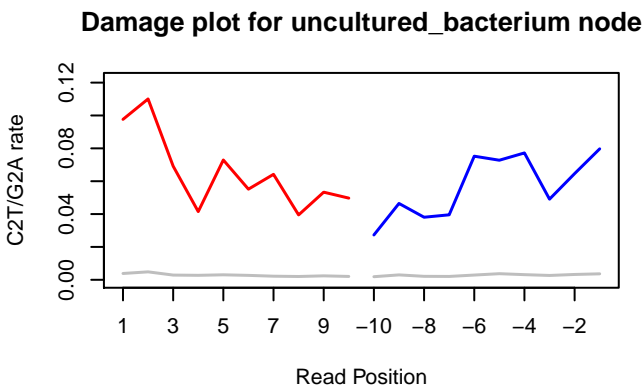
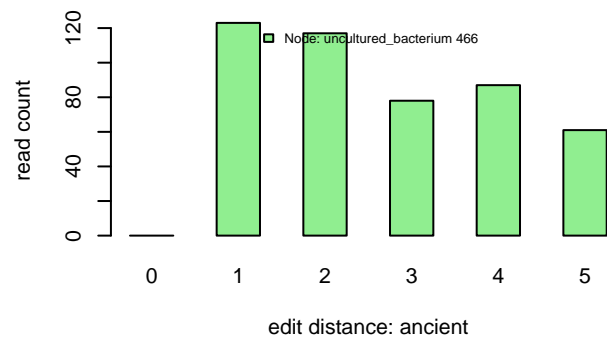
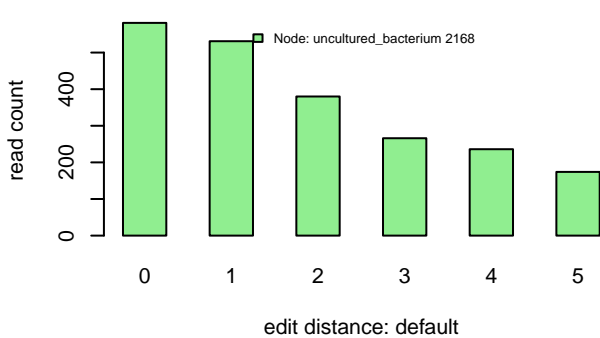


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	2593
all reads	81
nonDup	80
readDis	0.016
nonStacked	1
destacking?	Off
C>T_1	0.0976
G>A_-1	0.0797
mean length (sd)	50 (16.055)

uncultured_bacterium;_JQ696889.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
uncultured_Bacteroidales_bacterium;_FJ976400.1 100%
Bacteroidetes_bacterium_Oral_Taxon_274;_FJ577256.2 100%
Actinomyces_genomosp._C1;_AY278610.1 100%
uncultured_Actinomyces_sp.;_FJ976446.1 100%
uncultured_Porphyromonadaceae_bacterium;_AM420225.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%