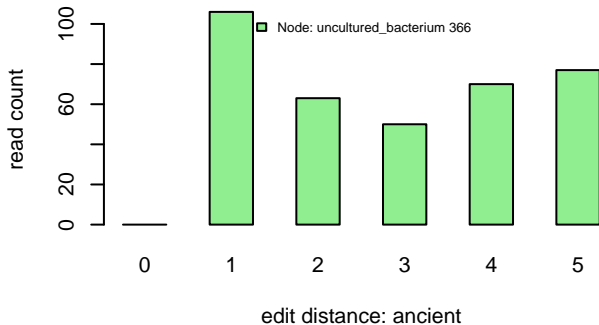
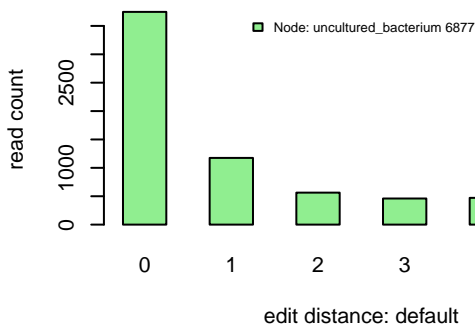
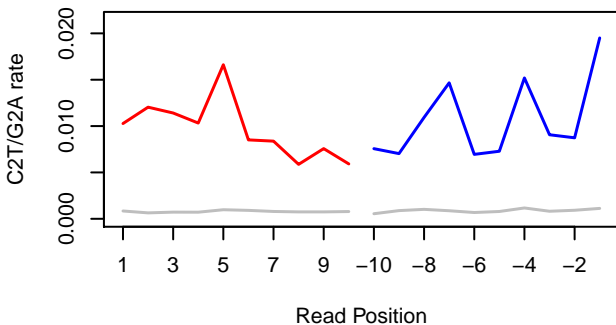


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for uncultured\_bacterium node



uncultured_bacterium;_JQ475838.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
actinomyces_genomsp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976255.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_AM420132.1 100%
Bacteroidetes_bacterium_Oral_Taxon_274;_FJ577256.2 100%
uncultured_Bacteroidales_bacterium;_FJ976400.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
Bacteroides-like_sp._oral_clone_AU126;_AY005072.1 100%
uncultured_Porphyromonadaceae_bacterium;_AM420225.1 100%

Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	77077
all reads	857
nonDup	847
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.0103
G>A_-1	0.0195
mean length (sd)	92 (28.505)