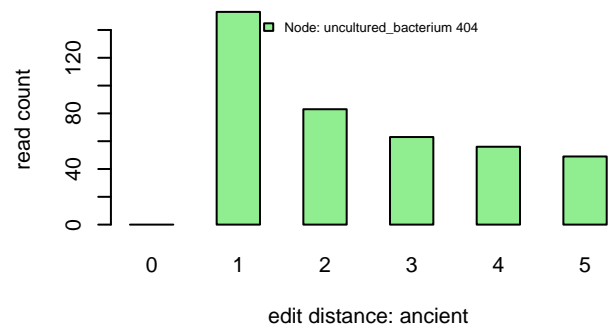
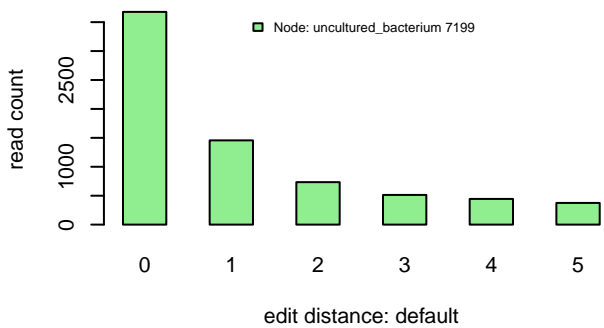
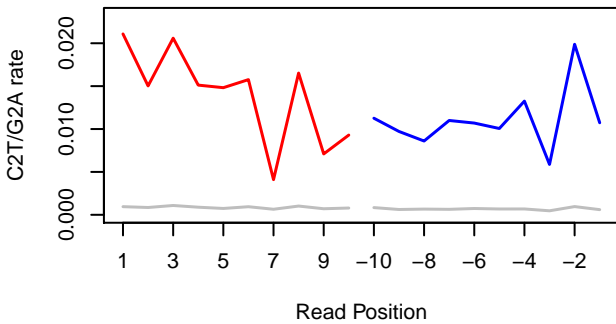


S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu



### Damage plot for uncultured\_bacterium node



uncultured_bacterium;_KF982315.1 100%
Bacteria;_FJ976437.1 100%
unidentified_proteobacterium;_X73223.1 100%
uncultured_Lautropia_sp.;_AM419986.1 100%
uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%
Ottowia_sp._oral_taxon_894;_CP012073.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
Lautropia_sp._TeTO;_FJ823153.1 100%
Lautropia_sp._oral_clone_FX006;_AY134902.1 100%

edit distance: ancient	
Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	43944
all reads	292
nonDup	287
readDis	0.072
nonStacked	3
destacking?	Off
C>T_1	0.0211
G>A_1	0.0107
mean length (sd)	114 (41.251)