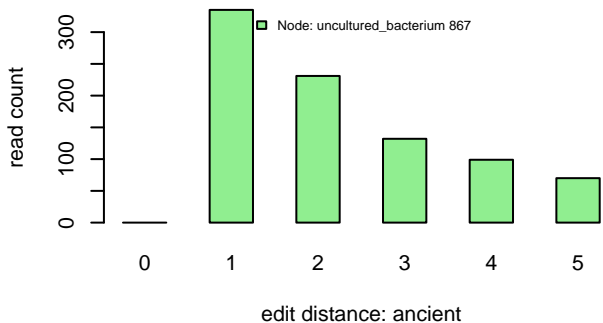
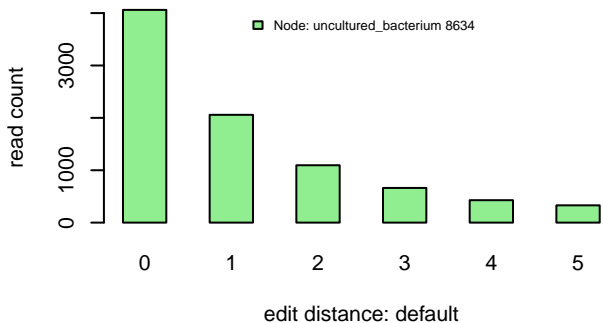
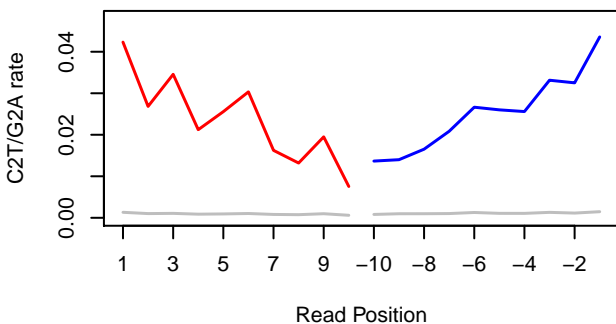


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	66127
all reads	455
nonDup	449
readDis	0.072
nonStacked	10
destacking?	Off
C>T_1	0.0423
G>A_-1	0.0436
mean length (sd)	90 (27.904)

uncultured_bacterium;_KF982315.1 100%
Bacteria;_FJ976437.1 100%
uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%
Ottowia_sp._oral_taxon_894;_CP012073.1 100%
Bergeyella_sp._AF14;_DQ241813.1 100%
Bacteria;_FJ976371.1 100%
uncultured_Bergeyella_sp.;_EU644465.1 100%
uncultured_Lautropia_sp.;_AM419986.1 100%
Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%
Lautropia_sp._oral_clone_FX006;_AY134902.1 100%