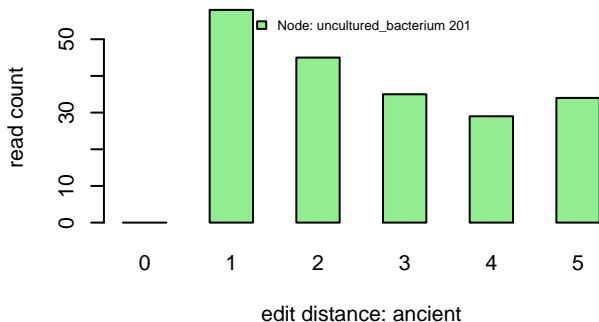
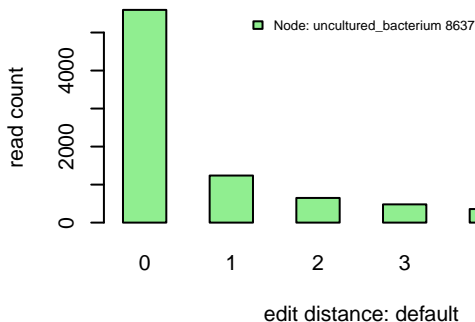
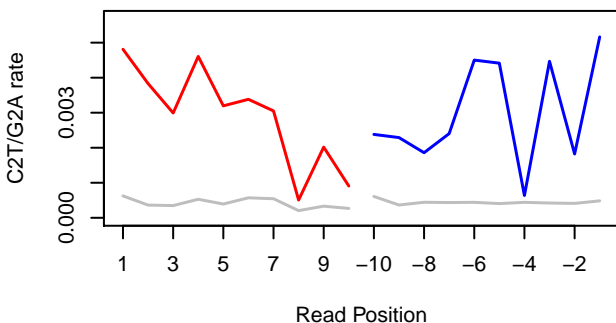


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	105442
all reads	711
nonDup	705
readDis	0.001
nonStacked	1
destacking?	Off
C>T_1	0.0048
G>A_-1	0.0052
mean length (sd)	94 (28.561)

uncultured_bacterium;_JQ472329.1 100%

Bacteria;_FJ976437.1 100%

uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%

uncultured_Lautropia_sp.;_AM419986.1 100%

Ottowia_sp._oral_taxon_894;_CP012073.1 100%

Lautropia_sp._oral_clone_FX006;_AY134902.1 100%

bacterium_str._86355;_AF227858.1 100%

Haemophilus_sp._Smarlab_3302188;_AY538693.1 100%

Aggregatibacter_aphrophilus_NJ8700;_CP001607.1 100%

Neisseria_elongata;_AJ239303.1 100%