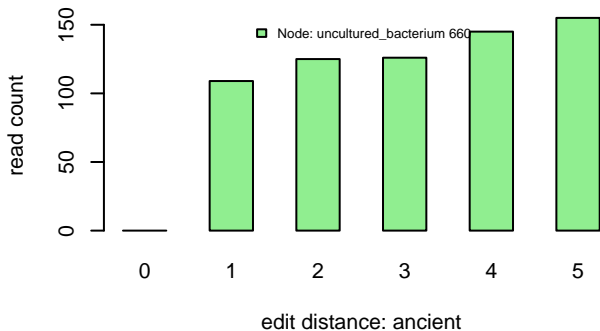
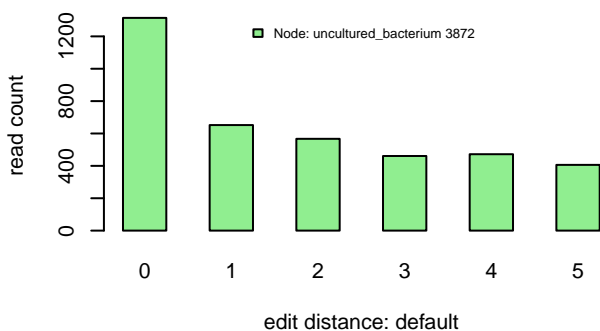
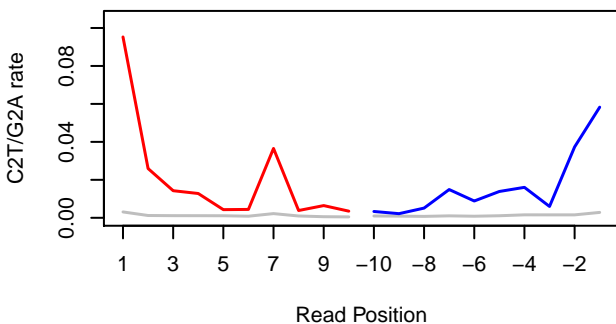


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	4872
all reads	254
nonDup	159
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.0953
G>A_-1	0.0583
mean length (sd)	58 (23.028)

uncultured_bacterium;_GQ014687.1 100%

uncultured_microorganism;_EU931404.1 100%

Porphyromonas_bennonis;_NR_044491.1 100%

uncultured_Porphyromonas_sp.;_DQ130022.1 100%

uncultured_Porphyromonadaceae_bacterium;_EU572512.1 100%

Bacteria;_EU702984.1 100%

Porphyromonas_sp._TSW25CA2;_FM945401.2 100%

uncultured_organism;_AY897754.1 100%

uncultured_Bacteroidales_bacterium;_HM080431.1 100%

uncultured_Anaerococcus_sp.;_DQ130021.1 100%