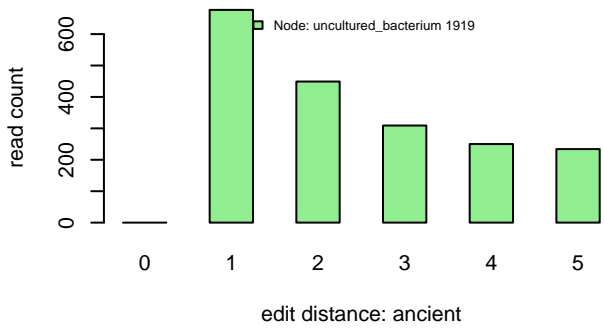
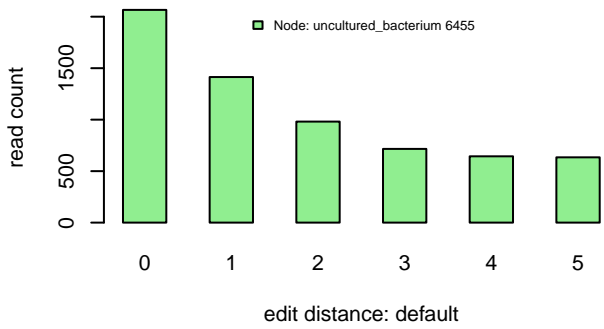
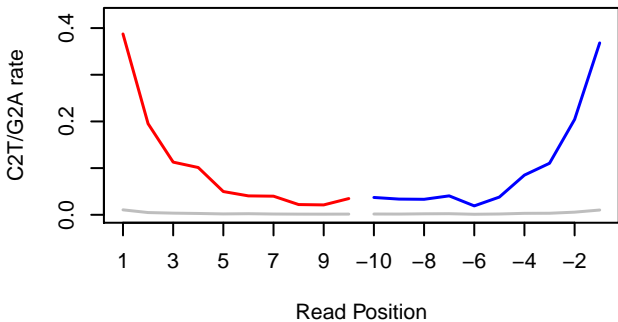


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	10531
all reads	242
nonDup	180
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.3874
G>A_-1	0.3682
mean length (sd)	71 (25.903)

uncultured_bacterium;_JQ696889.1 100%
Ottowia_sp._oral_taxon_894;_CP012073.1 100%
Bacteria;_FJ976437.1 100%
uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%
uncultured_Lautropia_sp.;_AM419986.1 100%
Lautropia_sp._oral_clone_FX006;_AY134902.1 100%
Atopobium_parvulum_DSM_20469;_CP001721.1 100%
Abiotrophia_defectiva;_AY879307.1 100%
Abiotrophia_defectiva;_AY879306.1 100%
uncultured_Gemella_sp.;_EU071466.1 100%