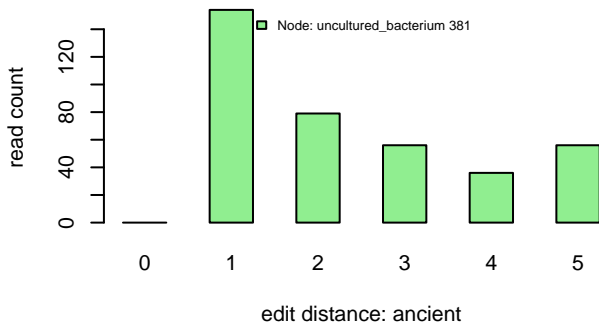
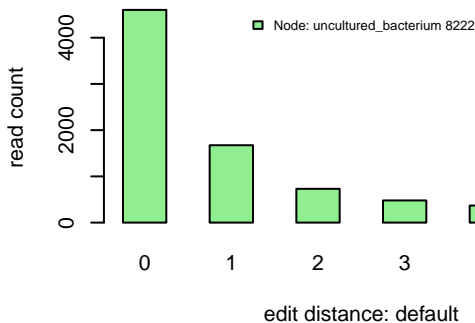
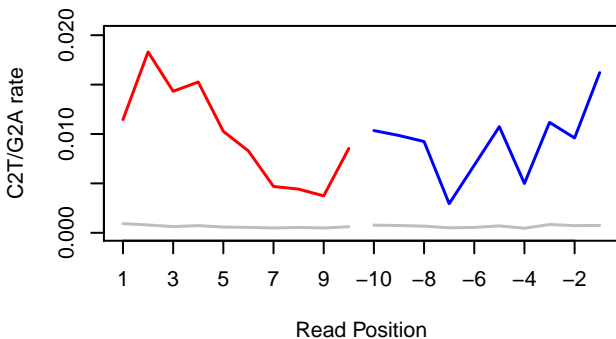


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



edit distance: ancient	
Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	70877
all reads	1084
nonDup	1076
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.0115
G>A_-1	0.0162
mean length (sd)	95 (28.131)

uncultured_bacterium;_JQ451620.1 100%

Bacteria;_FJ976437.1 100%

uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%

uncultured_Lautropia_sp.;_AM419986.1 100%

Lautropia_sp._oral_clone_FX006;_AY134902.1 100%

Ottowia_sp._oral_taxon_894;_CP012073.1 100%

uncultured_Lautropia_sp.;_AY806314.1 100%

bacterium;_MF801248.1 100%

uncultured_Brachymonas_sp.;_KR852686.1 100%

uncultured_Leptotrichia_sp.;_FJ976247.1 100%