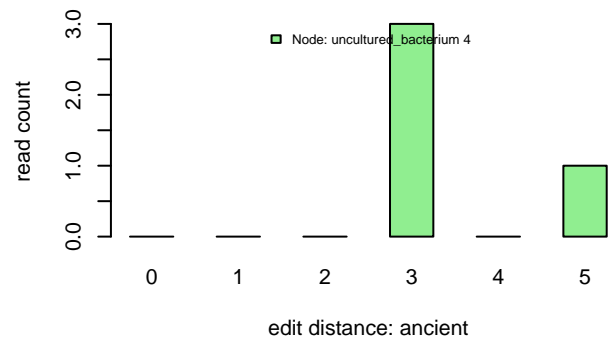
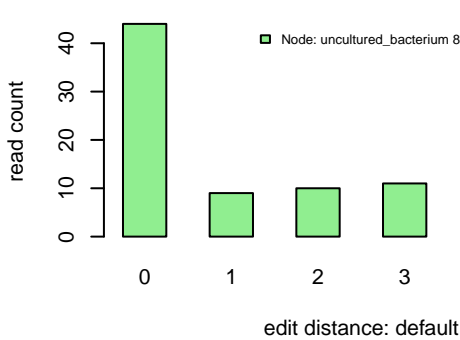
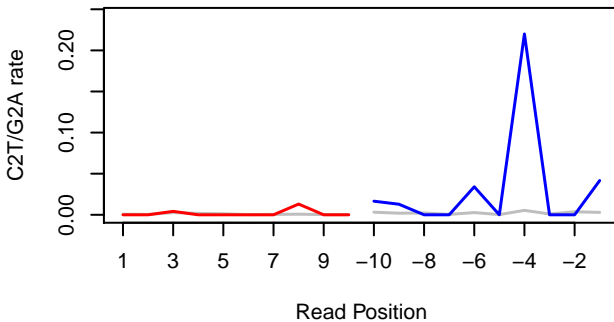


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	98
all reads	16
nonDup	10
readDis	0.806
nonStacked	8
destacking?	Off
C>T_1	0
G>A_-1	0.0417
mean length (sd)	73 (23.532)

uncultured_bacterium;_JX194159.1 100%
Prevotella_sp._SN-2;_KM016086.1 050%
Prevotella_sp._NAIP3C;_JQ797589.1 050%
uncultured_Prevotella_sp.;_KJ590244.1 050%
uncultured_Bacteroidetes_bacterium;_AB874502.1 050%
Pseudomonas_lini;_LT629746.1 012%
uncultured_Pseudomonas_sp.;_DQ426915.1 012%
uncultured_Pseudomonas_sp.;_DQ426916.1 012%
bacterium_NLAE-zl-G96;_JX048144.1 012%
uncultured_Firmicutes_bacterium;_KJ597304.1 012%