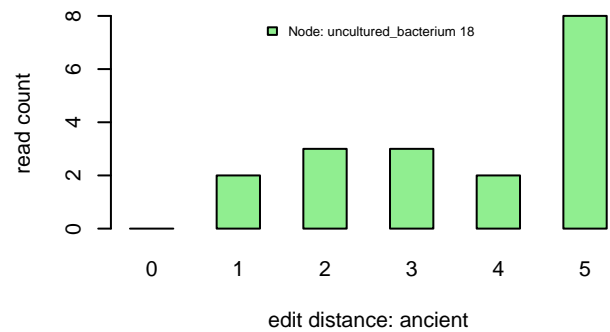
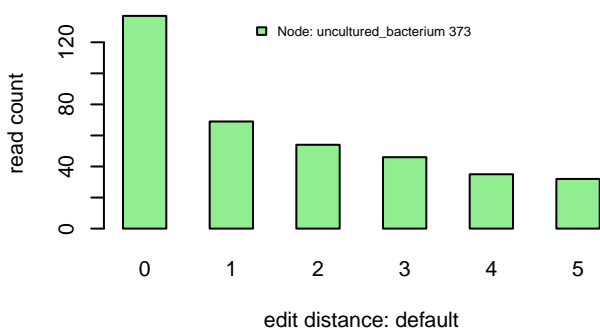
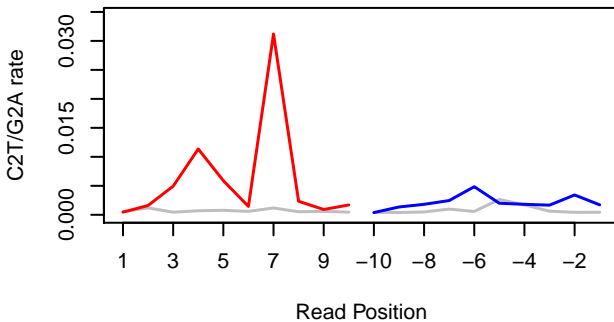


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	490
all reads	77
nonDup	76
readDis	0.018
nonStacked	0
destacking?	Off
C>T_1	5e-04
G>A_-1	0.0017
mean length (sd)	73 (27.016)

uncultured_bacterium;_EU059182.1 100%

Actinomyces_viscosus;_FJ823139.1 100%

Actinomyces_oris;_GU561322.1 100%

Actinomyces_viscosus;_AJ234056.1 100%

Actinomyces_sp._oral_clone_JA063;_AY349367.1 100%

Actinomyces_viscosus;_AJ234046.1 100%

Actinomyces_sp._oral_taxon_175_str._F0384;_HM596281.1 100%

Propionibacterium_sp._MA;_EU337118.1 100%

Actinomyces_oris;_GQ421311.1 100%

uncultured_Actinomyces_sp.;_JN802684.1 100%