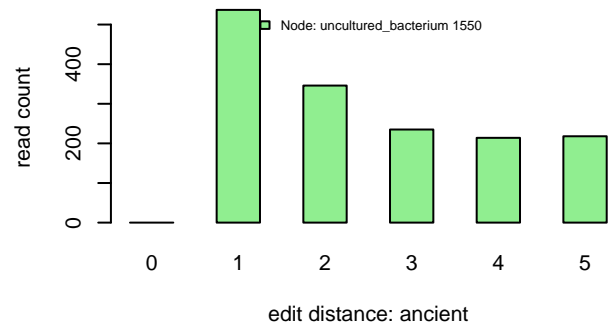
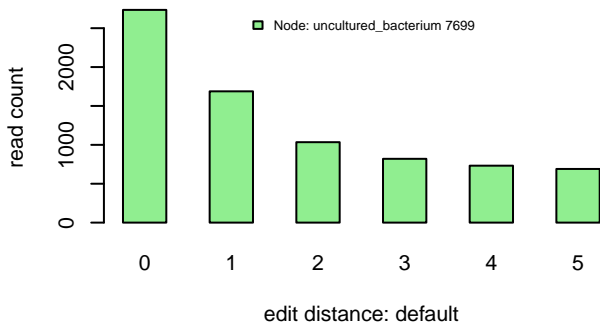
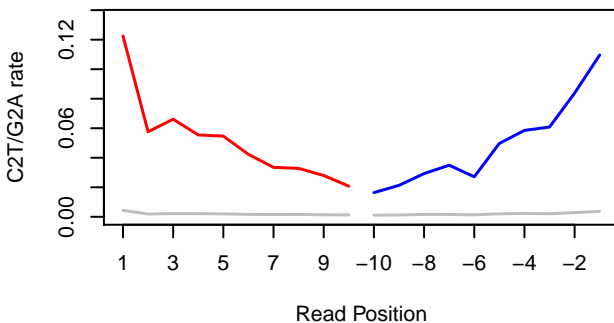


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



uncultured_bacterium;_JQ696889.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_AY144358.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976255.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_GU227159.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_GU227157.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_AM420132.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976288.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_GU227158.1 100%

Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	11062
all reads	455
nonDup	451
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.1225
G>A_-1	0.1096
mean length (sd)	59 (22.854)