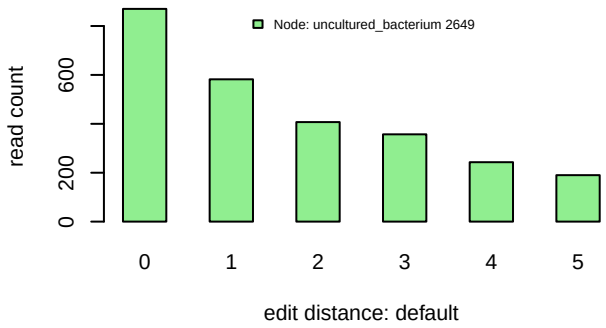
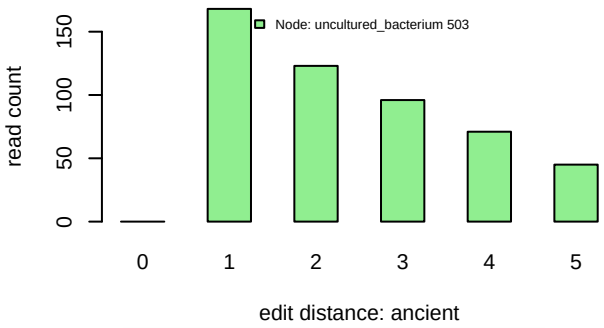
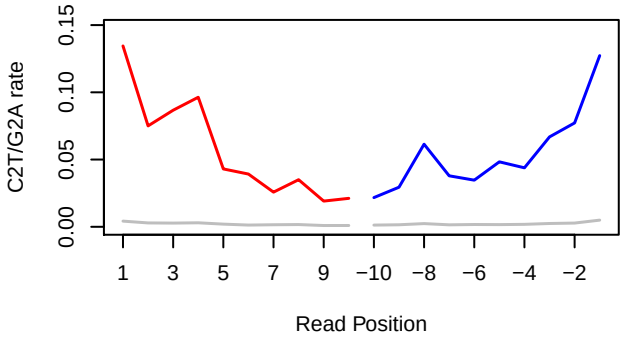
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur





Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	3162
all reads	406
nonDup	400
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.1345
G>A_−1	0.1273
mean length (sd)	54 (14.752)

uncultured_bacterium;_JQ998551.1 100%
Actinomyces_massiliensis_4401292;_NR_044288.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976257.1 100%
Actinomyces_massiliensis_4401292;_NR_113059.1 100%
uncultured_Candidatus_Saccharibacteria_bacterium;_HQ616338.1 100%
uncultured_human_oral_bacterium_A4;_AF201995.1 100%
Clostridiales_bacterium_canine_oral_taxon_157;_JN713323.1 100%
Actinomyces_sp._oral_taxon_180_str._F0310;_GQ131415.1 100%
Candidatus_Saccharibacteria;_KM462163.1 100%
Actinomyces_oris;_GQ421309.1 100%