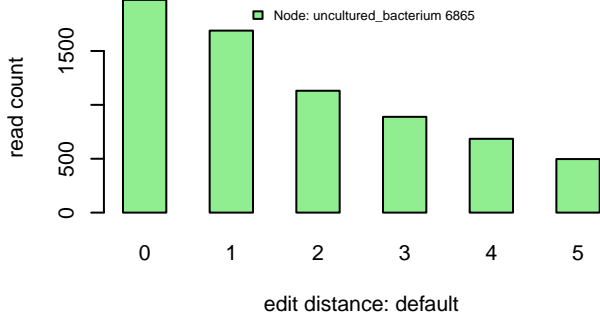
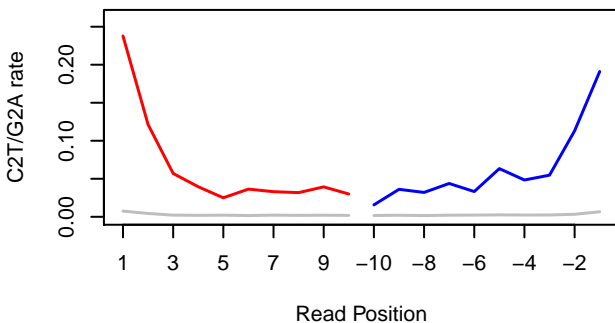


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	8228
all reads	1339
nonDup	1206
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.2379
G>A_-1	0.1911
mean length (sd)	59 (21.461)

uncultured_bacterium;_JQ216016.1 100%
uncultured_Actinomyces_sp.;_FJ976253.1 100%
Actinomyces;_NR_025633.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
Actinomyces_gerencseriae;_NR_029280.1 100%
Actinomyces_gerencseriae;_X80414.1 100%
Actinomyces_israelii;_AF479270.1 100%