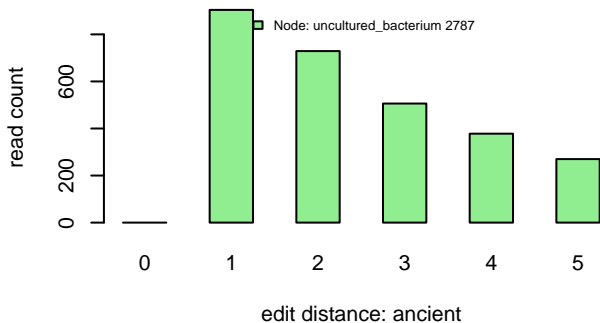
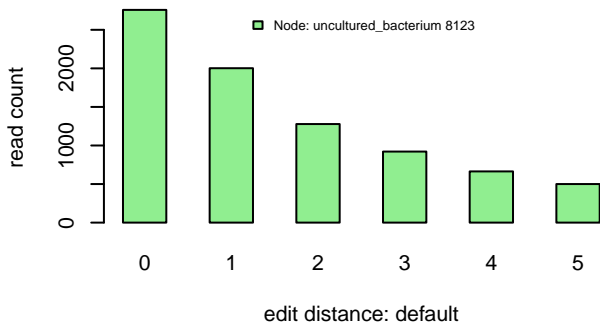
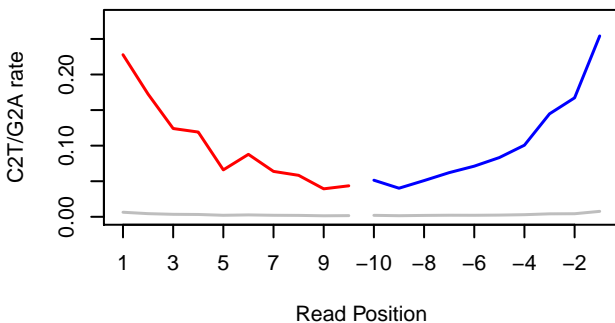


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	9193
all reads	330
nonDup	322
readDis	0.004
nonStacked	0
destacking?	Off
C>T_1	0.2278
G>A_-1	0.2542
mean length (sd)	51 (18.888)

uncultured_bacterium;_JF231237.1 100%
uncultured_Actinobaculum_sp.;_FJ976277.1 100%
Actinomycetaceae_bacterium_BA112;_JX171198.2 100%
Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%
Actinomyces_sp_oral_taxon_848_str_F0332;_GQ131416.1 100%
uncultured_human_oral_bacterium_A11;_AF202002.1 100%
Pseudopropionibacterium_propionicum;_HQ413290.1 100%
uncultured_Propionibacterium_sp.;_AM420135.1 100%
Pseudopropionibacterium_propionicum;_NR_114803.1 100%
uncultured_Propionibacterium_sp.;_FJ976211.1 100%