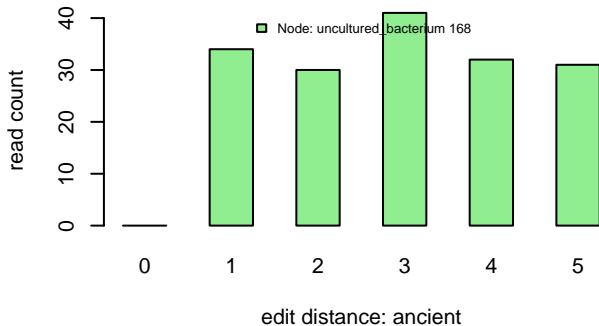
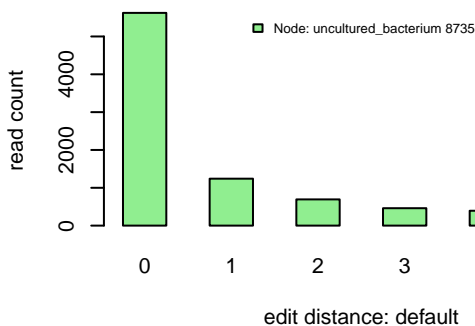
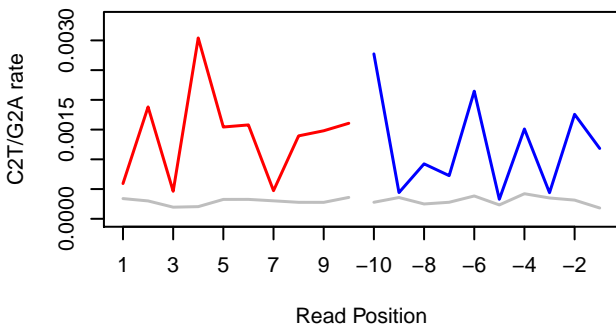


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for uncultured\_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	88604
all reads	830
nonDup	822
readDis	0
nonStacked	0
destacking?	Off
C>T_1	6e-04
G>A_-1	0.0012
mean length (sd)	93 (28.026)

uncultured_bacterium;_HM263417.1 100%
Aggregatibacter_aphrophilus_NJ8700;_CP001607.1 100%
bacterium_str._86355;_AF227858.1 100%
Aggregatibacter_aphrophilus;_GU470905.1 100%
Aggregatibacter_aphrophilus_NJ8700;_EF605278.1 100%
Haemophilus_sp._Smarlab_3302188;_AY538693.1 100%
uncultured_Haemophilus_sp.;_AM419971.1 100%
Bacteria;_FJ976437.1 100%
Aggregatibacter_aphrophilus;_HQ009508.1 100%
laemophilus_genomosp._P2_oral_clone_MB3_C24;_DQ003621.1 100%