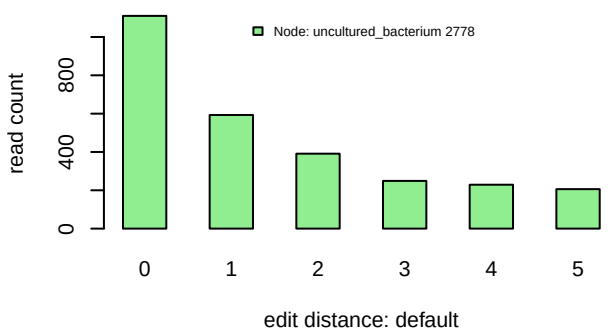
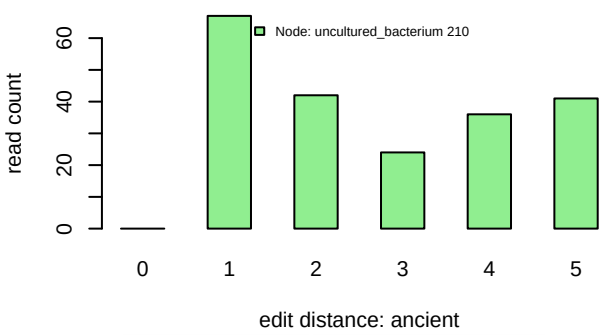
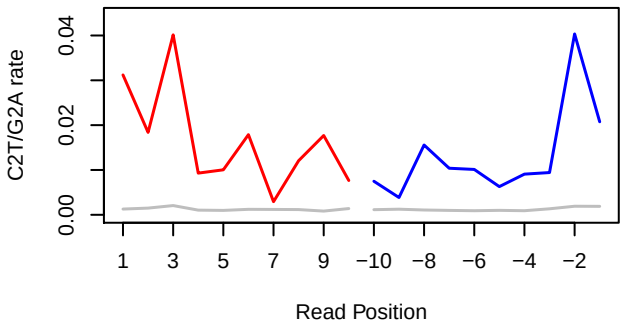
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un





Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	3441
all reads	81
nonDup	74
readDis	0.044
nonStacked	1
destacking?	Off
C>T_1	0.0312
G>A_−1	0.0207
mean length (sd)	60 (19.325)

uncultured_bacterium;_JF229185.1 100%
Haemophilus_simiae;_DQ851143.1 100%
Actinomyces_gerencseriae;_X80414.1 100%
uncultured_Pasteurellaceae_bacterium;_KC675367.1 100%
Actinomyces_gerencseriae;_NR_029280.1 100%
uncultured_Aggregatibacter_sp.;_KC675780.1 100%
Actinomyces_sp._2002−42188;_AY244765.1 100%
Aggregatibacter_aphrophilus_NJ8700;_CP001607.1 100%
uncultured_Actinomyces_sp.;_FJ976253.1 100%
uncultured_Haemophilus_sp.;_GU902763.1 100%