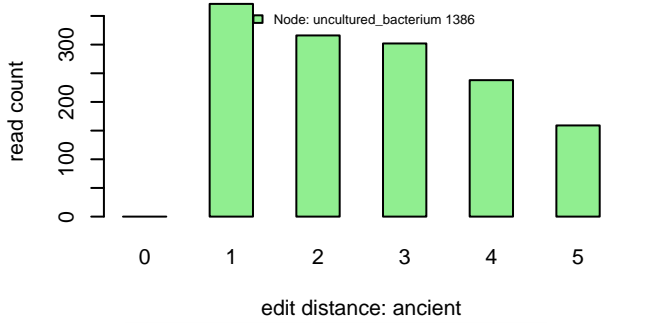
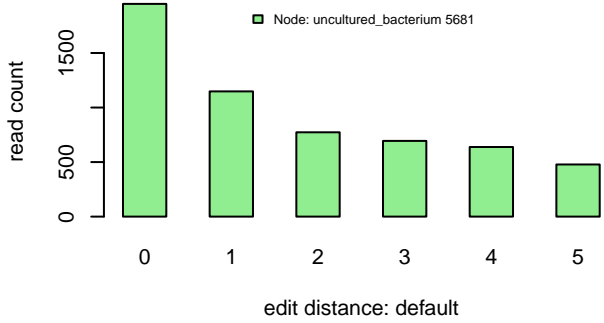
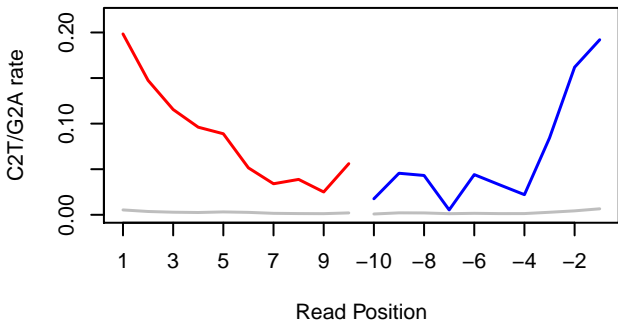


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	7287
all reads	679
nonDup	362
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.1984
G>A_-1	0.192
mean length (sd)	61 (22.619)

- uncultured_bacterium;_JQ696889.1 100%
- uncultured_Actinomyces_sp.;_FJ976253.1 100%
- uncultured_eubacterium_WCHB1-63;_AF050568.1 100%
- uncultured_Actinomyces_sp.;_FJ976431.1 100%
- uncultured_Actinomyces_sp.;_AM420166.1 100%
- Actinomyces;_NR_025633.1 100%
- Actinomyces_gerencseriae;_NR_029280.1 100%
- Actinomyces_gerencseriae;_X80414.1 100%
- Actinomyces_israelii;_NR_026227.1 100%
- Actinomyces_genomosp._C1;_AY278610.1 100%