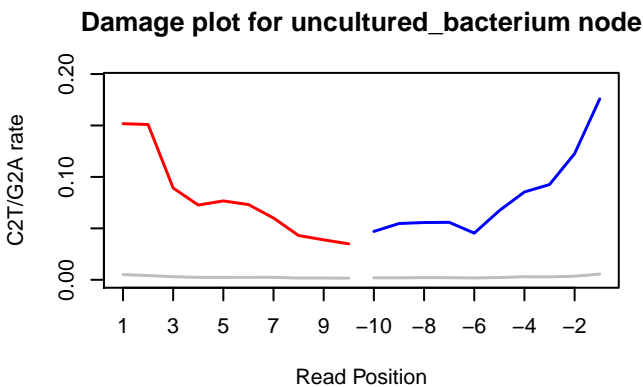
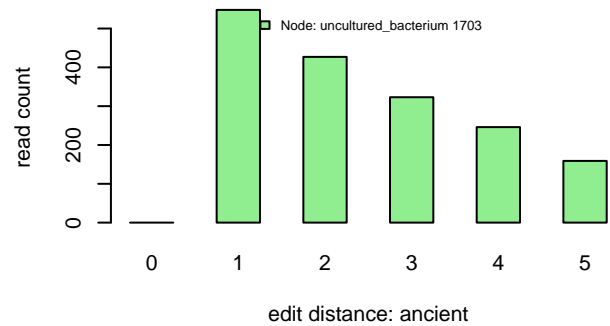
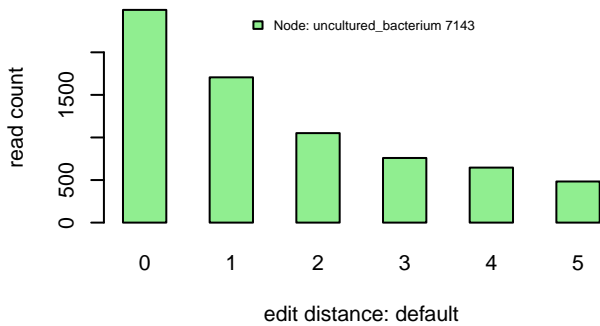


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	8933
all reads	849
nonDup	830
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.1518
G>A_1	0.1758
mean length (sd)	65 (25.057)

uncultured_bacterium;_FM873742.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
uncultured_Actinomyces_sp.;_JQ406543.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
uncultured_Actinomyces_sp.;_FJ976253.1 100%
Clostridiales_bacterium_oral_taxon_G74;_GU432185.1 100%
Actinomyces_genomosp._C1;_AY278610.1 100%
Actinomyces_gerencseriae;_X80414.1 100%