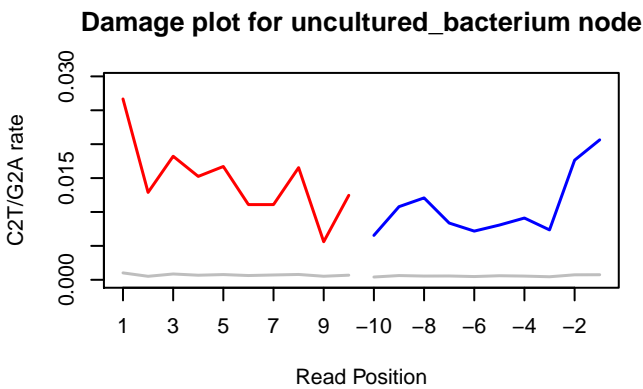
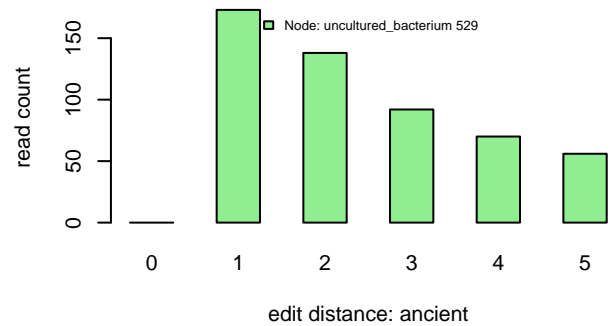
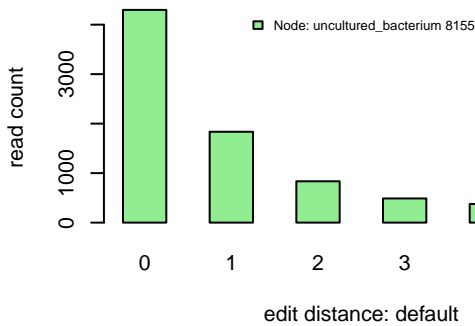


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	39975
all reads	518
nonDup	511
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.0267
G>A_-1	0.0206
mean length (sd)	117 (41.678)

uncultured_bacterium;_HM305213.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Corynebacterium_sp.;_FJ976287.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
uncultured_Corynebacterium_sp.;_AM420019.1 100%
Corynebacterium_matruchotii;_AF543288.1 100%
Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%
uncultured_Corynebacterium_sp.;_AM420027.1 100%
Corynebacterium_matruchotii;_AF543287.1 100%
uncultured_actinobacterium;_DQ424220.1 100%