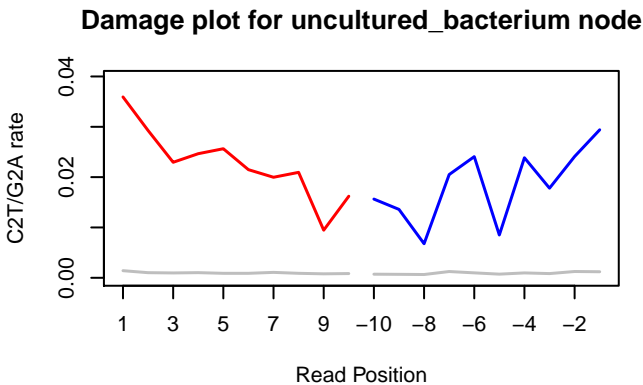
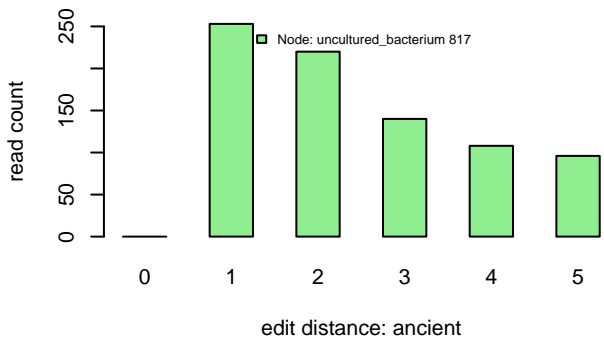
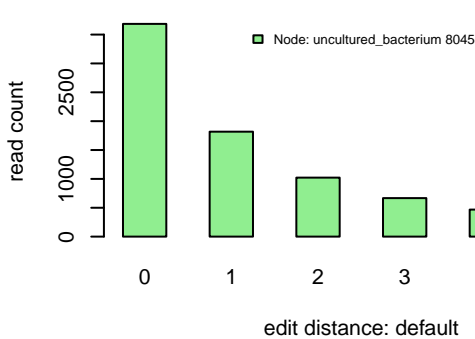


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



edit distance: ancient	
Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	19176
all reads	559
nonDup	539
readDis	0.003
nonStacked	1
destacking?	Off
C>T_1	0.0359
G>A_-1	0.0294
mean length (sd)	109 (40.817)

uncultured_bacterium;_HM273608.1 100%
Actinomyces;_NR_025633.1 100%
uncultured_Corynebacterium_sp.;_FJ976287.1 100%
uncultured_Actinomyces_sp.;_AM420166.1 100%
Corynebacterium_matruchotii;_AF543288.1 100%
actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%
Corynebacterium_matruchotii;_AF543287.1 100%
Bacteria;_FJ976437.1 100%
uncultured_Corynebacterium_sp.;_AM420211.1 100%