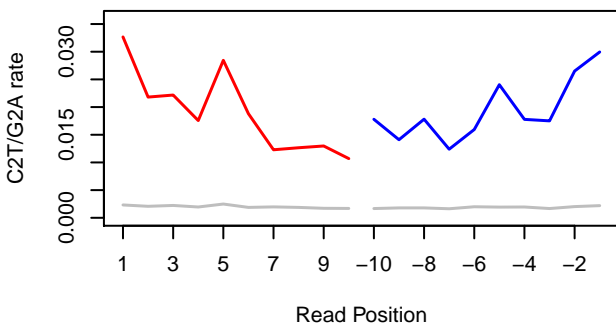
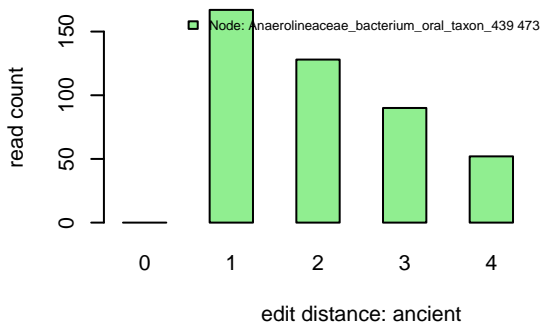
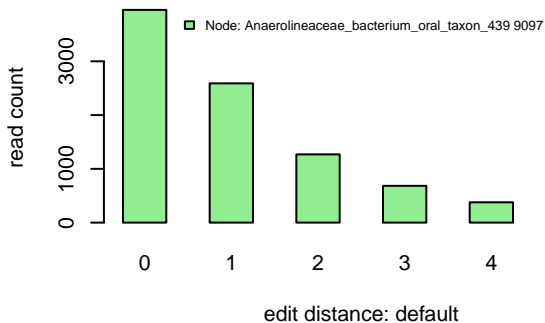


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	95773
all reads	10000
nonDup	9858
readDis	0.677
nonStacked	4696
destacking?	Off
C>T_1	0.0327
G>A_-1	0.03
mean length (sd)	100 (39.625)

[illegible]