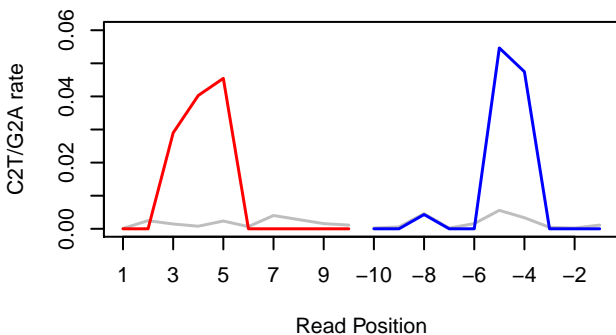
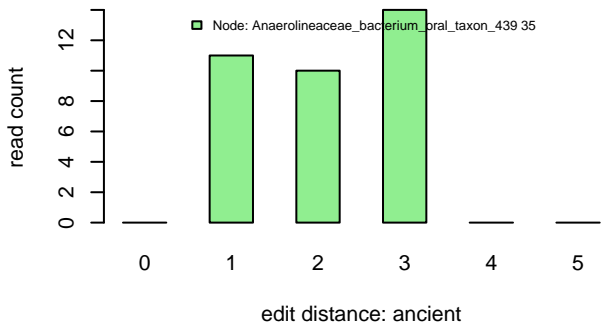
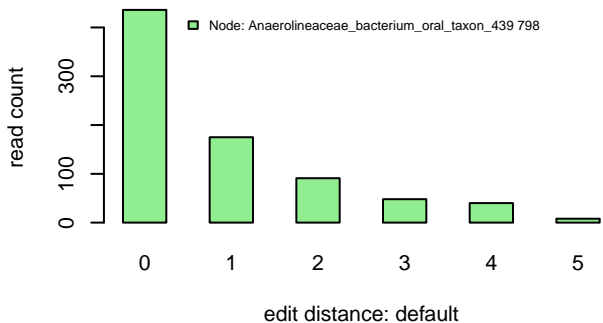


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	835
all reads	835
nonDup	162
readDis	0.22
nonStacked	41
destacking?	Off
C>T_1	0
G>A_-1	0
mean length (sd)	74 (23.542)

[illegible]