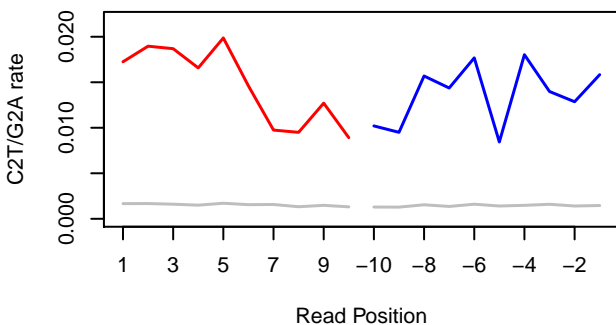
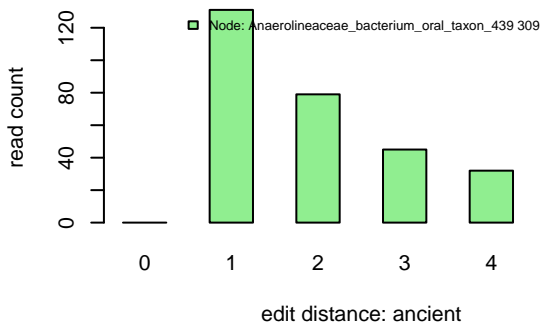
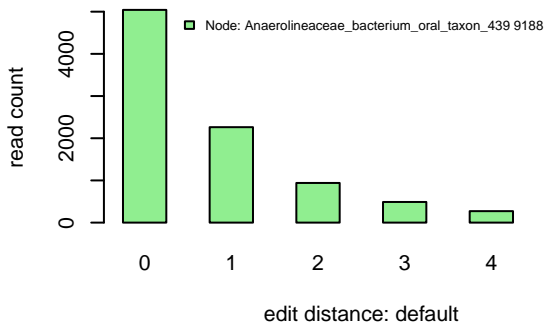


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	381921
all reads	10000
nonDup	9880
readDis	0.677
nonStacked	4621
destacking?	Off
C>T_1	0.0172
G>A_-1	0.0158
mean length (sd)	106 (41.128)

[illegible]