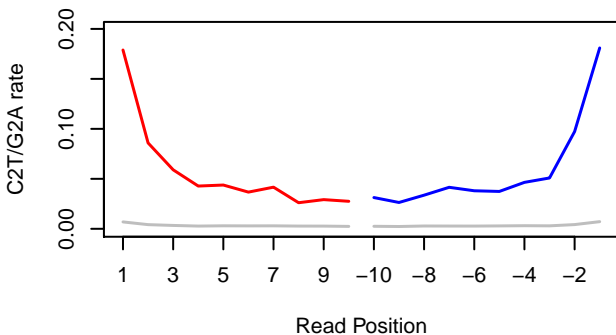
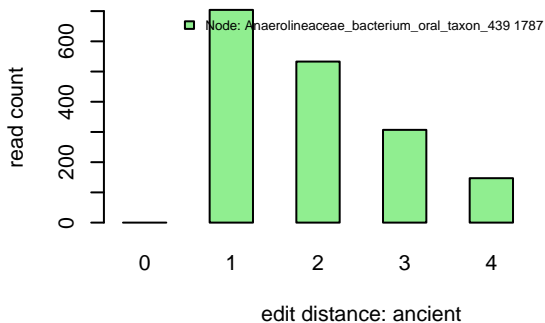
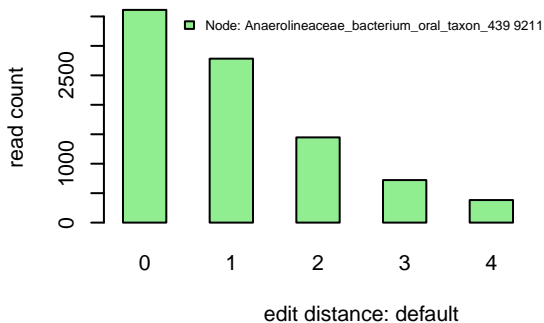


**S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu**



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	99631
all reads	10000
nonDup	9865
readDis	0.781
nonStacked	6151
destacking?	Off
C>T_1	0.1789
G>A_-1	0.1809
mean length (sd)	60 (24.008)

[illegible]