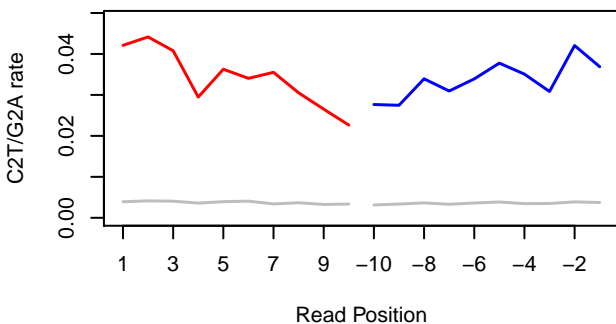
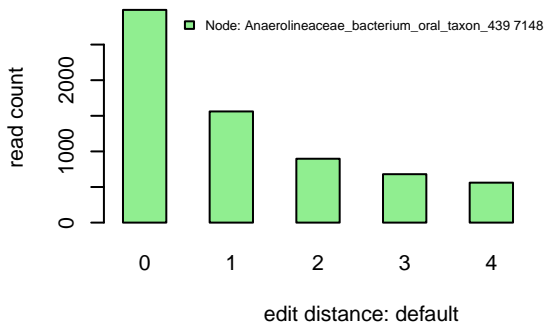


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	165940
all reads	10000
nonDup	9875
readDis	0.66
nonStacked	4491
destacking?	Off
C>T_1	0.0421
G>A_-1	0.0369
mean length (sd)	103 (40.245)

[illegible]