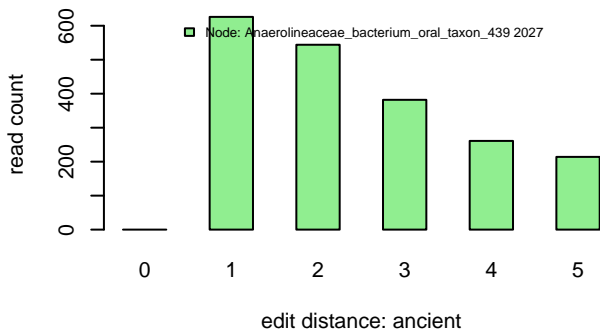
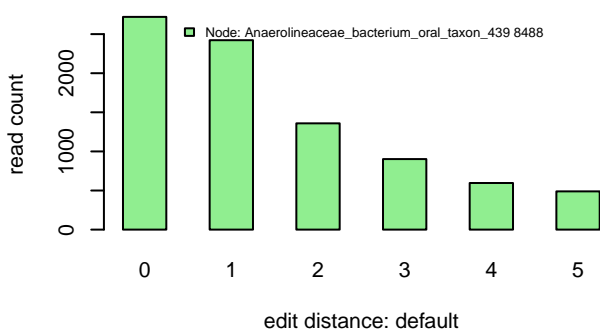
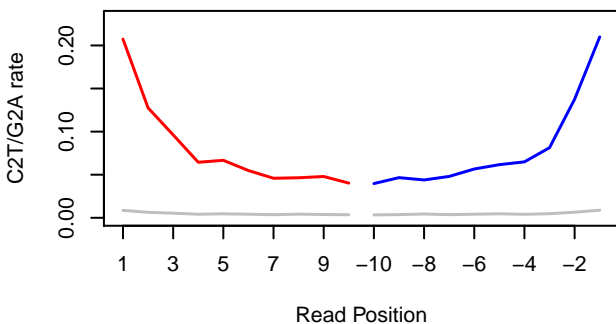


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



age plot for Anaerolineaceae_bacterium_oral_taxon_4



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	275560
all reads	10000
nonDup	9878
readDis	0.726
nonStacked	5581
destacking?	Off
C>T_1	0.2073
G>A_-1	0.2099
mean length (sd)	62 (23.99)

[illegible]