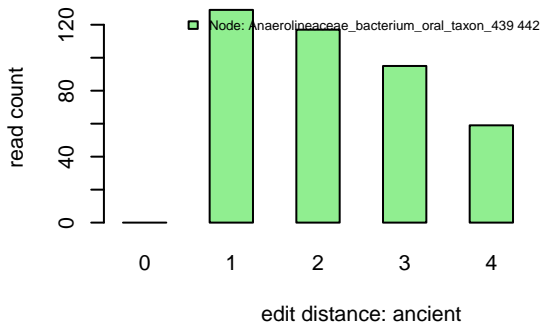
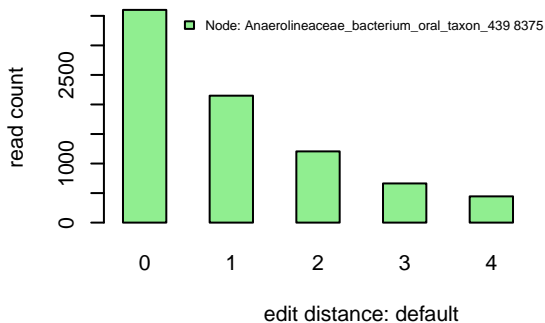
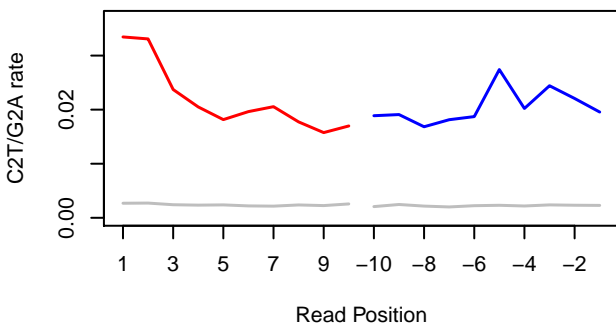


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



age plot for Anaerolineaceae_bacterium_oral_taxon_4



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	491002
all reads	10000
nonDup	9889
readDis	0.646
nonStacked	4237
destacking?	Off
C>T_1	0.0334
G>A_1	0.0196
mean length (sd)	109 (40.914)

[illegible]