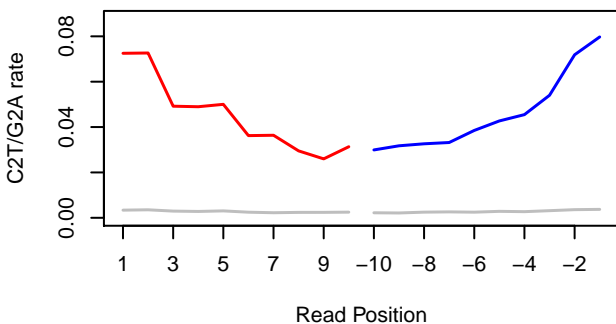
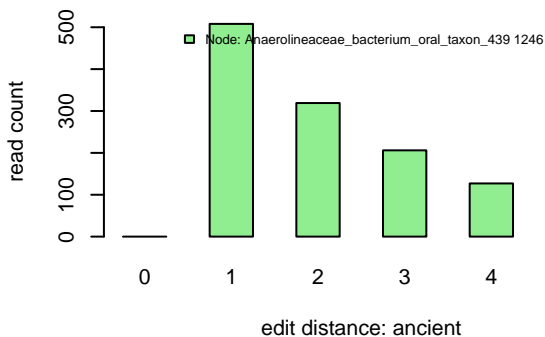
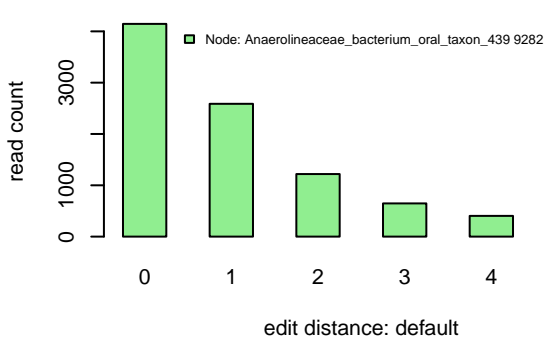


101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	279415
all reads	10000
nonDup	9906
readDis	0.808
nonStacked	6546
destacking?	Off
C>T_1	0.0725
G>A_-1	0.0797
mean length (sd)	58 (22.313)

[illegible]