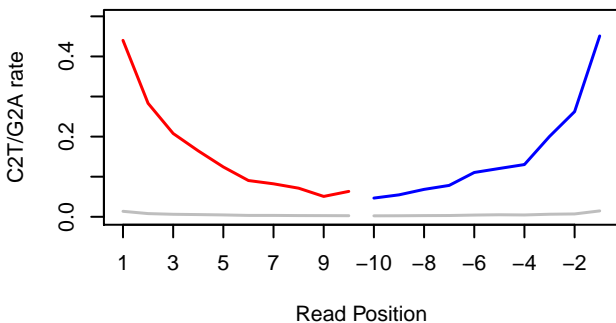
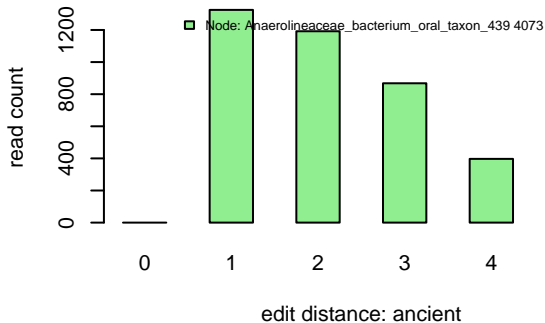
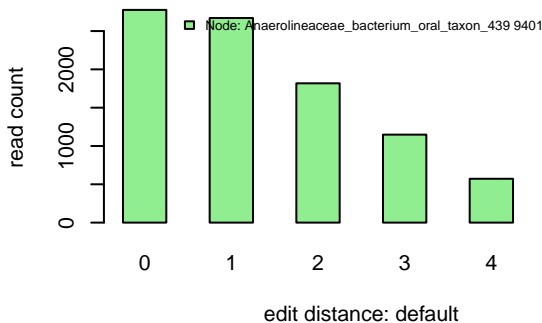


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	142824
all reads	10000
nonDup	5683
readDis	0.755
nonStacked	3992
destacking?	Off
C>T_1	0.4402
G>A_-1	0.4511
mean length (sd)	51 (18.51)

[illegible]