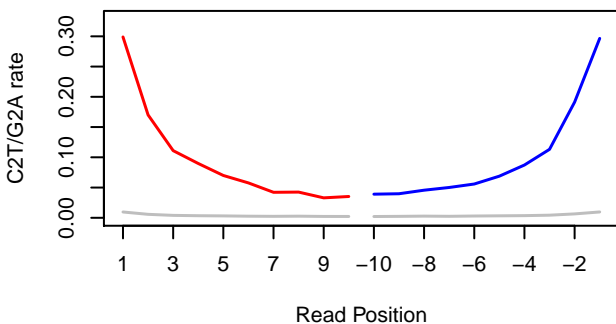
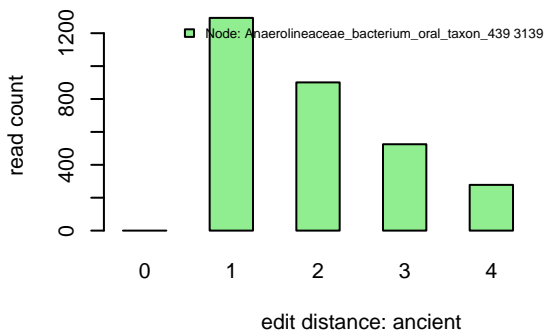
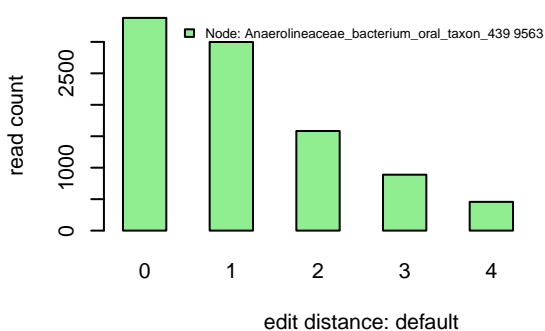


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	212854
all reads	10000
nonDup	9907
readDis	0.831
nonStacked	6918
destacking?	Off
C>T_1	0.299
G>A_-1	0.2966
mean length (sd)	49 (16.102)

[illegible]