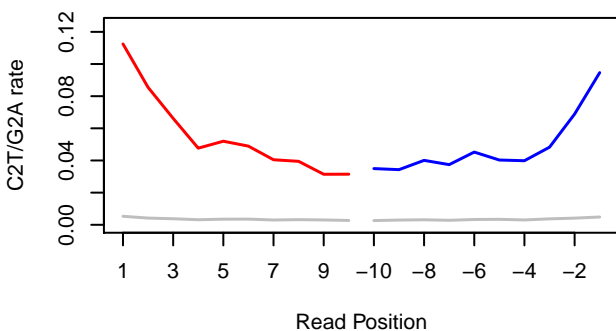
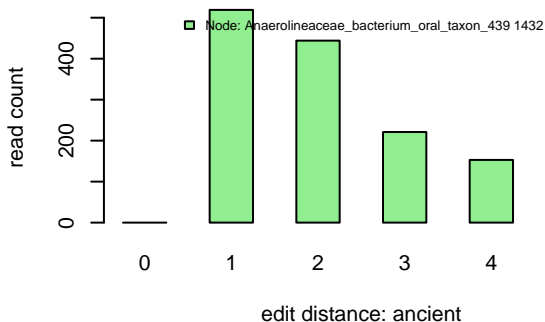
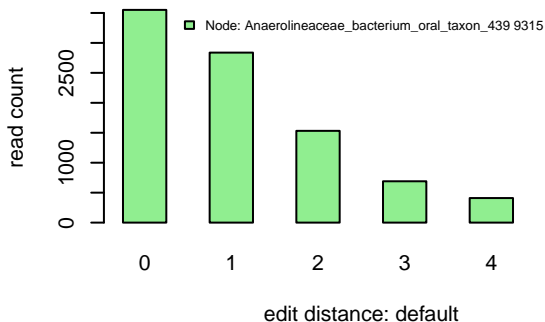


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	37407
all reads	10000
nonDup	9899
readDis	0.8
nonStacked	6432
destacking?	Off
C>T_1	0.1125
G>A_-1	0.0946
mean length (sd)	52 (14.455)

[illegible]