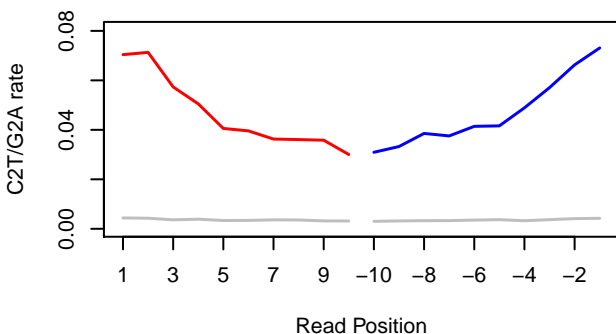
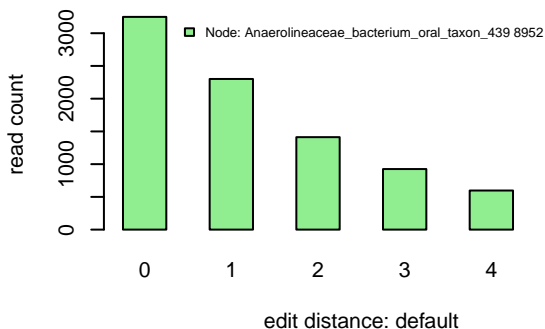


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	280108
all reads	10000
nonDup	9916
readDis	0.804
nonStacked	6475
destacking?	Off
C>T_1	0.0704
G>A_1	0.0731
mean length (sd)	58 (20.568)

[illegible]