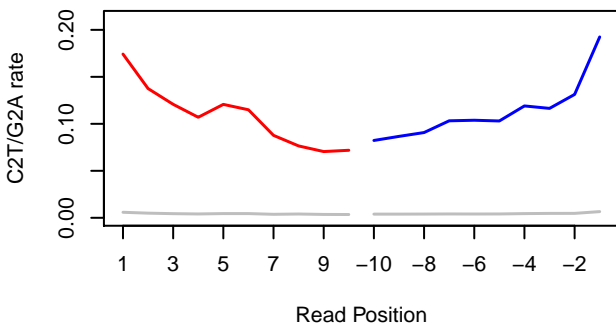
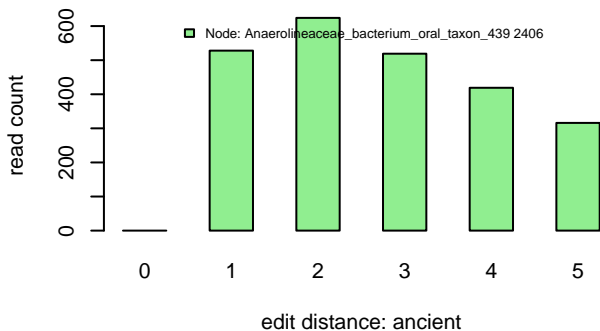
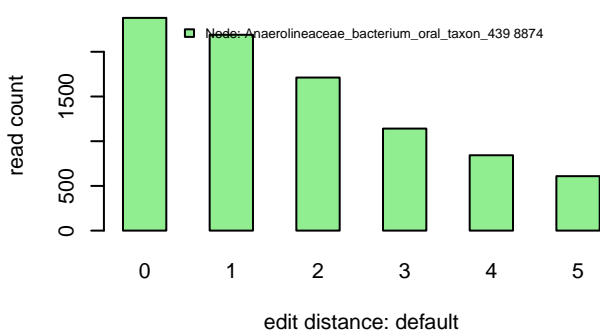


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	242164
all reads	10000
nonDup	9909
readDis	0.794
nonStacked	6297
destacking?	Off
C>T_1	0.1742
G>A_-1	0.1924
mean length (sd)	61 (22.965)

[illegible]