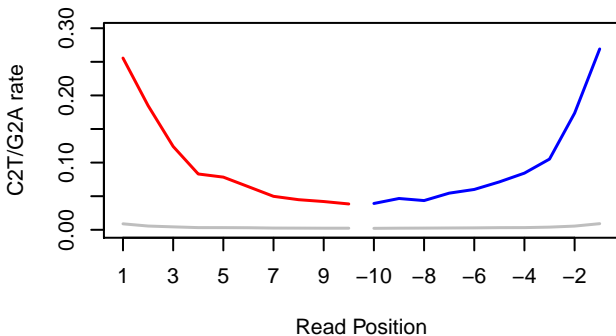
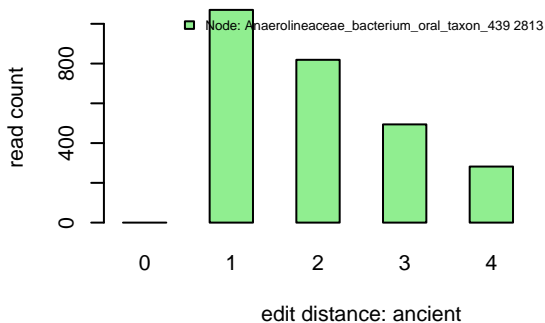
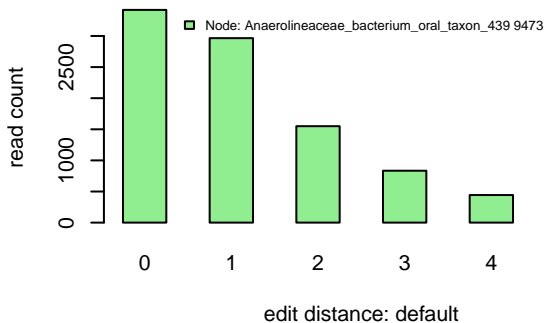


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	12864
all reads	10000
nonDup	9787
readDis	0.818
nonStacked	6625
destacking?	Off
C>T_1	0.2556
G>A_1	0.2692
mean length (sd)	54 (20.439)

[illegible]