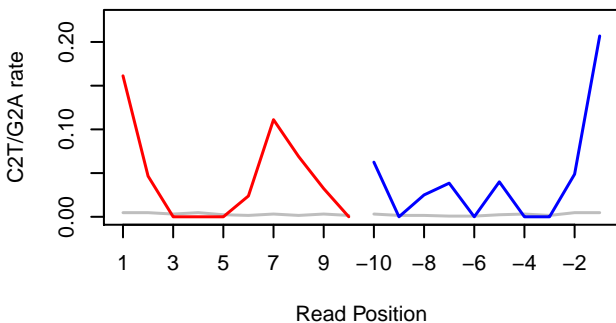
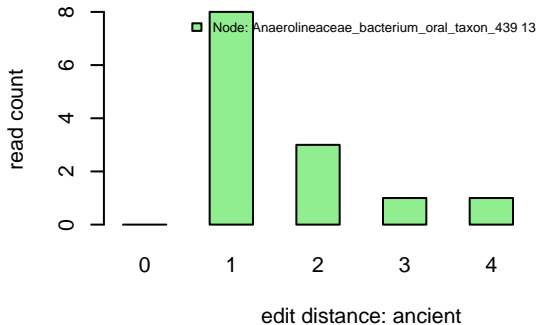
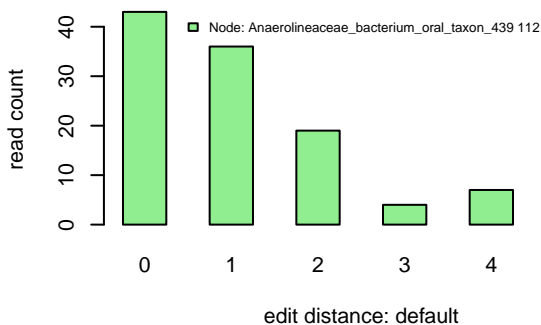


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	121
all reads	121
nonDup	106
readDis	0.899
nonStacked	96
destacking?	Off
C>T_1	0.1613
G>A_-1	0.2069
mean length (sd)	62 (23.105)

[illegible]