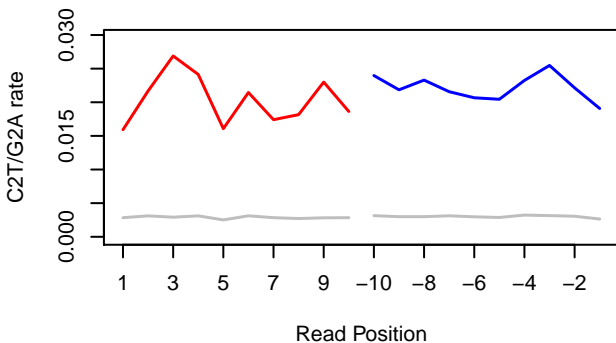
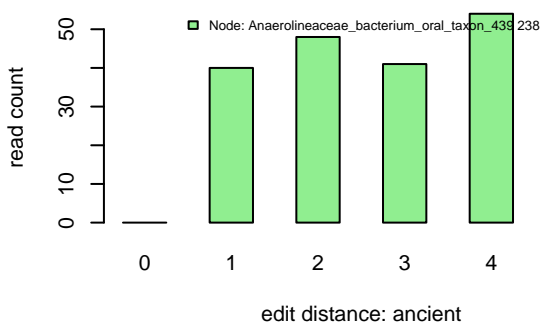
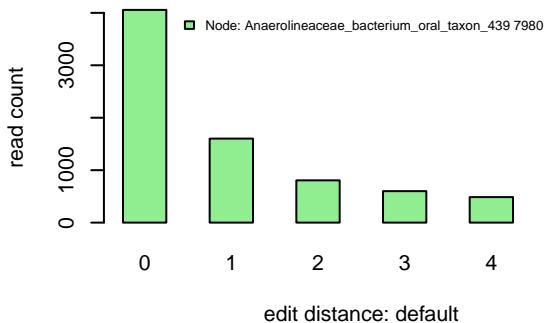


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	63158
all reads	10000
nonDup	9891
readDis	0.72
nonStacked	5229
destacking?	Off
C>T_1	0.0159
G>A_-1	0.0191
mean length (sd)	88 (28.623)

[illegible]