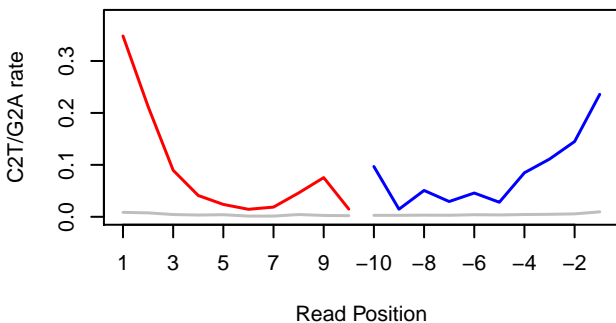
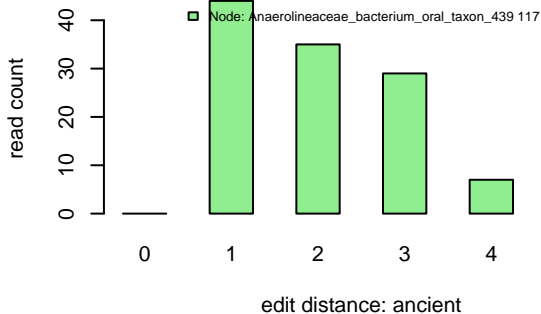
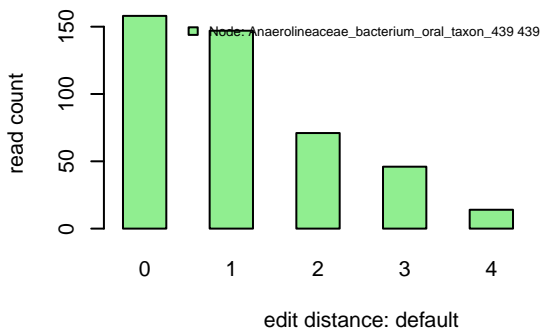


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	458
all reads	458
nonDup	273
readDis	0.889
nonStacked	244
destacking?	Off
C>T_1	0.3482
G>A_-1	0.2357
mean length (sd)	39 (8.499)

[illegible]