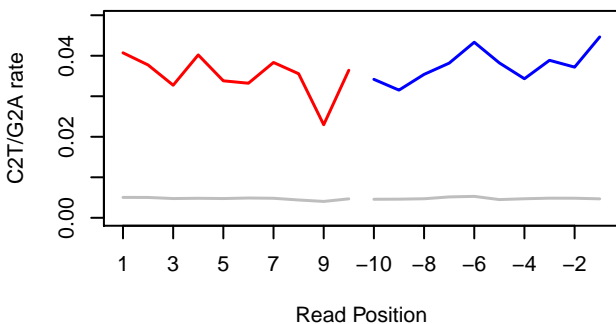
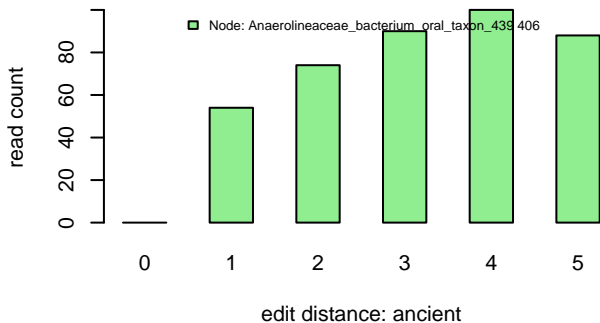
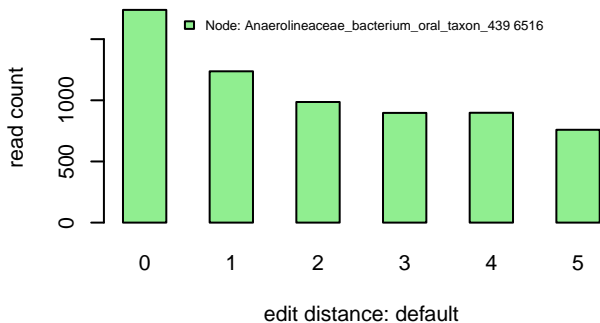


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	305792
all reads	10000
nonDup	9929
readDis	0.679
nonStacked	4726
destacking?	Off
C>T_1	0.0407
G>A_-1	0.0446
mean length (sd)	90 (27.81)

[illegible]