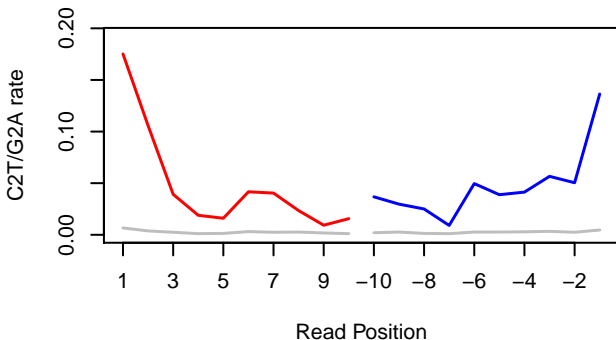
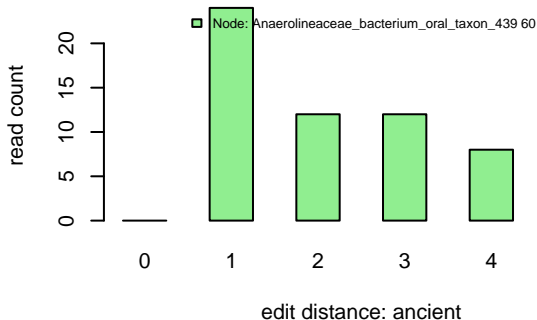
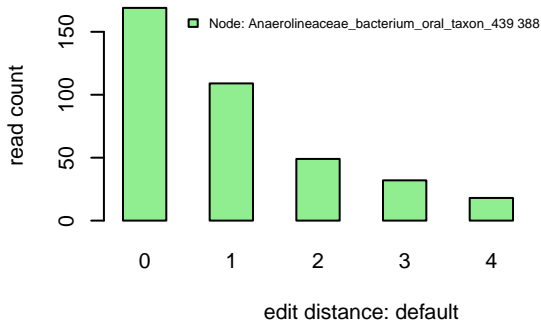


102_S0_L000_R1_000.fastq.merged.prefixed.hg19un)102_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	411
all reads	411
nonDup	390
readDis	0.956
nonStacked	370
destacking?	Off
C>T_1	0.1752
G>A_-1	0.1364
mean length (sd)	66 (13.09)

[illegible]