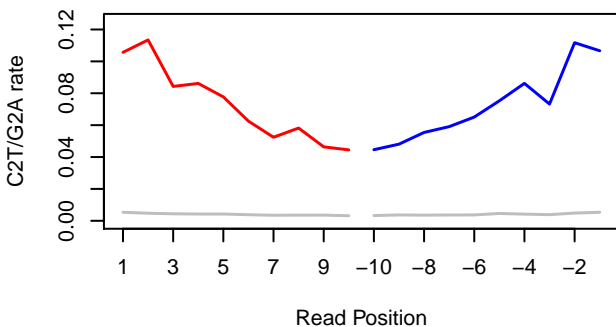
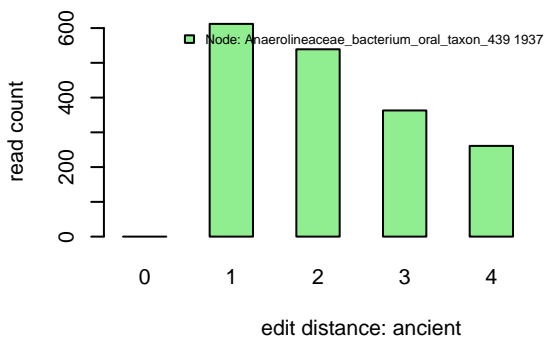
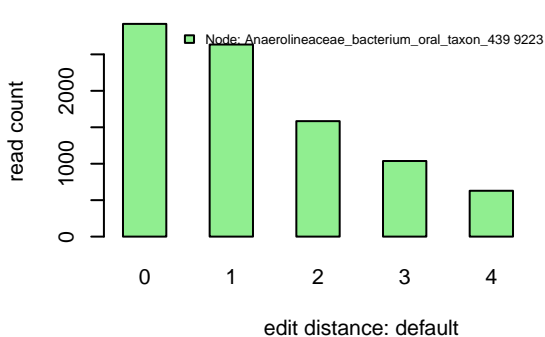


101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	110012
all reads	10000
nonDup	9887
readDis	0.812
nonStacked	6694
destacking?	Off
C>T_1	0.1057
G>A_-1	0.1067
mean length (sd)	50 (16.885)

[illegible]