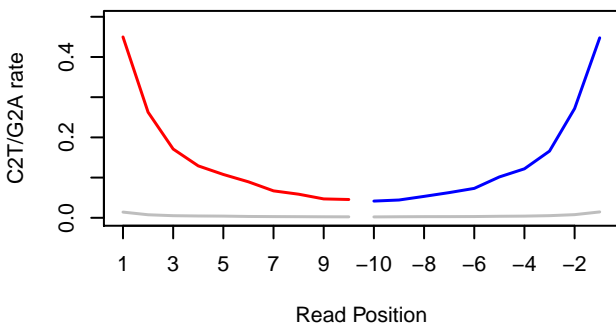
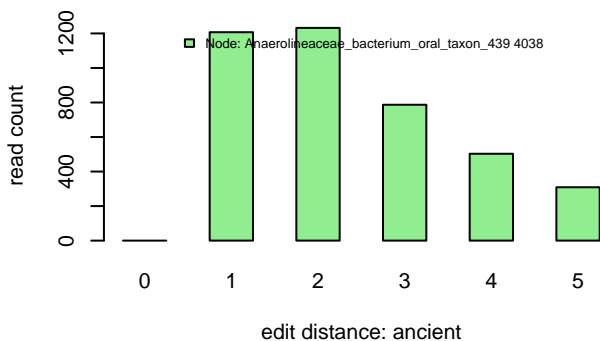
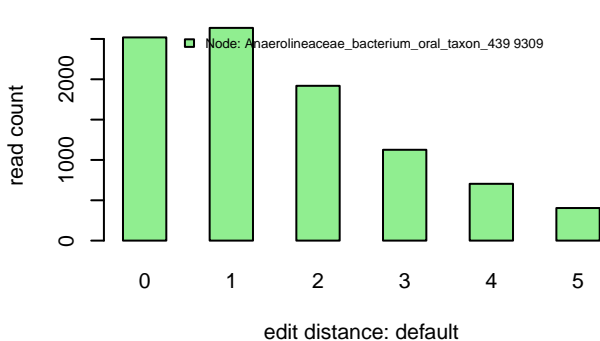


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	60022
all reads	10000
nonDup	8644
readDis	0.736
nonStacked	5119
destacking?	Off
C>T_1	0.4499
G>A_-1	0.4477
mean length (sd)	65 (20.115)

[illegible]