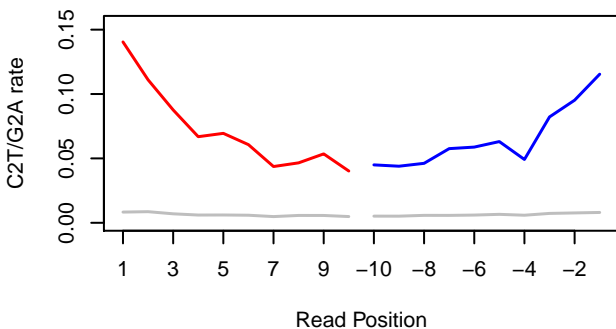
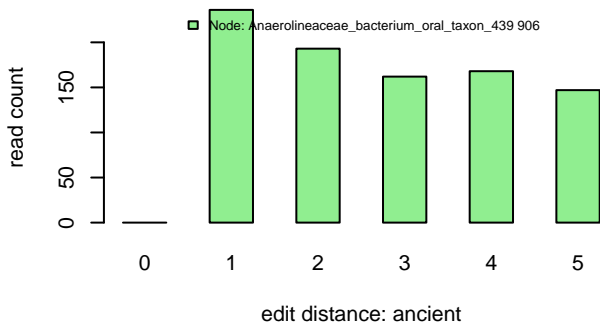
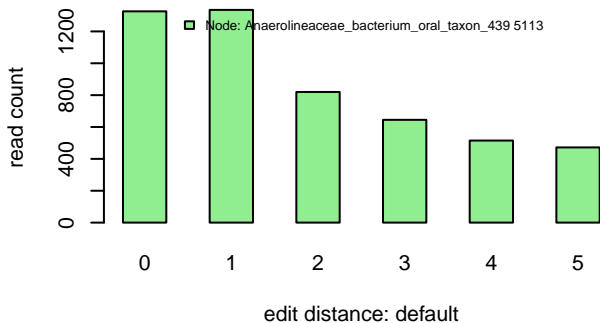


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	6563
all reads	6563
nonDup	6409
readDis	0.665
nonStacked	3460
destacking?	Off
C>T_1	0.1404
G>A_1	0.1154
mean length (sd)	52 (14.182)

[illegible]