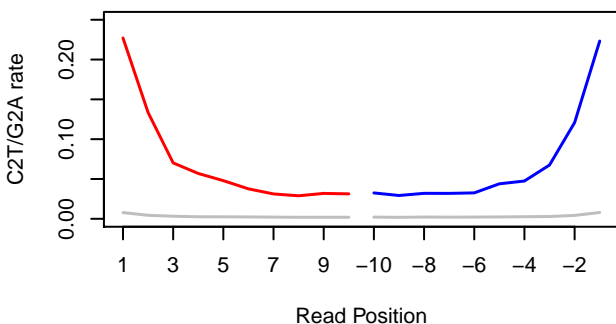
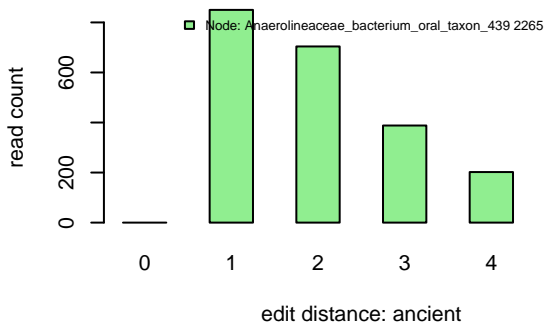
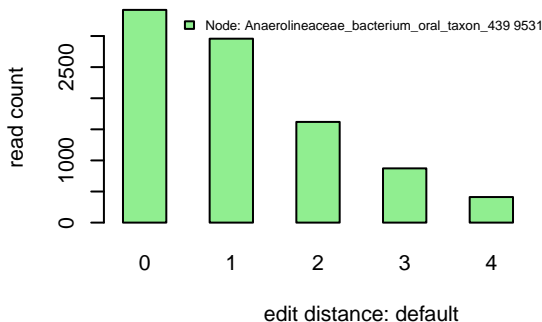


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	1223254
all reads	10000
nonDup	9915
readDis	0.78
nonStacked	6123
destacking?	Off
C>T_1	0.2271
G>A_-1	0.2233
mean length (sd)	65 (25.147)

[illegible]