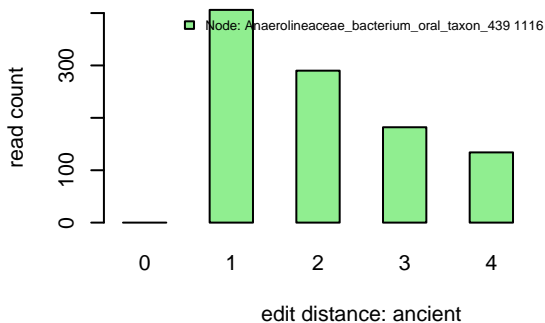
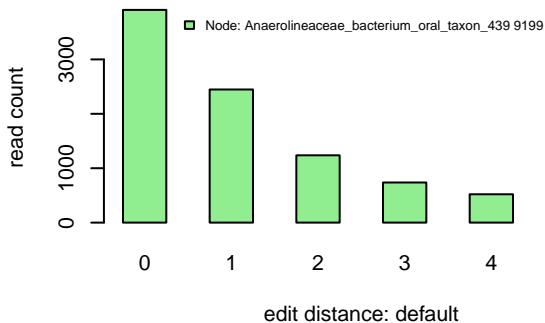
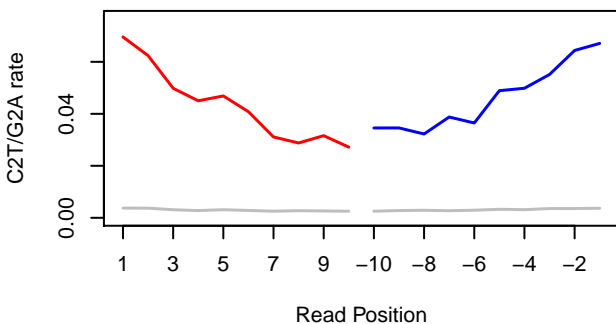


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



age plot for Anaerolineaceae_bacterium_oral_taxon_4



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	366658
all reads	10000
nonDup	9915
readDis	0.802
nonStacked	6519
destacking?	Off
C>T_1	0.0696
G>A_-1	0.0671
mean length (sd)	58 (21.21)

[illegible]