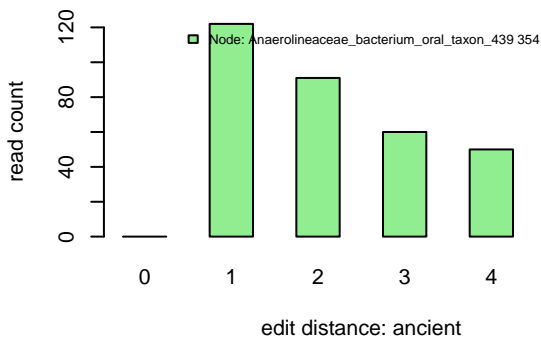
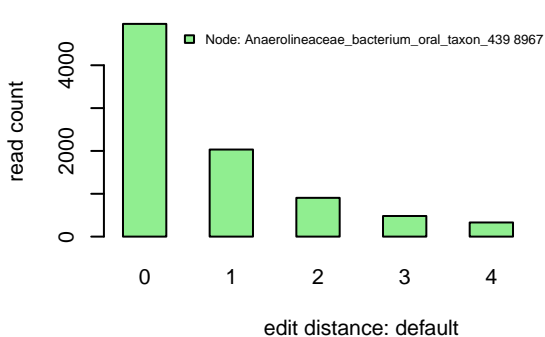
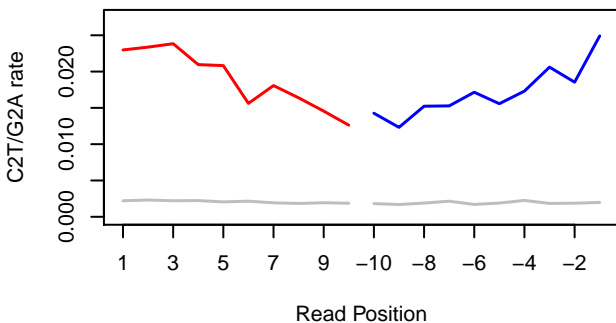


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



age plot for Anaerolineaceae_bacterium_oral_taxon_4



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	1950522
all reads	10000
nonDup	9970
readDis	0.723
nonStacked	5286
destacking?	Off
C>T_1	0.023
G>A_-1	0.0249
mean length (sd)	88 (28.266)

[illegible]