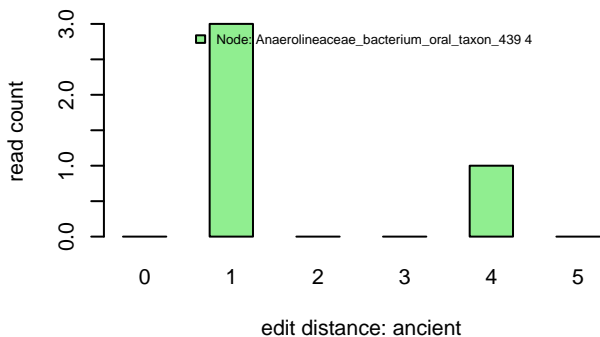
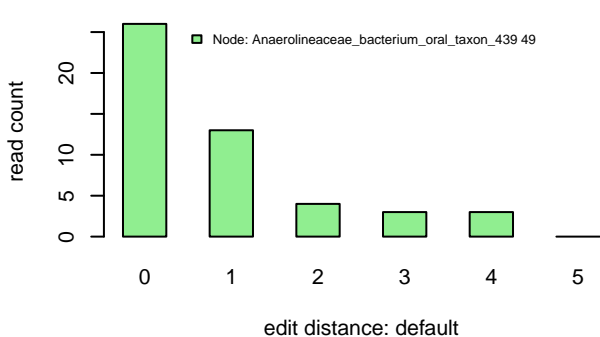
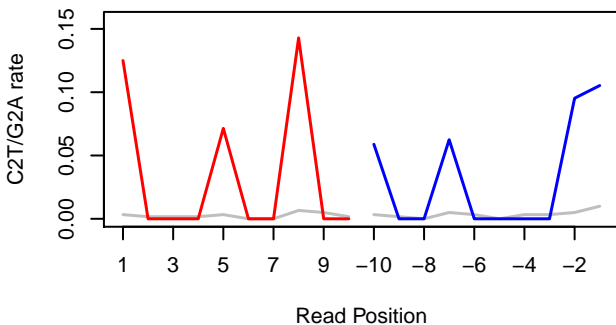


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



age plot for Anaerolineaceae_bacterium_oral_taxon_4



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	55
all reads	55
nonDup	45
readDis	0.806
nonStacked	37
destacking?	Off
C>T_1	0.125
G>A_-1	0.1053
mean length (sd)	55 (19.411)

[illegible]