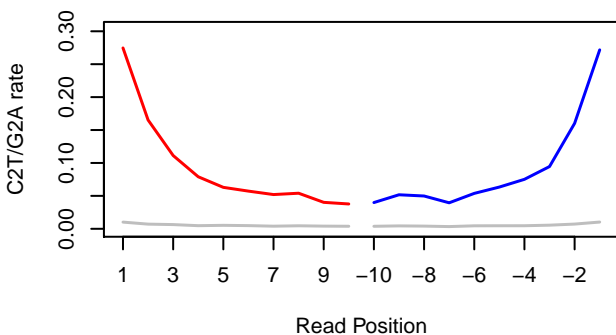
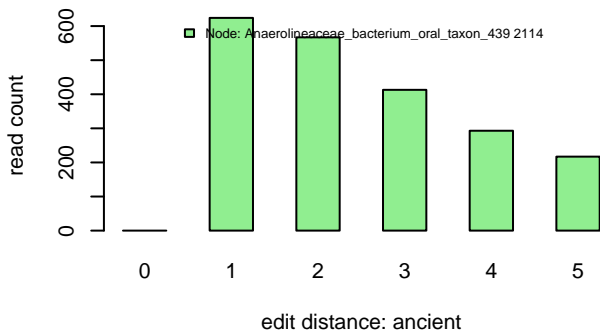
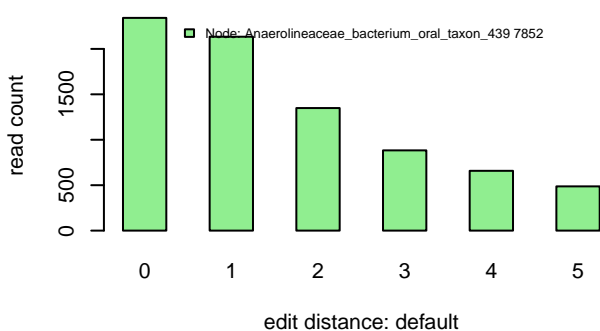


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Node	Anaerolineaceae_bacterium_oral_taxon_439
Top Reference	Anaerolineaceae_bacterium_oral_taxon_439
assigned reads	63087
all reads	10000
nonDup	9901
readDis	0.667
nonStacked	4989
destacking?	Off
C>T_1	0.2747
G>A_-1	0.2718
mean length (sd)	70 (26.117)

[illegible]