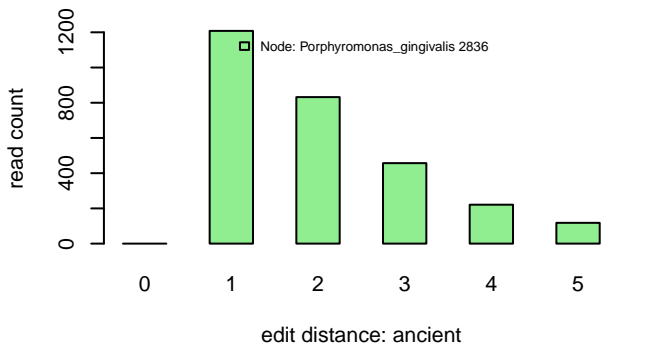
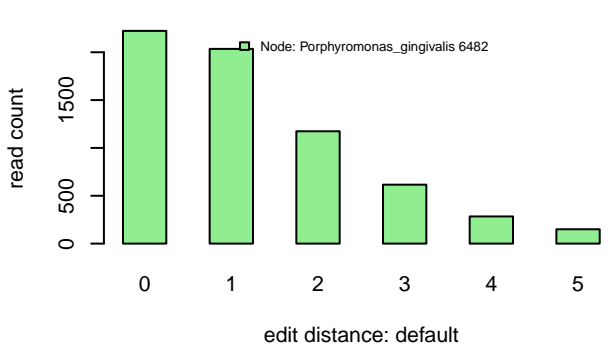
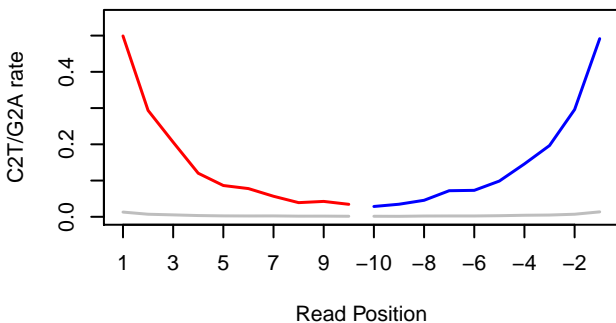


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_TDC60
assigned reads	6658
all reads	4403
nonDup	3782
readDis	0.586
nonStacked	1882
destacking?	Off
C>T_1	0.499
G>A_1	0.4915
mean length (sd)	43 (11.916)

Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis;_U77885.1 005%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 005%