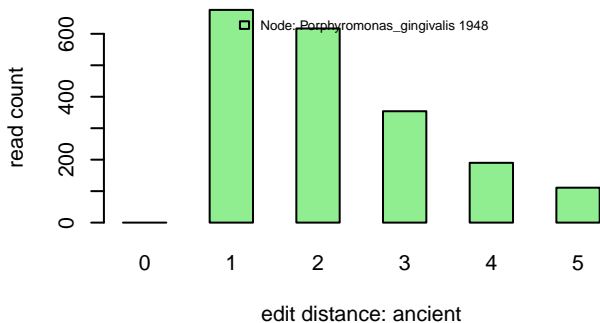
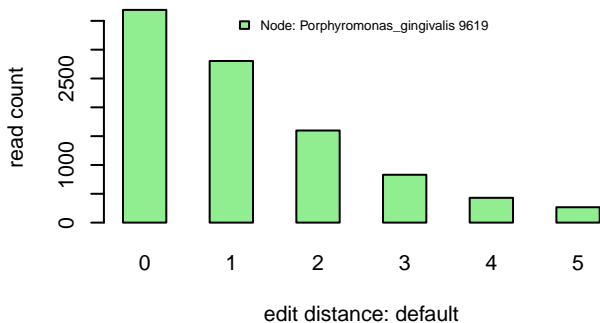
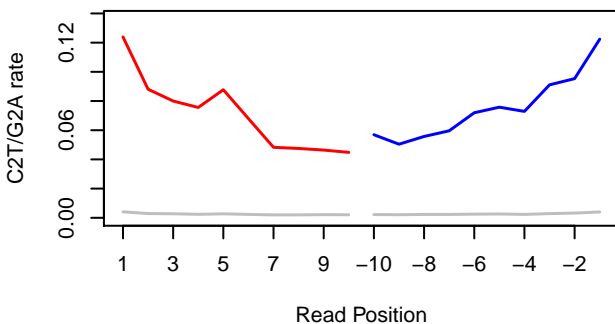


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	26087
all reads	7128
nonDup	7017
readDis	0.66
nonStacked	3744
destacking?	Off
C>T_1	0.1239
G>A_-1	0.1223
mean length (sd)	54 (19.948)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 010%
Porphyromonas_gingivalis_TDC60;_AB757136.1 008%
Tannerella_forsythia_92A2;_CP003191.1 001%