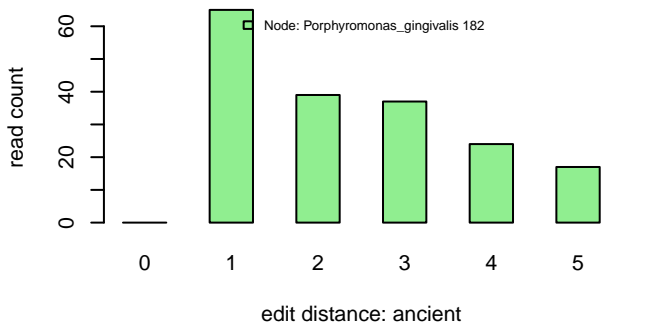
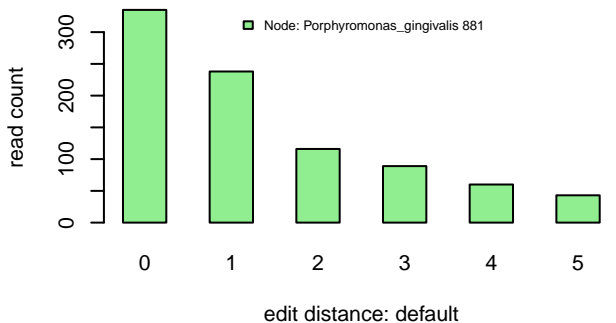
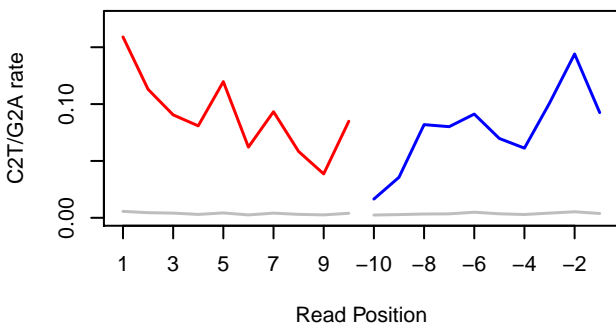


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_AJW4
assigned reads	961
all reads	518
nonDup	504
readDis	0.951
nonStacked	464
destacking?	Off
C>T_1	0.159
G>A_-1	0.0925
mean length (sd)	50 (17.373)

Porphyromonas_gingivalis_AJW4;_CP011996.1	100%
Porphyromonas_gingivalis_TDC60;_AP012203.1	100%
Porphyromonas_gingivalis;_CP007756.1	098%
Porphyromonas_gingivalis;_CP011995.1	097%
Porphyromonas_gingivalis_W83;_AE015924.1	092%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1	090%
Porphyromonas_gingivalis_381;_CP012889.1	090%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1	089%
Porphyromonas_gingivalis;_NR_076194.1	001%
Porphyromonas_gingivalis;_AB757234.1	001%