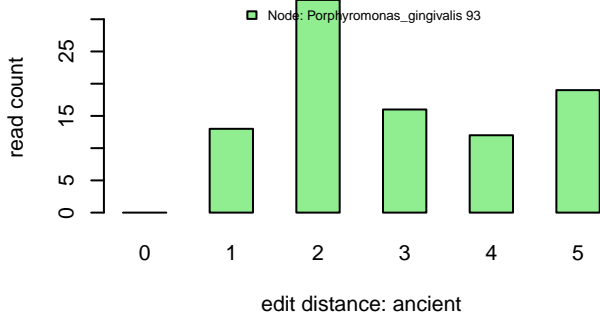
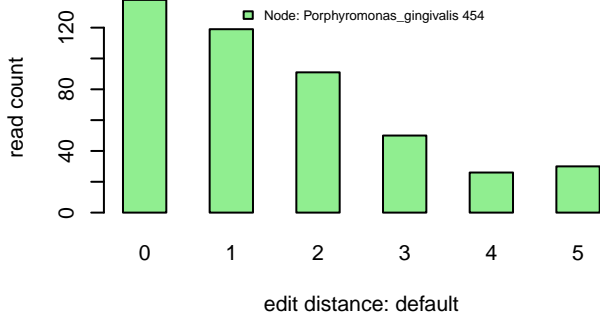
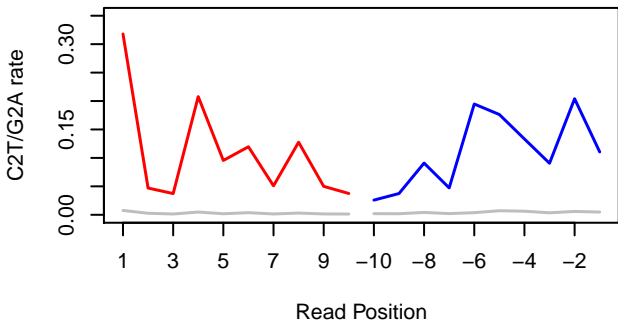


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_AJW4
assigned reads	494
all reads	342
nonDup	202
readDis	0.562
nonStacked	127
destacking?	Off
C>T_1	0.318
G>A_-1	0.1105
mean length (sd)	57 (19.502)

Porphyromonas_gingivalis_AJW4;_CP011996.1	100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1	100%
Porphyromonas_gingivalis_TDC60;_AP012203.1	100%
Porphyromonas_gingivalis;_CP007756.1	100%
Porphyromonas_gingivalis;_CP011995.1	100%
Porphyromonas_gingivalis_W83;_AE015924.1	100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1	100%
Porphyromonas_gingivalis_381;_CP012889.1	100%
Porphyromonas_gingivalis;_U54691.1	016%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1	015%