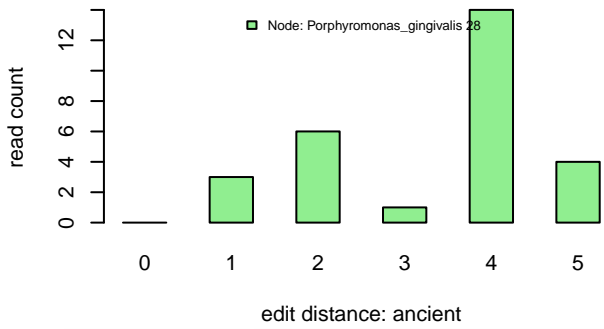
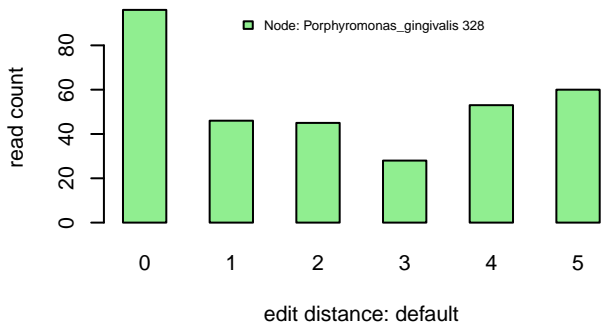
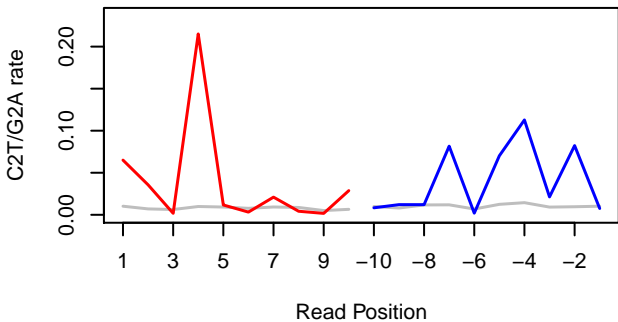


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_TDC60
assigned reads	543
all reads	291
nonDup	187
readDis	0.339
nonStacked	63
destacking?	Off
C>T_1	0.065
G>A_1	0.0075
mean length (sd)	65 (25.58)

Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis_ATCC_33277;_NR_076276.1 065%
Porphyromonas_gingivalis;_NR_076194.1 061%