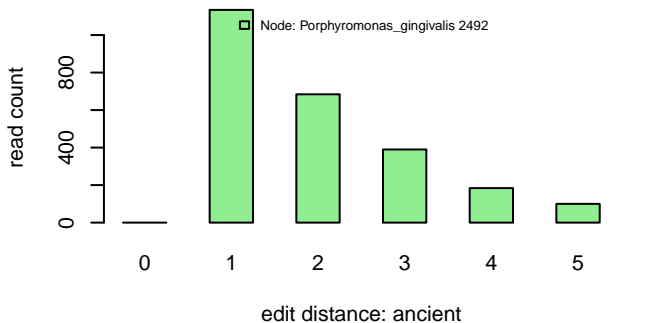
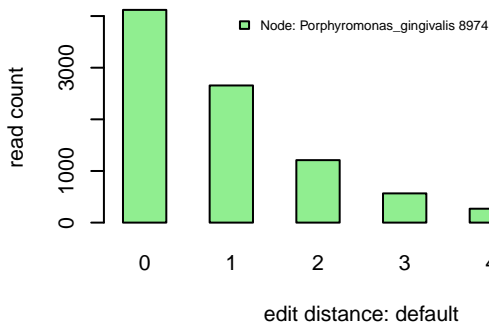
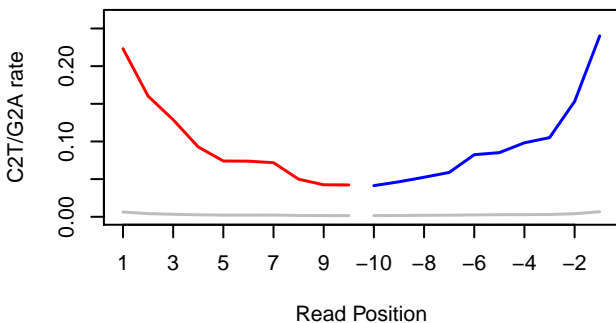


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	9178
all reads	6276
nonDup	6152
readDis	0.751
nonStacked	3898
destacking?	Off
C>T_1	0.2232
G>A_-1	0.2401
mean length (sd)	45 (13.25)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Tannerella_forsythia_92A2;_CP003191.1 002%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 001%
Tannerella_forsythia_KS16;_AP013045.1 000%