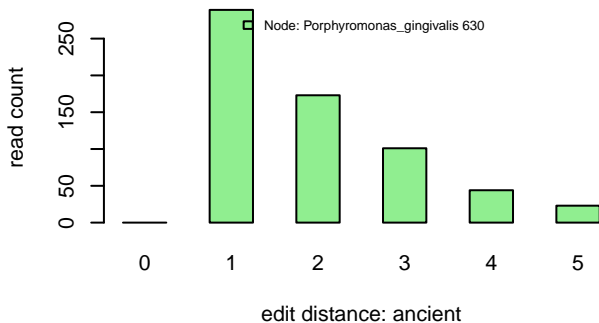
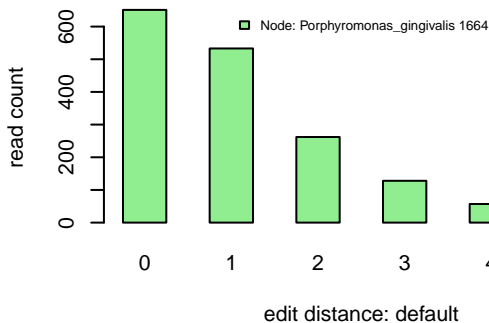
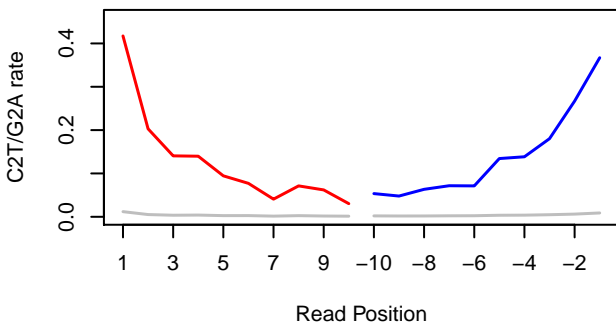


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	1710
all reads	1126
nonDup	1101
readDis	0.898
nonStacked	914
destacking?	Off
C>T_1	0.4173
G>A_-1	0.367
mean length (sd)	42 (11.625)

Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 001%
Porphyromonas_gingivalis_TDC60;_AB757412.1 000%
Porphyromonas_gingivalis_TDC60;_AB757136.1 000%