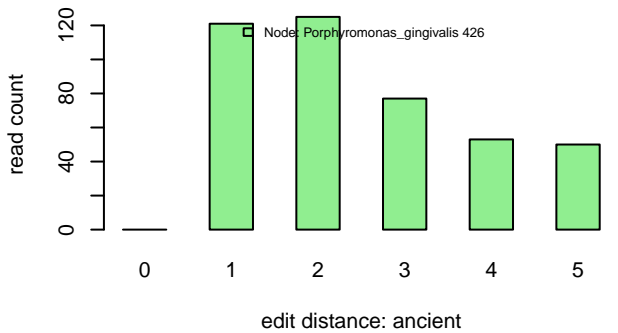
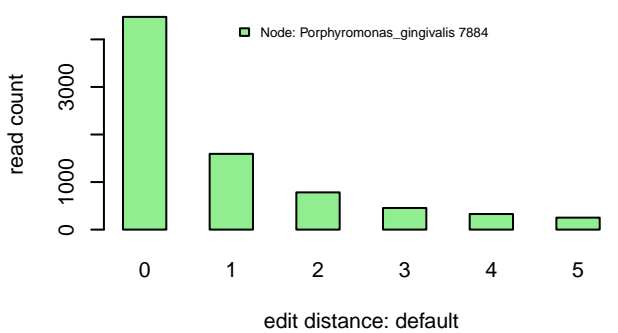
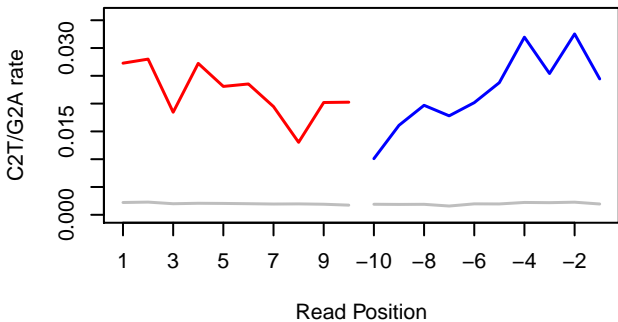


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_381
assigned reads	9039
all reads	5089
nonDup	4973
readDis	0.676
nonStacked	2813
destacking?	Off
C>T_1	0.0273
G>A_1	0.0244
mean length (sd)	89 (36.457)

Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_ATCC_33277;_NR_076276.1 004%
Porphyromonas_gingivalis;_NR_076194.1 003%