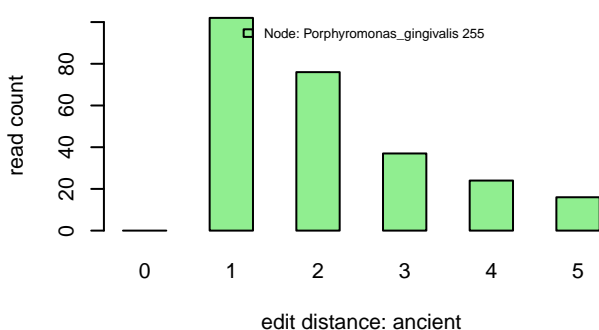
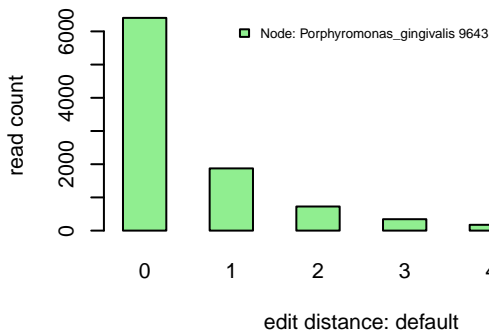
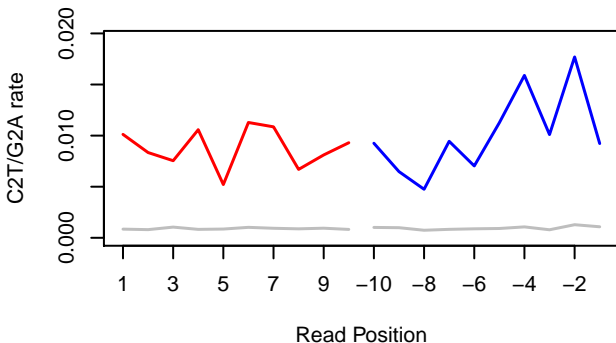


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	124920
all reads	6642
nonDup	6595
readDis	0.7
nonStacked	3522
destacking?	Off
C>T_1	0.0101
G>A_-1	0.0092
mean length (sd)	83 (28.384)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_ATCC_33277;_NR_076276.1 001%
Tannerella_forsythia_92A2;_CP003191.1 001%
Prevotella_dentalis_DSM_3688;_CP003368.1 000%