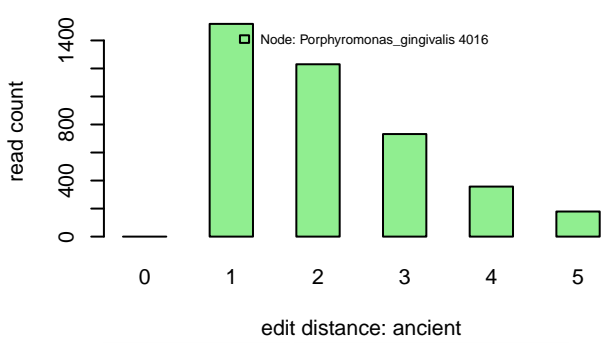
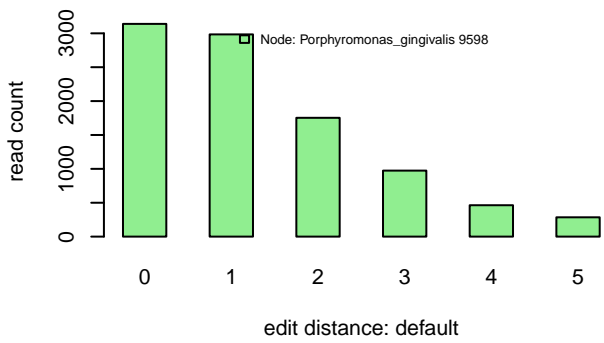
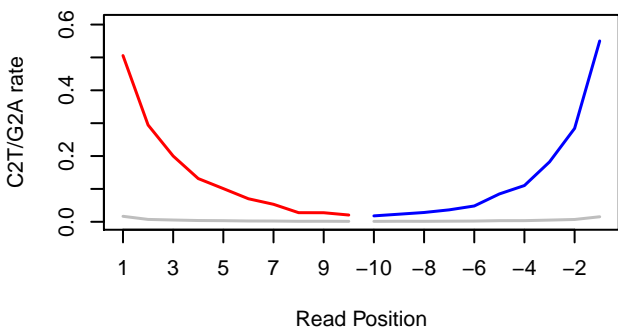


mn1\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19unmn1\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



Damage plot for Porphyromonas\_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	17354
all reads	6576
nonDup	5029
readDis	0.462
nonStacked	1887
destacking?	Off
C>T_1	0.5057
G>A_-1	0.5502
mean length (sd)	51 (11.908)

Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 007%
Porphyromonas_gingivalis_TDC60;_AB757136.1 005%
Tannerella_forsythia_92A2;_CP003191.1 003%