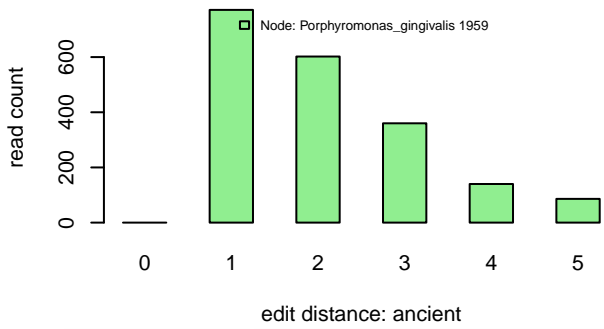
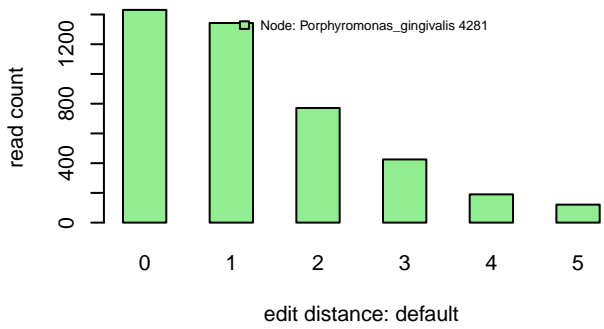
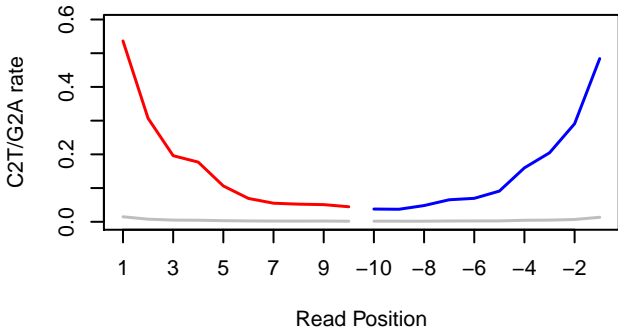


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



Damage plot for Porphyromonas\_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_TDC60
assigned reads	4459
all reads	3076
nonDup	2176
readDis	0.81
nonStacked	1690
destacking?	Off
C>T_1	0.5364
G>A_-1	0.484
mean length (sd)	48 (15.078)

Porphyromonas_gingivalis_TDC60;_AP012203.1	100%
Porphyromonas_gingivalis_AJW4;_CP011996.1	100%
Porphyromonas_gingivalis;_CP011995.1	100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1	100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1	100%
Porphyromonas_gingivalis_W83;_AE015924.1	100%
Porphyromonas_gingivalis_381;_CP012889.1	100%
Porphyromonas_gingivalis;_CP007756.1	100%
Porphyromonas_gingivalis;_AB757234.1	002%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1	002%