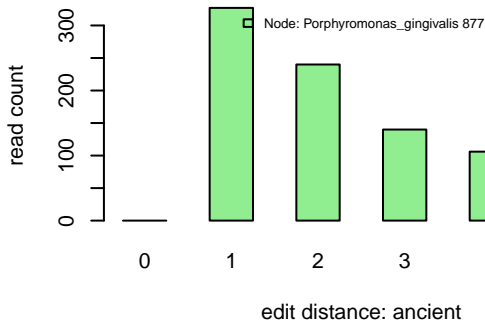
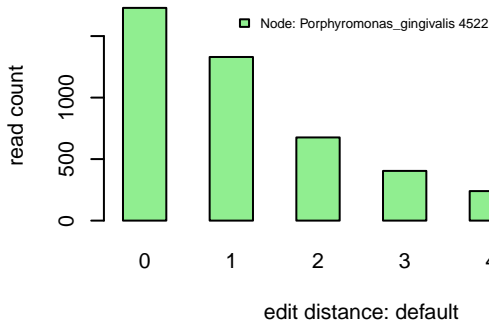
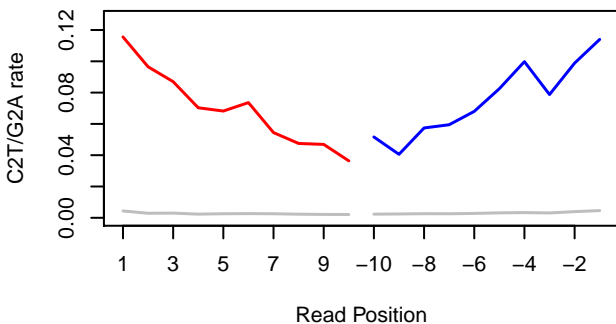


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for Porphyromonas\_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	4861
all reads	2883
nonDup	2826
readDis	0.809
nonStacked	2024
destacking?	Off
C>T_1	0.1156
G>A_-1	0.114
mean length (sd)	57 (20.923)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 001%
Porphyromonas_gingivalis_TDC60;_AB757136.1 001%
Porphyromonas_gingivalis_ATCC_33277;_NR_076276.1 001%