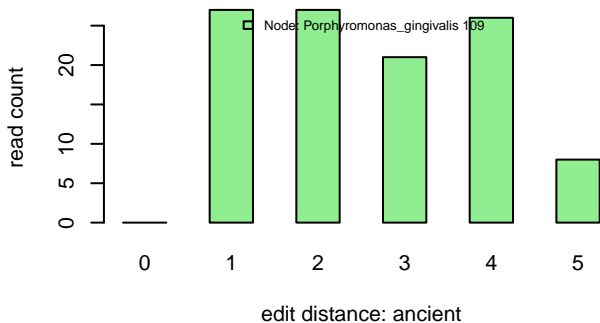
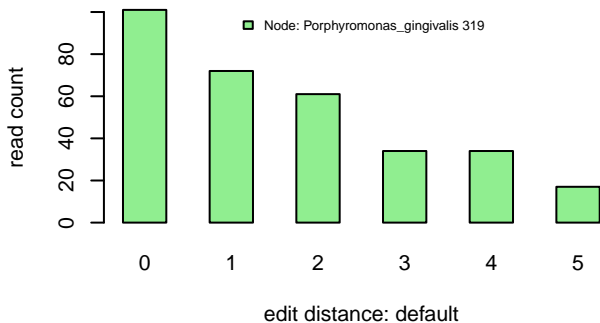
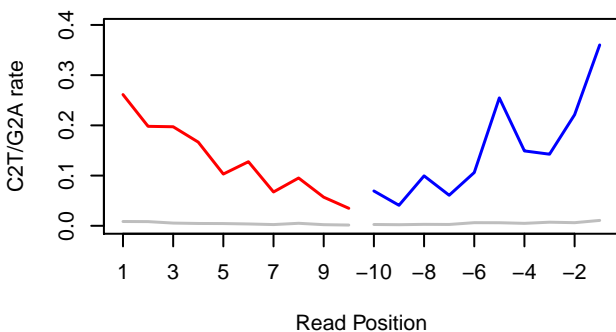


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### Damage plot for Porphyromonas\_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	342
all reads	191
nonDup	185
readDis	0.955
nonStacked	174
destacking?	Off
C>T_1	0.2613
G>A_-1	0.3602
mean length (sd)	46 (11.828)

Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 098%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 095%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 013%
Porphyromonas_gingivalis_TDC60;_AB757136.1 008%
NA;_3ZGZ_B 000%