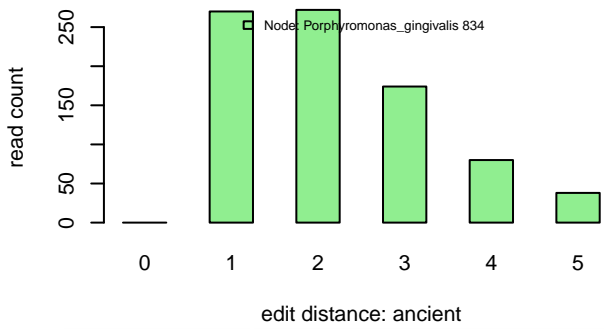
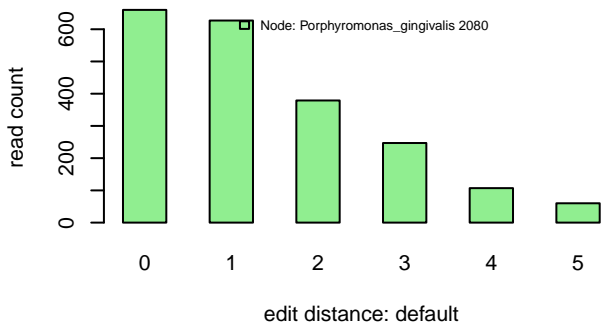
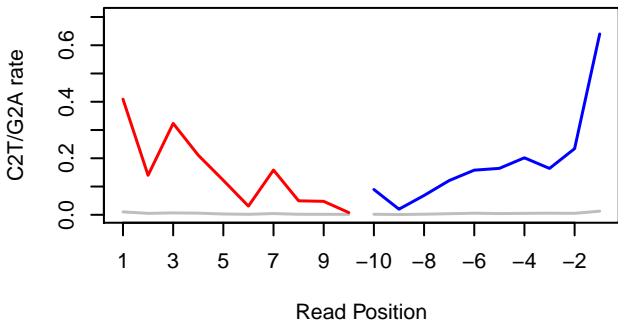


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



Damage plot for Porphyromonas\_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis_TDC60
assigned reads	4198
all reads	2862
nonDup	349
readDis	0.579
nonStacked	204
destacking?	Off
C>T_1	0.4091
G>A_1	0.64
mean length (sd)	44 (12.956)

Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis;_CP007756.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis;_AF155223.1 036%
Porphyromonas_gingivalis;_AJ969093.1 036%