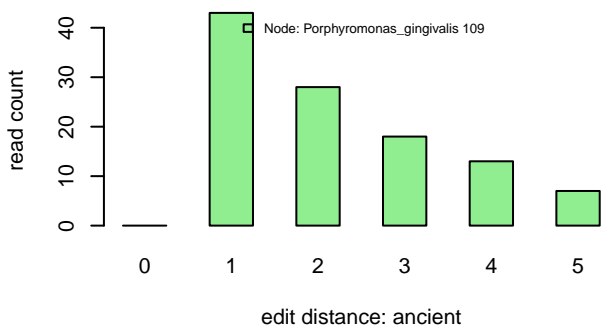
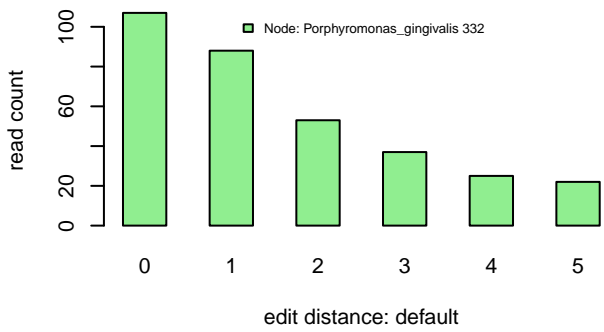
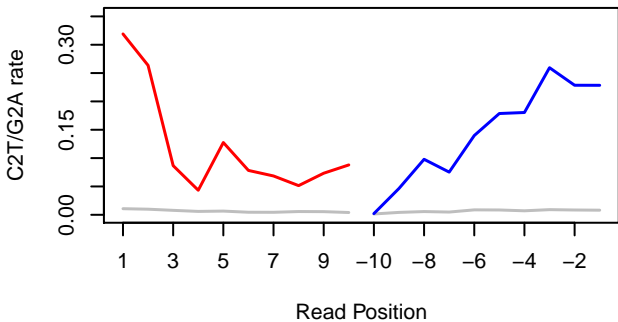


S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu



Damage plot for Porphyromonas\_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	425
all reads	249
nonDup	235
readDis	0.873
nonStacked	204
destacking?	Off
C>T_1	0.319
G>A_-1	0.2284
mean length (sd)	47 (15.393)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_ATCC_33277;_NR_076276.1 006%
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 003%
Tannerella_forsythia_92A2;_CP003191.1 002%