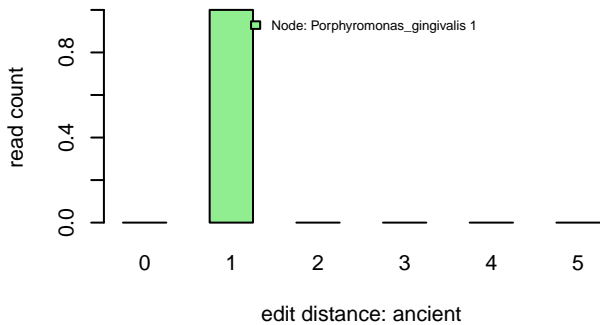
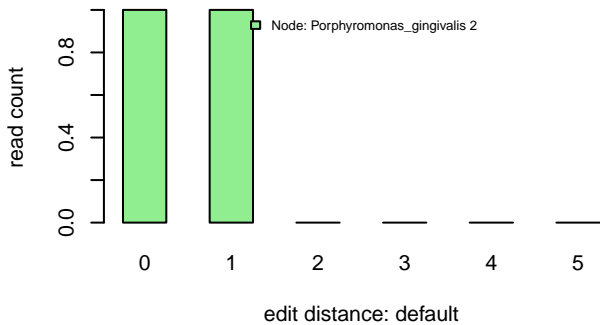
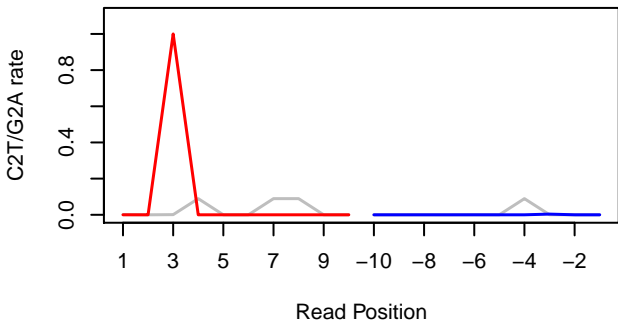


w_S0_L000_R1_000.fastq.merged.prefixed.hg19unmaw_S0_L000_R1_000.fastq.merged.prefixed.hg19unma



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	270
all reads	270
nonDup	10
readDis	0.239
nonStacked	2
destacking?	Off
C>T_1	0
G>A_-1	0
mean length (sd)	49 (1.852)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 050%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 050%
NA%
NA%
NA%