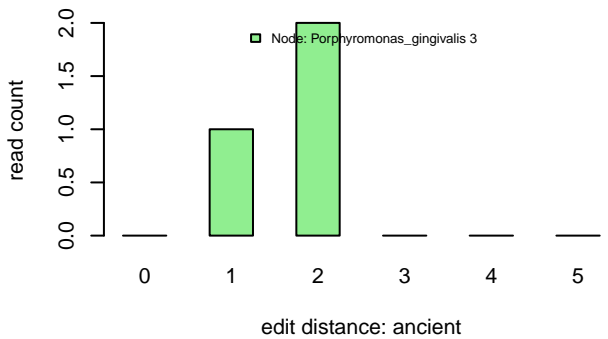
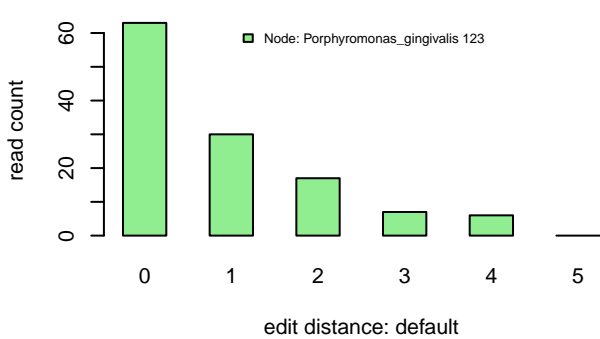
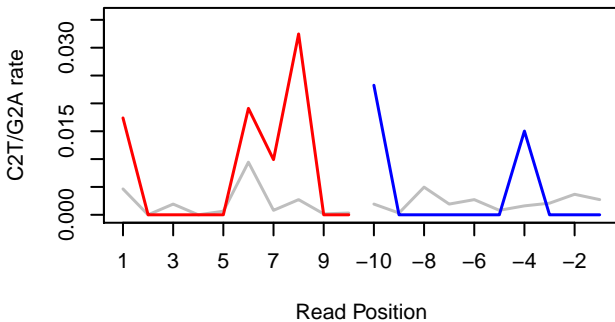


0401_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0401_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Porphyromonas_gingivalis node



Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	135
all reads	80
nonDup	53
readDis	0.64
nonStacked	34
destacking?	Off
C>T_1	0.0174
G>A_-1	0
mean length (sd)	69 (23.79)

Porphyromonas_gingivalis;_CP011995.1 100%
Porphyromonas_gingivalis_W83;_AE015924.1 100%
Porphyromonas_gingivalis_TDC60;_AP012203.1 100%
Porphyromonas_gingivalis_AJW4;_CP011996.1 100%
Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%
Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%
Porphyromonas_gingivalis_381;_CP012889.1 100%
Porphyromonas_gingivalis_W83;_LK012724.1 005%
Proteiniphilum_saccharofermentans;_LT605205.1 005%
Tannerella_forsythia_KS16;_AP013045.1 002%