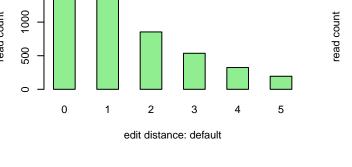
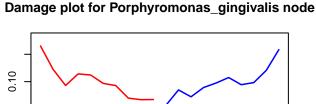
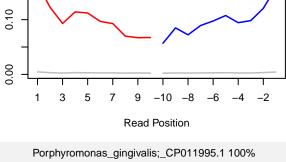


100

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







C2T/G2A rate

Read Position

Porphyromonas\_gingivalis;\_CP011995.1 100%

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_A7A1-28;_CP013131.1 100%

Porphyromonas\_gingivalis\_ATCC\_33277;\_AB015879.1 001%

Tannerella\_forsythia\_92A2;\_CP003191.1 001%

Porphyromonas\_gingivalis\_TDC60;\_AB757136.1 001%

0 1 2 3 4 5

edit distance: ancient

Node Porphyromonas\_gingivalis

Top Reference Porphyromonas\_gingivalis

assigned reads 5143

	3 3 3
Top Reference	Porphyromonas_gingivalis
assigned reads	5143
all reads	3894
nonDup	3545
readDis	0.864
nonStacked	2734
destacking?	Off
C>T_1	0.1652
G>A1	0.1585
mean length (sd)	59 (22.196)