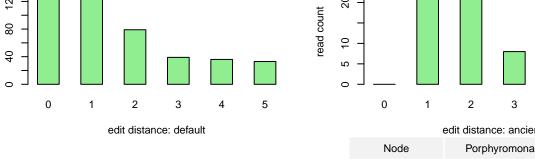
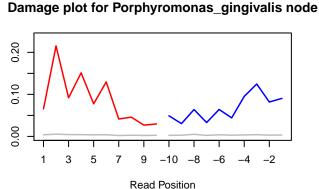
## Porphyromonas\_gingivalis 80 Node: Porphyromonas\_gingivalis 481

0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un)101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un





C2T/G2A rate

Read Position Porphyromonas\_gingivalis\_W83;\_AE015924.1 100% Porphyromonas\_gingivalis;\_CP011995.1 100% Porphyromonas\_gingivalis;\_CP007756.1 100% Porphyromonas\_gingivalis\_A7A1-28;\_CP013131.1 100% Porphyromonas\_gingivalis\_AJW4;\_CP011996.1 100% Porphyromonas\_gingivalis\_TDC60;\_AP012203.1 100%

Porphyromonas\_gingivalis\_ATCC\_33277;\_AP009380.1 044% Porphyromonas\_gingivalis\_381;\_CP012889.1 043% Tannerella\_forsythia\_92A2;\_CP003191.1 007% Porphyromonas\_gingivalis;\_NR\_076194.1 003%

0 1	2	3	4	5
	edit distance: ancient			
Node	Porphyromonas_gingivalis			
Top Reference	Porphyromonas_gingivalis_W83			
assigned reads	543			
all reads	258			
nonDup	251			
readDis	0.681			
nonStacked	140			
destacking?	Off			
C>T_1	0.0655			
G>A1	0.0904			
mean length (sd)	63 (24.18)			