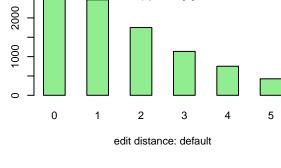
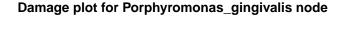
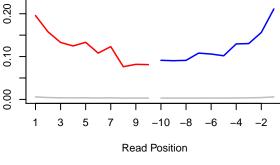


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur







C2T/G2A rate

Porphyromonas_gingivalis_AJW4;_CP011996.1 100%

Porphyromonas_gingivalis_TDC60;_AP012203.1 100% Porphyromonas_gingivalis;_CP011995.1 100%

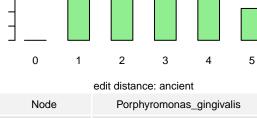
Porphyromonas_gingivalis_W83;_AE015924.1 100%

Porphyromonas_gingivalis;_CP007756.1 100% Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%

Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100%

Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 010%

Porphyromonas_gingivalis_381;_CP012889.1 100% Porphyromonas_gingivalis;_Z35494.1 011%



N	-					
0						
	0 1	2	3	4	5	
	edit distance: ancient					
	Node	Porphyromonas_gingivalis				
	Top Reference	Porphyromonas_gingivalis_AJW4				
	assigned reads		281	03		
	all reads	6297				

all reads	6297
nonDup	6215
readDis	0.409
nonStacked	1936
destacking?	Off

C>T_1 0.1957 G>A_-1 0.2111 57 (21.16) mean length (sd)