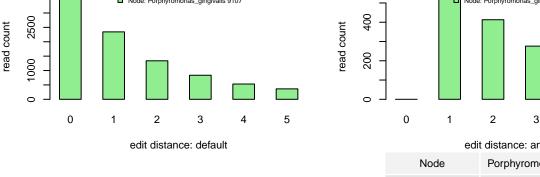
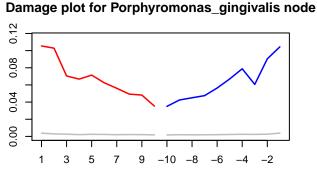
Node: Porphyromonas_gingivalis 9107 Node: Porphyromonas_gingivalis 1588

0102_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0102_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_381;_CP012889.1 100% Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100% Porphyromonas_gingivalis_TDC60;_AP012203.1 100% Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%

edit distance: ancient	
Node	Porphyromonas_gingivalis
Top Reference	Porphyromonas_gingivalis
assigned reads	21230
all reads	6401
nonDup	6295
readDis	0.644
nonStacked	3351
destacking?	Off
C>T_1	0.1054
G>A1	0.1044
mean length (sd)	71 (26.293)

Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 002% Tannerella_forsythia_92A2;_CP003191.1 002%

Porphyromonas_gingivalis_TDC60;_AB757136.1 001%