9 Node: Porphyromonas_gingivalis 853 Node: Porphyromonas_gingivalis 139 300 ead count 4

20

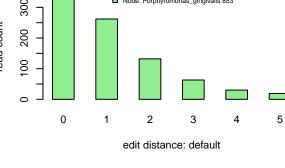
0

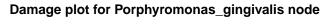
C>T_1

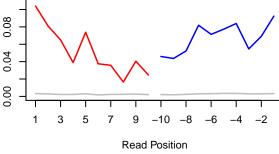
G>A_-1

mean length (sd)

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_W83;_AE015924.1 100%

Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100% Porphyromonas_gingivalis_TDC60;_AP012203.1 100%

Porphyromonas_gingivalis;_CP007756.1 100%

Porphyromonas_gingivalis_381;_CP012889.1 073% Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 073%

Tannerella_forsythia_92A2;_CP003191.1 035%

Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 015%

Porphyromonas_gingivalis_TDC60;_AB757136.1 009%

edit distance: ancient Node Porphyromonas_gingivalis Top Reference Porphyromonas_gingivalis assigned reads 907 all reads 531 nonDup 516 readDis 0.404 nonStacked 122 Off destacking?

2

3

0.1041

0.0924

52 (19.038)