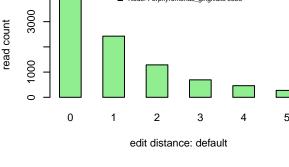
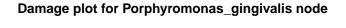
Node: Porphyromonas_gingivalis 9356 Node: Porphyromonas_gingivalis 1882 ead count

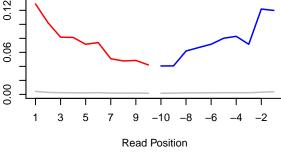
400

200

0102_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0102_S0_L000_R1_000.fastq.merged.prefixed.hg19ur







C2T/G2A rate

Porphyromonas_gingivalis_W83;_AE015924.1 100%

Porphyromonas_gingivalis_AJW4;_CP011996.1 100% Porphyromonas_gingivalis_A7A1-28;_CP013131.1 100%

Porphyromonas_gingivalis;_CP011995.1 100%

Porphyromonas_gingivalis_TDC60;_AP012203.1 100%

Porphyromonas_gingivalis_ATCC_33277;_AP009380.1 100% Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 002%

Porphyromonas_gingivalis_TDC60;_AB757136.1 001%

Porphyromonas_gingivalis_381;_CP012889.1 100%

Tannerella_forsythia_92A2;_CP003191.1 001%

0 2 3 edit distance: ancient Node Porphyromonas_gingivalis Top Reference Porphyromonas_gingivalis assigned reads 36849 all reads 7758 nonDup 7664 readDis 0.686 nonStacked 4231 Off destacking? C>T_1 0.1291 G>A_-1 0.1198

66 (23.93)

mean length (sd)