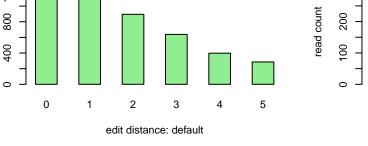
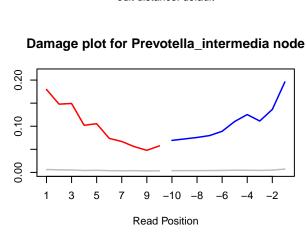


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur





C2T/G2A rate

Prevotella_intermedia;_AP014597.1 100% Prevotella_intermedia_17;_CP003503.1 100%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019300.1 085%

Prevotella_intermedia_17;_CP003502.1 035% revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019301.1 024%

Prevotella_enoeca;_CP013195.1 000%

revotella_intermedia_ATCC_25611_=_DSM_20706;_AB618289.1 000%

Prevotella_fusca_JCM_17724;_CP012075.1 000%

Prevotella_fusca_JCM_17724;_CP012074.1 000%

Porphyromonas_gingivalis;_CP011995.1 000%

0 2 3 edit distance: ancient Node Prevotella_intermedia Top Reference Prevotella_intermedia assigned reads 5654 all reads 2935 nonDup 2881 readDis 0.813 nonStacked 2111 destacking? Off C>T_1 0.1797

0.1959

57 (19.575)

G>A_-1

mean length (sd)