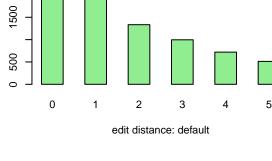
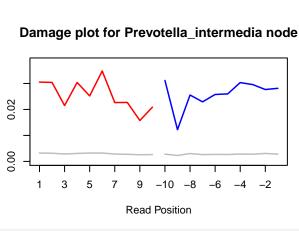
■ Node: Prevotella_intermedia 378 ■ Node: Prevotella_intermedia 7901 80 read count ead count

9 4

20

S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu





C2T/G2A rate

Prevotella_intermedia;_CP019302.1 100% Prevotella_intermedia_17;_CP003503.1 100%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019300.1 100%

Prevotella_intermedia_17;_CP003502.1 056%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019301.1 032% Prevotella_dentalis_DSM_3688;_CP003369.1 001%

Prevotella_fusca_JCM_17724;_CP012074.1 001% Porphyromonas_gingivalis;_CP011995.1 001%

revotella_intermedia_ATCC_25611_=_DSM_20706;_AY734539.1 000%

Tannerella_forsythia_KS16;_AP013045.1 000%

0 2 3 edit distance: ancient Node Prevotella_intermedia Top Reference Prevotella_intermedia assigned reads 47041 all reads 4920

nonDup 4851 readDis 0.557 nonStacked 2000 destacking? Off C>T_1 0.0305 G>A_-1 0.0281

105 (40.634)

mean length (sd)