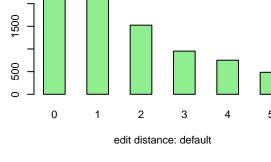
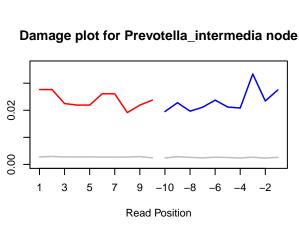
100 ■ Node: Prevotella_intermedia 396 Node: Prevotella_intermedia 8170 read count

ead count 9

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





C2T/G2A rate

Prevotella_intermedia;_AP014597.1 100% Prevotella_intermedia_17;_CP003503.1 100% revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019300.1 096%

Prevotella_intermedia_17;_CP003502.1 051%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019301.1 029%

Porphyromonas_gingivalis;_CP011995.1 001%

Prevotella_dentalis_DSM_3688;_CP003369.1 001% Prevotella_fusca_JCM_17724;_CP012074.1 000%

revotella_intermedia_ATCC_25611_=_DSM_20706;_AB618289.1 000%

Tannerella_forsythia_KS16;_AP013045.1 000%

	0	1	2	3	4	5
	edit distance: ancient					
	Node		F	Prevotella_intermedia		
	Top Reference assigned reads all reads nonDup readDis nonStacked destacking? C>T_1 G>A1			Prevotella_intermedia		
				40599		
				4905 4803		
				0.577		
				2140		
				Off		
				0.0277		
				0.0275		

107 (41.064)

mean length (sd)