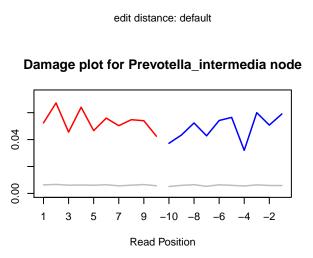


5

0

0

S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu



2

3

0

C2T/G2A rate

Prevotella\_intermedia;\_AP014597.1 100% Prevotella\_intermedia\_17;\_CP003503.1 100% revotella\_intermedia\_ATCC\_25611\_=\_DSM\_20706;\_CP019300.1 100%

Prevotella\_intermedia\_17;\_CP003502.1 064% revotella\_intermedia\_ATCC\_25611\_=\_DSM\_20706;\_CP019301.1 042% Prevotella\_fusca\_JCM\_17724;\_CP012074.1 001%

edit distance: ancient Node Prevotella\_intermedia Top Reference Prevotella\_intermedia assigned reads 60702 all reads 5160 nonDup 5081 readDis 0.412 nonStacked 1460 destacking? Off C>T\_1 0.0524 G>A\_-1 0.059 mean length (sd) 108 (40.606)

2

3

5

Porphyromonas\_gingivalis;\_CP011995.1 001%

Prevotella\_dentalis\_DSM\_3688;\_CP003369.1 000%

Prevotella\_enoeca;\_CP013195.1 000%

Prevotella\_fusca\_JCM\_17724;\_CP012075.1 000%