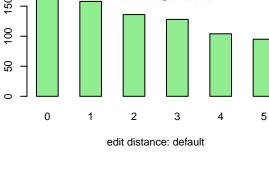
■ Node: Prevotella intermedia 90 Node: Prevotella_intermedia 798 150

read count

10

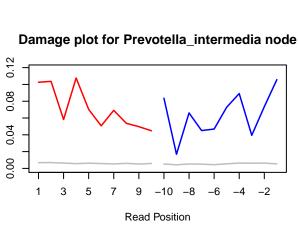
Ω

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



read count

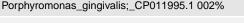
C2T/G2A rate



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

	_						
revotella_intermedia_	ATCC_2561	1_=_	DSM_2	20706;_	CP0193	300.1 0	83%
revotella_intermedia_	_ATCC_2561	11_=_l	DSM_2	20706;_	CP0193	801.1 0	55%

Prevotella_intermedia_17;_CP003502.1 023%



Prevotella_fusca_JCM_17724;_CP012074.1 001%

Prevotella_dentalis_DSM_3688;_CP003369.1 001%

Tannerella_forsythia_KS16;_AP013045.1 000%

Prevotella_enoeca;_CP013195.1 000%

0 2 3 5 edit distance: ancient Node Prevotella_intermedia Top Reference Prevotella_intermedia assigned reads 1529 all reads 615 nonDup 600 readDis 0.707 363

Off

0.1026 0.1057

104 (40.471)

nonStacked

destacking?

C>T_1

G>A_-1

mean length (sd)