

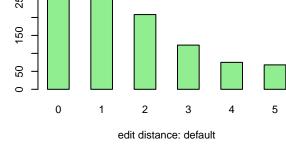
40 20

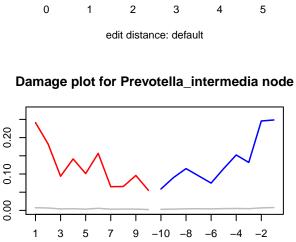
0

Node Top Reference

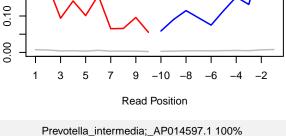
mean length (sd)

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





C2T/G2A rate



Prevotella_intermedia_17;_CP003503.1 100%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019300.1 076% revotella_intermedia_ATCC_25611_=_

Prevotella_dentalis_DSM_3688;_CP003369.1 000%

	_	_		_	.—		
termed	dia_ATC	C_256	11_=_DSI	M_2070	6;_CP0	19301.1	026%
Prevo	tella_int	ermedi	a_17;_CP	003502	2.1 020%		

Porphyromonas_gingivalis;_CP011995.1 003%
Tannerella_forsythia_KS16;_AP013045.1 001%



Prevotella_fusca_JCM_17724;_CP012074.1 000% Prevotella_fusca_JCM_17724;_CP012075.1 000% assigned reads 1148 all reads 556 nonDup 541 readDis 0.892 nonStacked 458 destacking? Off C>T_1 0.2412 G>A_-1 0.2484

2

edit distance: ancient

3

Prevotella_intermedia

Prevotella_intermedia

48 (15.126)