S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu ■ Node: Treponema_denticola 438 ■ Node: Treponema_denticola 3475 800 read count 9 400 20 0 2 3 0 1 2 3 edit distance: default edit distance: ancient Treponema_denticola Node Top Reference Treponema_denticola_ATCC_35405 Damage plot for Treponema_denticola node assigned reads 4204 all reads 3306 0.08 3222 nonDup readDis 0.039 0.04 nonStacked 65 Off destacking? C>T_1 0.1046

G>A_-1

mean length (sd)

0.0695

94 (39.003)

	0	I	- 1	ı	ı	- 1	ı	1	- 1	- 1			
		1	3	5	7	9	-10	-8	-6	-4	-2		
		Read Position											
	Tre	Treponema_denticola_ATCC_35405;_AE017226.1 100%											
		Т	repor	nema_	_AF12	22909	9.1 10	00%					
Treponema_denticola;_AF083949.1													
Treponema_denticola;_AF042861.1 1										00%			
)%											
		Т	repor	ema_	ma_denticola;_AF074950.1 100%								
		Т	repor	32219	9.1 10	00%							
Treponema_denticola_ATCC_33520;_AF04											100%		
Treponema_denticola;_NR_0762									2.1 1	00%			
		Т	repor	nema_	_dent	icola	_AF08	3251 ²	1.1 10	00%			