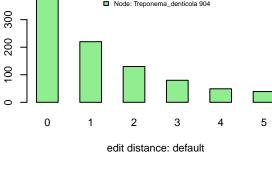
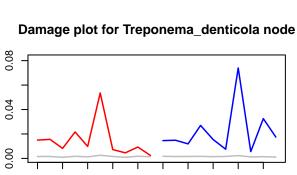
■ Node: Treponema_denticola 27 ■ Node: Treponema_denticola 904

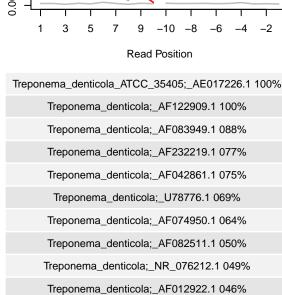
read count

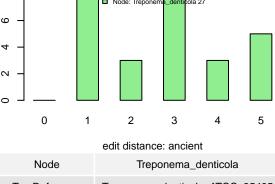
S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu





C2T/G2A rate





•••							
0 -	<u> </u>						
	0	1	2	2	3	4	5
edit distance: ancient							
Node			Treponema_denticola				
To	p Referer	nce	Treponema_denticola_ATCC_35405				
assigned reads			1046				
all reads			818				
nonDup			803				
readDis			0.26				
nonStacked			89				

nonStacked 89 destacking? Off C>T_1 0.015 G>A_-1 0.0175

mean length (sd)

105 (40.152)