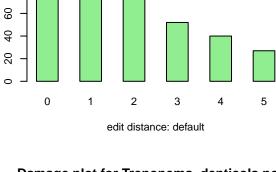
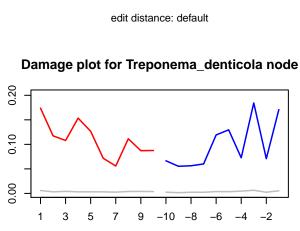
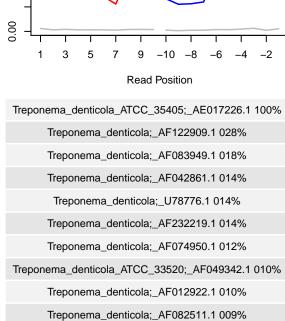
## ■ Node: Treponema\_denticola 112 Node: Treponema\_denticola 385 8 9

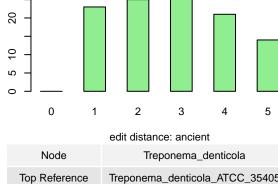
S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu





C2T/G2A rate





47 —							
0 -	J						
	0	1	2		3	4	5
			edit distance: ancient				
Node			Treponema_denticola				
Top Reference			Treponema_denticola_ATCC_35405				
assigned reads			421				
all reads			328				
nonDup			322				
readDis			0.773				
nonStacked			174				

1/4 nonStacked destacking? Off C>T\_1 0.1739 G>A\_-1 0.1709

mean length (sd)

58 (20.674)