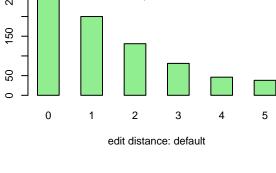
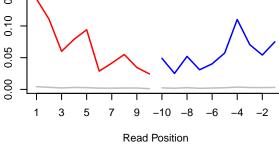
Lead Count reponema_denticola 766 Node: Treponema_denticola 109 Node: Treponema_denticola 109

0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un

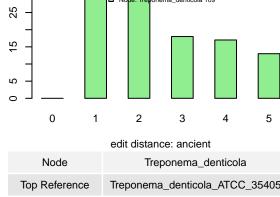






00]						
o						
1 3 5 7 9 -10 -8 -6 -4 -2						
Read Position						
Treponema_denticola_ATCC_35405;_AE017226.1 1						
Treponema_denticola;_AF122909.1 049%						
Treponema_denticola;_AF083949.1 040%						
Treponema_denticola;_AF042861.1 034%						
Treponema_denticola;_U78776.1 030%						
Treponema_denticola;_AF074950.1 020%						
Treponema_denticola;_AF232219.1 020%						
Treponema_denticola;_AF082511.1 020%						
Treponema_denticola;_NR_076212.1 018%						

Treponema_denticola_ATCC_33520;_AF049342.1 017%



	0	1	2	3	4	5
	edit distance: ancient					
	Node		Treponema_denticola			
	Top Reference		Treponema_denticola_ATCC_35405			
	assigned reads		861			
	all reads		685			
	nonDup		673			
	readDis			0.52	2	
	nonStacked	i		198		
	destacking?	?		Off		
	C>T_1			0.142	25	
					_	

C>T_1 0.1425
G>A_-1 0.075
mean length (sd) 68 (25.081)