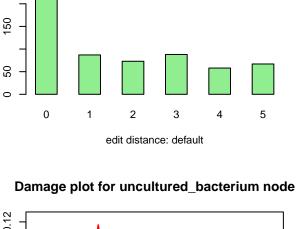
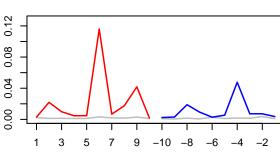
12 ■ Node: uncultured_bacterium 40 Node: uncultured_bacterium 608 read count ead count

 ∞ 9

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





Read Position uncultured_bacterium;_EU474155.1 100% Lactobacillus_gorillae;_AB904719.1 100% Lactobacillus_gorillae;_NR_134066.1 100% Lactobacillus_gorillae;_AB904718.1 100% Streptococcus_danieliae;_NR_117375.1 100% uncultured_Acetobacterium_sp.;_AY185310.1 100% uncultured_Alkaliphilus_sp.;_KF362056.1 100% uncultured_Streptococcus_sp.;_KJ910368.1 100% Bacteria; KM974662.1 100%

uncultured_Chloroflexi_bacterium;_HQ147612.1 100%

0	1		2		3		4	
edit distance: ancient								
Node			uncultured_bacterium					
Top Reference			uncultured_bacterium					
assigned reads			810					
all reads			21					
nonDup			5					
readDis			0.391					
nonStacked			1					
destacking?			Off					
C>T_1			0.003					
G>A1			0.0036					
mean length (sd)			64 (22.904)					