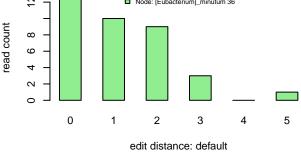
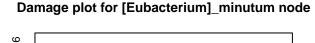
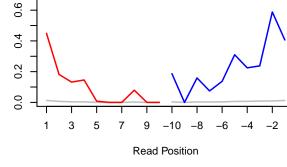
■ Node: [Eubacterium]_minutum 36 Node: [Eubacterium]_minutum 19 ead count

0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur







C2T/G2A rate

[Eubacterium]_minutum;_NR_036949.1 100% uncultured_Eubacterium_sp.;_AM420183.1 100%

Eubacterium_minutum_ATCC_700079;_NR_112052.1 100%

Eubacterium_minutum_ATCC_700079;_CP016202.1 100%

Eubacterium_minutum_ATCC_700079;_NR_114689.1 100%

uncultured_bacterium;_KF506073.1 100%

uncultured_Clostridium_sp.;_EF704000.1 100%

Eubacterium_sp._oral_taxon_G75;_GU432190.1 100%

uncultured_Eubacterium_sp.;_AM419990.1 100%

NA%

0	1	2	3	4	5	
edit distance: ancient						
Node		[Eu	[Eubacterium]_minutum			

edit distance: ancient				
Node	[Eubacterium]_minutum			
Top Reference	[Eubacterium]_minutum			
assigned reads	36			
all reads	34			
nonDup	33			
readDis	0.064			
nonStacked	1			
destacking?	Off			
C>T_1	NA			
G>A1	NA			
mean length (sd)	40 (8.605)			