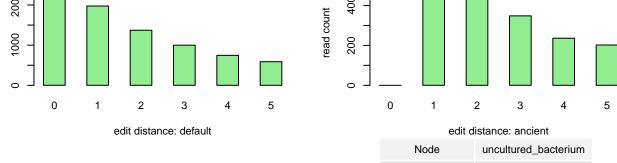
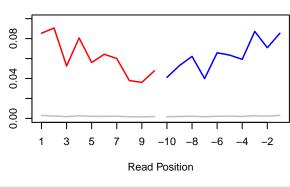
Node: uncultured_bacterium 8326 Node: uncultured_bacterium 1765

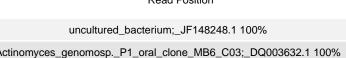
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un)101_S0_L000_R1_000.fastq.merged.prefixed.hg19un





C2T/G2A rate

Damage plot for uncultured_bacterium node



Bacteria;_FJ976437.1 100%
uncultured_Actinomyces_sp.;_FJ976431.1 100%

uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%

Ottowia_sp._oral_taxon_894;_CP012073.1 100%

Lautropia_sp._oral_clone_FX006;_AY134902.1 100%

uncultured_Lautropia_sp.;_AM419986.1 100%

uncultured_Actinomyces_sp.;_JQ406543.1 100%

Actinomyces;_NR_025633.1 100%

0	1	2	3	4
edit distance: ancient				
Node		uncultured_bacterium		
Top Reference		uncultured_bacterium		
assigned reads		10783		
all reads		239		
nonDup		236		
readDis		0.004		
nonStacked		0		
destacking?		Off		
C>T_1		0.0854		
G>A1		0.0854		
mean length (sd)		65 (26.092)		