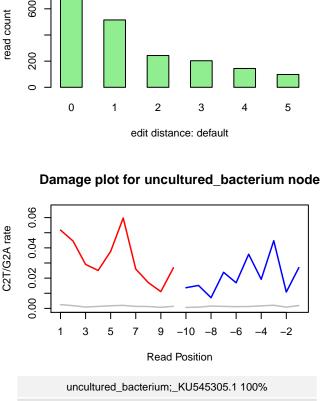
## ■ Node: uncultured\_bacterium 2016 Node: uncultured\_bacterium 227

ead count

9

0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



0	1		2	3	4	
	е	dit d	istance: ancient			
Node			uncultured_bacterium			
Top Reference			uncultured_bacterium			
assigned reads			2413			
all reads			133			
nonDup			123			
readDis			0.009			
nonStacked			0			
destacking?			Off			
C>T_1			0.0518			
G>A1			0.027			
mean length (sd)			59 (22.122)			

Actinomyces_spCM84;_HQ610192.1 100%
Actinomyces_sporal_clone_EP011;_AY008315.1 100%
Actinomyces_naeslundii;_GU561309.1 100%

Actinomyces\_naeslundii;\_AB062278.1 100% Actinomyces\_naeslundii;\_FJ823149.1 100%

Actinomyces\_naeslundii;\_AJ234045.1 100%

uncultured\_Rothia\_sp.;\_KC675369.1 100% Actinomyces\_naeslundii;\_NR\_037033.1 100%

uncultured\_Actinomycetales\_bacterium;\_HQ848539.1 100%